Table _____. U-Pb geochronologic analyses.

Trial 1: CP-40 Nu Plasma HR TC

						Isotope ratios					Apparent ages (Ma)									
Analysis	U (ppm)	206Pb / 204Pb	U/Th	206Pb* / 207Pb*	± (%)	207Pb* / 235U*	± (%)	206Pb* / 238U	± (%)	error corr.	206Pb* / 238U*	± (Ma)	207Pb* / 235U	± (Ma)	206Pb* / 207Pb*	± (Ma)	Best age (Ma)	± (Ma)	Conc (%)
Trial-1_0001	131	231347	0.9	8.7160	2.1	5.3927	2.8	0.3409	1.8	0.64	1891.0	29.6	1883.7	24.0	1875.7	38.6	1875.7	38.6	100.8
Trial-1_0002	53	15705	2.0	12.4451	5.0	2.1312	6.4	0.1924	4.0	0.63	1134.2	41.4	1159.0	44.0	1205.6	97.8	1205.6	97.8	94.1
Trial-1_0003	182	336854	1.0	5.8101	1.5	11.2225	2.0	0.4729	1.3	0.66	2496.3	27.3	2541.8	18.6	2578.3	25.0	2578.3	25.0	96.8
Trial-1_0004	74	30673	1.9	10.0016	3.1	3.9334	4.1	0.2853	2.6	0.64	1618.1	37.2	1620.6	33.0	1623.8	58.4	1623.8	58.4	99.7
Trial-1_0005	67	60994	0.9	18.3419	7.5	0.4856	9.4	0.0646	5.7	0.60	403.5	22.3	401.9	31.3	392.6	169.0	403.5	22.3	102.8
Trial-1_0006	244	176926	3.3	14.2836	2.5	1.4752	3.2	0.1528	2.0	0.62	916.7	16.7	920.3	19.1	928.6	50.8	928.6	50.8	98.7
Trial-1_0007	50	302936	2.1	12.7120	4.8	2.0938	6.2	0.1930	3.9	0.63	1137.8	40.4	1146.8	42.6	1163.7	95.8	1163.7	95.8	97.8
Trial-1_0008	202	146780	2.3	13.3415	2.5	1.8766	3.2	0.1816	2.0	0.62	1075.6	19.6	1072.9	21.0	1067.3	50.0	1067.3	50.0	100.8
Trial-1_0009	179	106782	1.4	12.3822	2.5	2.2451	3.2	0.2016	2.0	0.63	1184.0	21.5	1195.3	22.3	1215.7	48.7	1215.7	48.7	97.4
Trial-1_0010	240	263580	9.4	12.8129	2.2	2.0751	2.8	0.1928	1.8	0.62	1136.7	18.4	1140.6	19.4	1148.1	43.9	1148.1	43.9	99.0
Trial-1_0011	144	453772	0.7	5.1413	1.6	14.3125	2.1	0.5337	1.4	0.67	2757.0	31.6	2770.6	20.1	2780.6	25.9	2780.6	25.9	99.1
Trial-1_0012	69	138466	0.6	8.8109	3.0	5.3167	3.9	0.3398	2.5	0.64	1885.5	41.1	1871.5	33.4	1856.1	53.9	1856.1	53.9	101.6
Trial-1_0013	38	79577	1.6	13.5238	5.8	1.8046	7.4	0.1770	4.6	0.62	1050.6	44.8	1047.1	48.5	1039.9	117.4	1039.9	117.4	101.0
Trial-1_0014	124	175480	1.2	13.1066	3.2	1.9099	4.0	0.1816	2.5	0.62	1075.5	25.0	1084.6	27.0	1102.9	63.2	1102.9	63.2	97.5
Trial-1_0015	36	102289	2.8	9.5520	4.4	4.3629	5.8	0.3023	3.7	0.64	1702.5	55.4	1705.3	47.8	1708.9	81.9	1708.9	81.9	99.6
Trial-1_0016	170	83954	5.8	13.6964	2.9	1.6884	3.6	0.1677	2.3	0.62	999.5	21.0	1004.1	23.2	1014.3	57.9	1014.3	57.9	98.5
Trial-1_0017	79	132434	0.7	5.2719	2.2	13.8578	2.9	0.5299	1.9	0.66	2740.9	42.7	2740.0	27.3	2739.4	35.4	2739.4	35.4	100.1
Trial-1_0018	217	27839	2.5	19.7489	5.3	0.2850	6.7	0.0408	4.0	0.60	258.0	10.1	254.7	15.0	224.3	123.5	258.0	10.1	NA
Trial-1_0019	41	16211	1.3	11.1023	4.6	3.1256	5.9	0.2517	3.7	0.63	1447.1	48.4	1439.0	45.4	1427.0	87.2	1427.0	87.2	101.4
Trial-1_0020	81	12922	2.5	20.0618	9.1	0.2671	11.3	0.0389	6.8	0.60	245.8	16.3	240.4	24.2	187.8	211.2	245.8	16.3	NA
Trial-1_0021	44	26611	0.6	12.8574	5.1	2.1456	6.5	0.2001	4.1	0.63	1175.8	43.8	1163.6	45.2	1141.1	101.1	1141.1	101.1	103.0
Trial-1_0022	329	246795	4.5	13.6796	2.1	1.6500	2.6	0.1637	1.6	0.62	977.3	14.9	989.5	16.7	1016.7	41.9	1016.7	41.9	96.1
Trial-1_0023	125	80152	1.3	13.4163	3.1	1.9028	4.0	0.1852	2.5	0.62	1095.1	25.3	1082.1	26.8	1056.0	63.4	1056.0	63.4	103.7
Trial-1_0024	39	106952	1.2	17.7340	8.6	0.6548	10.8	0.0842	6.5	0.61	521.2	32.8	511.4	43.4	467.7	190.1	521.2	32.8	111.4
Trial-1_0025	91	17743	0.6	19.9049	7.8	0.3120	9.7	0.0450	5.8	0.60	284.0	16.2	275.7	23.5	206.1	181.5	284.0	16.2	NA
Trial-1_0026	196	36792	1.2	19.2405	5.4	0.3186	6.7	0.0445	4.0	0.60	280.5	11.0	280.9	16.4	284.2	122.7	280.5	11.0	NA
Trial-1_0027	44	8188	0.5	17.8897	7.7	0.7330	9.7	0.0951	5.9	0.61	585.7	32.8	558.3	41.6	448.3	171.4	585.7	32.8	130.6
Trial-1_0028	81	62592	2.7	12.7593	3.7	2.1521	4.8	0.1992	3.0	0.63	1170.8	32.1	1165.7	33.2	1156.4	74.1	1156.4	74.1	101.2
Trial-1_0029	108	79328	1.3	11.0618	2.8	3.1661	3.7	0.2540	2.3	0.63	1459.1	30.3	1448.9	28.2	1434.0	54.0	1434.0	54.0	101.8
Trial-1_0030	36	4316	1.6	19.8111	10.0	0.5003	12.5	0.0719	7.5	0.60	447.5	32.5	411.9	42.4	217.0	232.0	447.5	32.5	206.2
Trial-1_0031	71	167145	0.7	18.2329	7.2	0.5042	9.0	0.0667	5.5	0.60	416.1	22.0	414.5	30.8	405.9	161.4	416.1	22.0	102.5
Trial-1_0032	73	116930	2.0	4.9595	2.2	15.2316	2.9	0.5479	1.9	0.67	2816.3	44.4	2829.8	27.8	2839.4	35.4	2839.4	35.4	99.2
Trial-1_0033	27	28459	1.1	3.7560	3.2	25.1062	4.3	0.6839	2.9	0.68	3359.4	76.1	3312.4	41.9	3284.0	49.6	3284.0	49.6	102.3
Trial-1_0034	11	20103	0.5	10.8011	8.6	3.4200	11.2	0.2679	7.1	0.64	1530.2	96.6	1509.0	87.9	1479.4	163.7	1479.4	163.7	103.4
Trial-1_0035	105	216266	2.1	5.1377	1.8	14.7565	2.4	0.5499	1.6	0.67	2824.6	37.1	2799.7	23.2	2781.7	29.8	2781.7	29.8	101.5
Trial-1_0036	40	22547	1.7	13.8760	5.8	1.7177	7.5	0.1729	4.6	0.62	1027.9	44.0	1015.2	47.9	987.8	118.8	987.8	118.8	104.1
Trial-1_0037	129	209540	2.8	11.2588	2.7	2.9344	3.4	0.2396	2.2	0.63	1384.7	27.2	1390.8	26.1	1400.2	51.2	1400.2	51.2	98.9
Trial-1_0038	27	60626	0.8	5.4258	3.6	13.6025	4.8	0.5353	3.2	0.66	2763.6	72.3	2722.4	45.8	2692.0	59.7	2692.0	59.7	102.7
Trial-1_0039	288	154166	2.9	13.4259	2.1	1.8489	2.7	0.1800	1.7	0.62	1067.2	16.4	1063.0	17.6	1054.6	42.1	1054.6	42.1	101.2
Trial-1_0040	254	409311	3.1	12.6707	2.1	2.1495	2.7	0.1975	1.7	0.63	1162.0	18.1	1164.9	18.8	1170.2	42.0	1170.2	42.0	99.3
Trial-1_0041	300	13309	0.7	20.6777	5.6	0.1819	6.9	0.0273	4.1	0.60	173.5	7.1	169.7	10.9	117.0	131.7	173.5	7.1	NA
Trial-1_0042	344	64269	0.9	19.5134	4.2	0.2866	5.3	0.0406	3.2	0.60	256.3	8.0	255.9	12.0	252.0	97.7	256.3	8.0	NA
Trial-1_0043	46	18173	1.6	13.2085	5.1	1.9905	6.5	0.1907	4.1	0.62	1125.1	41.9	1112.3	44.0	1087.4	101.8	1087.4	101.8	103.5
Trial-1_0044	253	103297	18.0	13.0910	2.2	1.9537	2.8	0.1855	1.8	0.62	1096.9	17.7	1099.7	18.9	1105.3	44.0	1105.3	44.0	99.2
Trial-1_0045	118	120251	2.9	13.4921	3.3	1.8388	4.2	0.1799	2.6	0.62	1066.6	26.0	1059.4	27.9	1044.7	67.0	1044.7	67.0	102.1
Trial-1_0046	27	28221	0.5	5.3725	3.7	13.7522	4.9	0.5359	3.3	0.66	2766.1	73.1	2732.8	46.3	2708.3	60.3	2708.3	60.3	102.1
Trial-1_0047	285	648860	2.4	13.2499	2.1	1.8861	2.7	0.1812	1.7	0.62	1073.8	16.5	1076.2	17.8	1081.1	42.0	1081.1	42.0	99.3
Trial-1_0048	60	43134	2.4	13.1733	4.5	2.0186	5.7	0.1929	3.6	0.62	1136.9	37.0	1121.8	38.7	1092.7	89.2	1092.7	89.2	104.0
Trial-1_0049	74	27606	1.7	17.4165	6.2	0.6959	7.7	0.0879	4.7	0.61	543.1	24.5	536.3	32.3	507.6	135.5	543.1	24.5	107.0
Trial-1_0050	114	106086	1.8	17.4916	5.0	0.6918	6.3	0.0878	3.8	0.61	542.3	19.7	533.9	26.0	498.1	109.7	542.3	19.7	108.9
Trial-1_0051	101	21512	1.0	20.3982	7.7	0.2934	9.6	0.0434	5.8	0.60	273.9	15.5	261.2	22.2	149.0	181.6	273.9	15.5	NA
Trial-1_0052	469	115822	4.6	16.8531	2.3	0.7947	2.9	0.0971	1.8	0.61	597.6	10.2	593.8	13.2	579.5	50.6	597.6	10.2	103.1
Trial-1_0053	82	105095	4.3	12.7820	3.8	2.0950	4.8	0.1942	3.0	0.63	1144.1	31.8	1147.2	33.3	1152.8	75.1	1152.8	75.1	99.2
Trial-1_0054	280	143245	9.6	12.4893	2.1	2.1388	2.7	0.1937	1.7	0.63	1141.6	17.4	1161.4	18.4	1198.7	40.9	1198.7	40.9	95.2
Trial-1_0055	316	400455	1.9	8.9067	1.4	5.2776	1.9	0.3409	1.2	0.64	1891.1	19.6	1865.2	15.8	1836.5	25.7	1836.5	25.7	103.0
Trial-1_0056	27	38283	0.3	5.5539	3.8	12.8462	5.1	0.5175	3.4	0.66	2688.4	74.2	2668.4	48.0	2653.4	63.2	2653.4	63.2	101.3
Trial-1_0057	183	190739	1.1	13.4960	2.7	1.7956	3.5	0.1758	2.2	0.62	1043.7	20.8	1043.9	22.6	1044.1	54.7	1044.1	54.7	100.0
Trial-1_0058	38	19784	2.2	9.1800	4.2	4.9510	5.5	0.3296	3.5	0.64	1836.6	56.7	1811.0	46.7	1781.6	77.3	1781.6	77.3	103.1
Trial-1_0059	19	13281	1.3	13.2097	8.0	2.0000	10.3	0.1916	6.4	0.62	1130.1	66.7	1115.5	69.8	1087.2	161.4	1087.2	161.4	103.9
Trial-1_0060	220	238594	4.8	13.4724	2.4	1.8515	3.1	0.1809	1.9	0.62	1072.0	19.2	1064.0	20.6	1047.6	49.2	1047.6	49.2	102.3
Trial-1_0061	177	5341	1.0	22.4914	10.6	0.0916	12.9	0.0149	7.3	0.57	95.6	7.0	89.0	11.0	-85.1	260.6	95.6	7.0	NA
Trial-1_0062	55	34088	1.5	14.1065	5.2	1.6315	6.6	0.1669	4.1	0.62	995.1	37.7	982.4	41.5	954.2	105.8	954.2	105.8	104.3
Trial-1_0063	69	56847	0.8	10.8398	3.6	3.4201	4.6	0.2689	2.9	0.63	1535.1	39.9	1509.0	36.2	1472.6	67.6	1472.6	67.6	104.2
Trial-1_0064	108	209411	1.2	13.0633	3.4	1.9887	4.4	0.1884	2.7	0.62	1112.8	28.1	1111.7	29.7	1109.5	68.7	1109.5	68.7	100.3
Trial-1_0065	61	34768	0.9	13.2425	4.5	1.9803	5.8	0.1902	3.6	0.62	1122.4	37.4	1108.8	39.3	1082.2	91.3	1082.2	91.3	103.7
Trial-1_0066	15	9273	1.5	9.9512	7.1	4.0423	9.3	0.2917	5.9	0.64	1650.2	86.0	1642.7	75.5	1633.2	132.5	1633.2	132.5	101.0
Trial-1_0067	239	57478	1.2	18.0741	3.9	0.5740	4.9	0.0752	3.0	0.60	467.7	13.5	460.6	18.3	425.5	87.8	467.7	13.5	109.9
Trial-1_0068	17	6723	1.5	13.6903	9.2	1.7307	11.7	0.1718	7.3	0.62	1022.3	68.9	1020.0	75.5	1015.2	186.1	1015.2	186.1	100.7
Trial-1_0069	285	380943	5.4	9.3853	1.7	3.9172	2.3	0.2666	1.4	0.64	1523.7	19.6	1617.2	18.3	1741.2	31.8	1741.2	31.8	87.5
Trial-1_0070	398	61646	1.6	8.5354	1.4	4.6336	1.8	0.2868	1.2	0.64	1625.7	16.6	1755.3	15.0	1913.3	24.6	1913.3	24.6	85.0
Trial-1_0071	73	5668	1.4	21.5574	10.8	0.1885	13.4	0.0295	7.9	0.59	187.2	14.6	175.3	21.5	17.8	259.0	187.2	14.6	NA
Trial-1_0072	1219	259868	8.5	13.0048	1.1	1.7327	1.4	0.1634	0.9	0.62	975.8	7.9	1020.7	9.0	1118.5	21.8	1118.5	21.8	87.2
Trial-1_0073	99	237434	0.8	5.4925	2.0	13.2669	2.6	0.5285	1.7	0.66	2735.1	39.0	2698.8	24.9	2671.8	32.6	2671.8	32.6	102.4
Trial-1_0074	57	81839	1.0	13.6266	4.9	1.7833	6.2	0.1762	3.9	0.62	1046.4	37.5	1039.4	40.6	1024.6	98.9	1024.6	98.9	102.1
Trial-1_0075	280	109988	2.0	13.2614	2.1	1.9176	2.7	0.1844	1.7	0.62	1091.2	17.2	1087.2	18.3	1079.4	43.1	1079.4	43.1	101.1
Trial-1_0076	102	104741	1.2	5.8762	2.0	11.4902	2.7	0.4897	1.8	0.66	2569.4	37.6	2563.8	25.1	2559.4	33.7	2559.4	33.7	100.4
Trial-1_0077	200	40251	2.0	20.1489	6.8	0.1962	8.5	0.0287	5.0	0.60	182.2	9.1	181.9	14.1	177.7	158.4	182.2	9.1	NA
Trial-1_0078	380	125745	2.7	13.5265	1.9	1.7935	2.4	0.1759	1.5	0.62	1044.8	14.6	1043.1	15.9	1039.5	38.6	1039.5	38.6	100.5
Trial-1_0079	252	111314	2.2	13.3462	2.3	1.8682	2.9	0.1808	1.8	0.62	1071.5	17.8	1069.9	19.2	1066.6	45.6	1066.6	45.6	100.5
Trial-1_0080	151	35026	0.8	19.2209	5.9	0.3593	7.4	0.0501	4.5	0.60	315.1	13.7	311.7	19.9	286.6	135.4	315.1	13.7	NA
Trial-1_0081	48	48867	0.5	5.1336	2.7	14.7406	3.7	0.5488	2.4	0.67	2820.3	55.9	2798.6	34.9	2783.1	44.9	2783.1	44.9	101.3
Trial-1_0082	280	62808	3.7	13.9144	2.3	1.6650	2.9	0.1680	1.8	0.62	1001.2	16.7	995.3	18.4	982.2	46.2	982.2	46.2	101.9
Trial-1_0083	87	120193	1.5	13.1568	3.8	2.0153	4.8	0.1923	3.0	0.62	1133.8	31.3	1120.7	32.7	1095.3	75.4	1095.3	75.4	103.5
Trial-1_0084	210	97775	2.8	8.6145	1.7	5.5520	2.3	0.3469	1.5	0.64	1919.7	24.4	1908.7	19.6	1896.7	31.3	1896.7	31.3	101.2
Trial-1_0085	83	16079	2.1	17.5590	6.1	0.6776	7.6	0.0863	4.6	0.61	533.6	23.7	525.3	31.3	489.7	133.9	533.6	23.7	109.0
Trial-1_0086	48	23648	1.1	9.6499	3.8	4.3115	5.0	0.3018	3.2	0.64	1700.0	47.8	1695.6	41.3	1690.1	70.9	1690.1	70.9	100.6
Trial-1_0087	127	155627	1.8	12.1942	2.9	2.4249	3.7	0.2145	2.3	0.63	1252.5	26.5	1250.0	26.6	1245.6	56.5	1245.6	56.5	100.6
Trial-1_0088	238	26084	2.1	20.3922	6.0	0.2017	7.4	0.0298	4.4	0.60	189.5	8.3	186.6	12.7	149.7	140.3	189.5	8.3	NA
Trial-1_0089	255	55378	1.1	19.3797	5.0	0.2902	6.2	0.0408	3.7	0.60	257.7	9.4	258.7	14.2	267.7	114.2	257.7	9.4	NA
Trial-1_0090	93	175798	1.1	11.0093	3.0	3.2224	3.9	0.2573	2.5	0.63	1476.0	32.9	1462.6	30.5	1443.1	58.0	1443.1	58.0	102.3
Trial-1_0091	70	128849	1.8	9.3715	3.1	4.6835	4.0	0.3183	2.6	0.64	1781.6	40.2	1764.3	33.8	1743.9	56.7	1743.9	56.7	102.2
Trial-1_0092	187	31196	1.7	18.0845	4.2	0.5640	5.3	0.0740	3.2	0.60	460.1	14.2	454.1	19.4	424.2	94.4	460.1	14.2	108.5
Trial-1_0093	258	33756	3.4	19.7564	5.0	0.2750	6.3	0.0394	3.7	0.60	249.1	9.2	246.7	13.7	223.4	116.3	249.1	9.2	NA
Trial-1_0094	45	17508	1.5	10.4686	4.3	3.4947	5.6	0.2653	3.6	0.64	1517.1	48.0	1526.0	44.2	1538.4	81.2	1538.4	81.2	98.6
Trial-1_0095	45	119097	3.1	12.2384	4.8	2.4384	6.2	0.2164	3.9	0.63	1263.0	44.8	1254.0	44.7	1238.6	94.8	1238.6	94.8	102.0
Trial-1_0096	74	134151	3.1	9.0214	3.0	5.0716	3.9	0.3318	2.5	0.64	1847.3	39.9	1831.4	32.8	1813.3	53.8	1813.3	53.8	101.9
Trial-1_0097	118	537969	1.8	13.6993	3.0	1.8041	3.9	0.1792	2.4	0.62	1062.9	23.6	1046.9	25.2	1013.8	61.2	1013.8	61.2	104.8
Trial-1_0098	117	230291	0.8	8.7672	2.3	5.3766	3.0	0.3419	1.9	0.64	1895.7	31.8	1881.1	25.7	1865.1	41.5	1865.1	41.5	101.6
Trial-1_0099	352	161317	1.1	8.9742	1.4	5.0141	1.8	0.3264	1.2	0.64	1820.7	18.4	1821.7	15.2	1822.9	25.0	1822.9	25.0	99.9
Trial-1_0100	8	6373	1.6	7.7591	8.3	6.3993	10.9	0.3601	7.1	0.65	1982.7	120.7	2032.2	96.3	2082.8	147.0	2082.8	147.0	95.2
Trial-1_0101	188	107524	1.3	18.9559	5.0	0.4027	6.2	0.0554	3.7	0.60	347.4	12.6	343.6	18.1	318.2	112.8	347.4	12.6	NA
Trial-1_0102	175	676422	2.1	10.1571	2.1	3.7650	2.7	0.2774	1.8	0.64	1578.0	24.5	1585.3	22.1	1595.0	39.6	1595.0	39.6	98.9
Trial-1_0103	68	20854	1.7	13.3816	4.4	1.8732	5.6	0.1818	3.5	0.62	1076.8	34.4	1071.7	36.9	1061.3	87.8	1061.3	87.8	101.5
Trial-1_0104	32	30991	1.3	13.6832	6.4	1.8023	8.1	0.1789	5.1	0.62	1060.7	49.6	1046.3	53.3	1016.2	129.4	1016.2	129.4	104.4
Trial-1_0105	73	54203	1.4	13.3033	4.1	1.9550	5.3	0.1886	3.3	0.62	1113.9	33.5	1100.2	35.3	1073.1	82.5	1073.1	82.5	103.8
Trial-1_0106	632	90466	1.1	18.7101	2.7	0.4077	3.4	0.0553	2.0	0.60	347.1	6.8	347.2	9.9	347.8	60.6	347.1	6.8	NA
Trial-1_0107	54	27494	2.0	13.3247	4.8	1.9121	6.2	0.1848	3.9	0.62	1093.1	38.8	1085.3	41.3	1069.8	97.3	1069.8	97.3	102.2
Trial-1_0108	207	15170	1.2	20.2453	5.7	0.2619	7.1	0.0385	4.3	0.60	243.2	10.2	236.2	15.1	166.6	134.1	243.2	10.2	NA
Trial-1_0109	190	161064	0.7	10.9678	2.1	3.2826	2.7	0.2611	1.7	0.63	1495.6	23.2	1476.9	21.3	1450.3	40.4	1450.3	40.4	103.1
Trial-1_0110	112	46309	2.6	13.7972	3.5	1.7555	4.4	0.1757	2.7	0.62	1043.3	26.4	1029.2	28.5	999.4	70.2	999.4	70.2	104.4
Trial-1_0111	114	307632	1.0	10.9691	2.8	3.2405	3.6	0.2578	2.3	0.63	1478.6	30.0	1466.9	27.8	1450.0	52.8	1450.0	52.8	102.0
Trial-1_0112	139	396513	0.7	9.8072	2.3	4.1688	3.0	0.2965	1.9	0.64	1674.0	28.1	1667.9	24.4	1660.2	42.5	1660.2	42.5	100.8
Trial-1_0113	72	34063	1.5	11.1429	3.5	3.1937	4.5	0.2581	2.8	0.63	1480.1	37.6	1455.6	34.8	1420.1	66.5	1420.1	66.5	104.2
Trial-1_0114	157	133058	0.7	10.9552	2.4	3.2993	3.0	0.2621	1.9	0.63	1500.8	25.8	1480.9	23.8	1452.5	44.9	1452.5	44.9	103.3
Trial-1_0115	168	75511	3.1	13.4848	2.7	1.8873	3.5	0.1846	2.2	0.62	1091.9	21.8	1076.6	23.2	1045.8	55.1	1045.8	55.1	104.4
Trial-1_0116	65	8082	0.9	19.7941	9.8	0.2882	12.2	0.0414	7.3	0.60	261.3	18.8	257.1	27.8	219.0	226.9	261.3	18.8	NA
Trial-1_0117	123	67296	1.3	13.4447	3.2	1.9084	4.1	0.1861	2.5	0.62	1100.1	25.7	1084.0	27.2	1051.8	64.4	1051.8	64.4	104.6
Trial-1_0118	60	86572	1.0	13.5423	4.7	1.8108	6.0	0.1779	3.7	0.62	1055.2	36.2	1049.4	39.1	1037.2	94.7	1037.2	94.7	101.7
Trial-1_0119	89	193070	0.7	16.4634	5.1	0.8853	6.5	0.1057	3.9	0.61	647.8	24.3	643.8	30.8	630.1	110.3	647.8	24.3	102.8
Trial-1_0120	68	137463	0.9	5.4956	2.4	13.2164	3.1	0.5268	2.1	0.66	2727.8	46.4	2695.2	29.7	2670.9	39.0	2670.9	39.0	102.1
Trial-1_0121	176	136393	2.2	13.4353	2.7	1.8712	3.4	0.1823	2.1	0.62	1079.7	21.3	1071.0	22.8	1053.2	54.3	1053.2	54.3	102.5
Trial-1_0122	124	37563	1.1	13.3750	3.2	1.8871	4.1	0.1831	2.5	0.62	1083.6	25.4	1076.5	27.1	1062.2	64.3	1062.2	64.3	102.0
Trial-1_0123	64	53087	1.8	13.2595	4.5	1.8600	5.8	0.1789	3.6	0.62	1060.8	35.1	1067.0	38.0	1079.7	90.4	1079.7	90.4	98.3
Trial-1_0124	124	74415	2.6	13.4898	3.3	1.7782	4.2	0.1740	2.6	0.62	1034.0	24.9	1037.5	27.2	1045.0	66.1	1045.0	66.1	98.9
Trial-1_0125	98	508900	1.1	5.3530	2.0	13.2858	2.6	0.5158	1.7	0.66	2681.4	38.2	2700.2	24.8	2714.3	32.4	2714.3	32.4	98.8
Trial-1_0126	188	326831	0.9	13.1147	2.6	1.9292	3.3	0.1835	2.1	0.62	1086.1	20.6	1091.3	22.1	1101.7	51.7	1101.7	51.7	98.6
Trial-1_0127	106	171826	1.5	9.1149	2.5	4.8165	3.2	0.3184	2.1	0.64	1781.9	32.4	1787.8	27.2	1794.6	45.2	1794.6	45.2	99.3
Trial-1_0128	42	35478	1.9	13.0788	5.4	1.9914	6.9	0.1889	4.3	0.62	1115.4	44.0	1112.6	46.5	1107.1	107.5	1107.1	107.5	100.7
Trial-1_0129	714	274021	1.7	17.5497	2.1	0.5885	2.7	0.0749	1.6	0.61	465.7	7.3	469.9	10.1	490.8	47.0	465.7	7.3	94.9
Trial-1_0130	342	70627	1.1	18.3005	3.4	0.4805	4.2	0.0638	2.6	0.60	398.6	9.9	398.4	14.0	397.6	75.8	398.6	9.9	NA
Trial-1_0131	39	58442	1.6	5.7218	3.2	12.0175	4.2	0.4987	2.8	0.66	2608.2	59.8	2605.8	39.5	2603.8	52.6	2603.8	52.6	100.2

Trial-1_0132	181	98951	1.8	16.2338	3.5	0.9227	4.4	0.1086	2.7	0.61	664.8	17.0	663.8	21.4	660.2	74.5	664.8	17.0	100.7
Trial-1_0133	103	152094	1.6	3.6467	1.6	26.2020	2.2	0.6930	1.5	0.68	3394.0	38.7	3354.1	21.2	3330.3	24.9	3330.3	24.9	101.9
Trial-1_0134	110	6771	1.2	19.9535	7.7	0.2668	9.7	0.0386	5.8	0.60	244.2	13.8	240.2	20.7	200.4	180.0	244.2	13.8	NA
Trial-1_0135	89	126596	2.2	13.5220	3.8	1.8096	4.9	0.1775	3.0	0.62	1053.1	29.5	1048.9	31.9	1040.2	77.1	1040.2	77.1	101.2
Trial-1_0136	17	9377	1.7	12.8338	8.5	2.1058	10.9	0.1960	6.8	0.62	1153.8	71.6	1150.7	75.0	1144.8	169.4	1144.8	169.4	100.8
Trial-1_0137	237	76266	0.7	13.2965	2.3	1.9255	2.9	0.1857	1.8	0.62	1098.0	18.4	1090.0	19.6	1074.1	46.1	1074.1	46.1	102.2
Trial-1_0138	93	34990	1.6	13.7578	3.7	1.7498	4.8	0.1746	3.0	0.62	1037.4	28.4	1027.1	30.9	1005.2	75.9	1005.2	75.9	103.2
Trial-1_0139	63	150528	0.9	5.4624	2.4	13.4889	3.2	0.5344	2.1	0.66	2759.9	47.3	2714.5	30.0	2680.9	39.3	2680.9	39.3	102.9
Trial-1_0140	129	91110	1.7	8.4061	2.1	5.6529	2.8	0.3446	1.8	0.65	1908.9	29.9	1924.2	24.2	1940.7	38.3	1940.7	38.3	98.4
Trial-1_0141	52	8546	2.9	21.4169	13.2	0.1749	16.4	0.0272	9.7	0.59	172.8	16.5	163.7	24.8	33.5	317.4	172.8	16.5	NA
Trial-1_0142	51	8172	1.9	19.3319	8.3	0.4913	10.4	0.0689	6.3	0.60	429.4	26.0	405.8	34.8	273.4	190.7	429.4	26.0	157.1
Trial-1_0143	13	3960	0.9	13.3079	9.6	1.9556	12.3	0.1888	7.6	0.62	1114.6	78.2	1100.4	82.6	1072.4	193.3	1072.4	193.3	103.9
Trial-1_0144	399	57856	3.4	18.2300	3.0	0.5088	3.8	0.0673	2.3	0.60	419.7	9.3	417.6	12.9	406.3	67.2	419.7	9.3	103.3
Trial-1_0145	256	770788	0.7	11.2110	1.9	2.8924	2.5	0.2352	1.6	0.63	1361.6	19.0	1379.9	18.5	1408.4	36.4	1408.4	36.4	96.7
Trial-1_0146	82	82392	2.0	13.2059	3.8	2.0043	4.8	0.1920	3.0	0.62	1132.0	31.2	1117.0	32.6	1087.8	75.3	1087.8	75.3	104.1
Trial-1_0147	6	16292	3.8	8.4051	9.4	6.0714	12.3	0.3701	7.9	0.65	2029.9	138.2	1986.1	107.4	1940.9	167.8	1940.9	167.8	104.6
Trial-1_0148	284	182420	1.8	17.9968	3.5	0.5352	4.4	0.0699	2.7	0.60	435.3	11.2	435.3	15.6	435.0	77.9	435.3	11.2	100.1
Trial-1_0149	277	316787	2.0	9.6125	1.8	3.5693	2.3	0.2488	1.5	0.64	1432.5	18.7	1542.7	18.1	1697.2	32.4	1697.2	32.4	84.4
Trial-1_0150	88	71456	1.0	10.9992	3.1	3.2277	4.0	0.2575	2.5	0.63	1477.0	33.5	1463.8	31.1	1444.8	59.1	1444.8	59.1	102.2
Trial-1_0151	73	77164	1.6	9.7571	3.1	4.2429	4.1	0.3003	2.6	0.64	1692.6	38.6	1682.4	33.3	1669.7	57.7	1669.7	57.7	101.4
Trial-1_0152	247	129749	2.1	13.4855	2.3	1.8047	2.9	0.1765	1.8	0.62	1047.9	17.6	1047.2	19.0	1045.7	46.0	1045.7	46.0	100.2
Trial-1_0153	300	188757	1.4	18.1286	3.5	0.5148	4.4	0.0677	2.6	0.60	422.2	10.7	421.7	15.0	418.7	77.5	422.2	10.7	100.8
Trial-1_0154	104	259973	1.3	11.0420	2.9	3.1301	3.7	0.2507	2.4	0.63	1441.9	30.5	1440.1	28.7	1437.4	55.1	1437.4	55.1	100.3
Trial-1_0155	54	310611	2.1	9.4363	3.6	4.5487	4.6	0.3113	3.0	0.64	1747.1	45.6	1739.9	38.7	1731.3	65.5	1731.3	65.5	100.9
Trial-1_0156	75	796409	1.1	5.3388	2.2	13.6578	2.9	0.5288	2.0	0.66	2736.5	43.6	2726.3	27.8	2718.7	36.2	2718.7	36.2	100.7
Trial-1_0157	206	83589	1.1	19.5856	5.1	0.3308	6.4	0.0470	3.8	0.60	296.0	11.1	290.2	16.1	243.5	117.9	296.0	11.1	NA
Trial-1_0158	192	74916	2.6	13.6298	2.6	1.7643	3.3	0.1744	2.1	0.62	1036.4	19.9	1032.4	21.7	1024.1	53.0	1024.1	53.0	101.2
Trial-1_0159	57	46588	1.8	12.8617	4.4	2.1567	5.7	0.2012	3.6	0.63	1181.7	38.5	1167.2	39.5	1140.5	88.4	1140.5	88.4	103.6
Trial-1_0160	181	108601	1.7	9.1579	2.1	4.1296	2.7	0.2743	1.7	0.64	1562.5	24.0	1660.2	22.1	1786.0	37.7	1786.0	37.7	87.5
Trial-1_0161	443	565124	1.7	12.9605	1.7	1.8851	2.2	0.1772	1.4	0.62	1051.6	13.2	1075.9	14.5	1125.3	34.0	1125.3	34.0	93.5
Trial-1_0162	399	477341	1.4	11.0030	1.5	3.1474	1.9	0.2512	1.2	0.63	1444.5	15.8	1444.4	14.8	1444.2	28.4	1444.2	28.4	100.0
Trial-1_0163	28	37076	1.2	4.5006	3.4	18.2464	4.5	0.5956	3.0	0.67	3012.1	73.0	3002.8	43.6	2996.6	54.0	2996.6	54.0	100.5
Trial-1_0164	100	121812	0.9	9.7158	2.7	4.1301	3.5	0.2910	2.3	0.64	1646.7	32.8	1660.3	28.9	1677.5	50.2	1677.5	50.2	98.2
Trial-1_0165	172	70881	1.8	13.8386	2.8	1.7102	3.6	0.1717	2.2	0.62	1021.2	21.0	1012.4	23.0	993.3	57.1	993.3	57.1	102.8
Trial-1_0166	90	63508	1.7	14.1164	4.0	1.6109	5.0	0.1649	3.1	0.62	984.1	28.6	974.4	31.6	952.8	81.0	952.8	81.0	103.3
Trial-1_0167	513	55323	2.5	16.9059	2.9	0.4512	3.6	0.0553	2.2	0.61	347.1	7.4	378.1	11.4	572.7	62.6	347.1	7.4	NA
Trial-1_0168	165	126824	1.3	9.1038	2.0	4.8235	2.6	0.3185	1.7	0.64	1782.3	25.8	1789.0	21.7	1796.8	35.9	1796.8	35.9	99.2
Trial-1_0169	94	43185	1.9	13.7217	3.7	1.7577	4.7	0.1749	3.0	0.62	1039.2	28.3	1030.0	30.7	1010.5	75.4	1010.5	75.4	102.8
Trial-1_0170	67	63960	1.5	12.8459	4.1	2.1215	5.3	0.1977	3.3	0.63	1162.7	35.1	1155.8	36.4	1143.0	81.8	1143.0	81.8	101.7
Trial-1_0171	41	36456	2.3	13.7541	5.7	1.7235	7.3	0.1719	4.5	0.62	1022.7	42.6	1017.3	46.6	1005.7	115.4	1005.7	115.4	101.7
Trial-1_0172	111	54993	3.1	13.6577	3.5	1.6914	4.4	0.1675	2.8	0.62	998.6	25.6	1005.3	28.4	1020.0	70.5	1020.0	70.5	97.9
Trial-1_0173	186	89873	0.3	13.2883	2.6	1.8995	3.3	0.1831	2.0	0.62	1083.7	20.4	1080.9	21.8	1075.3	51.5	1075.3	51.5	100.8
Trial-1_0174	528	182274	4.3	13.3037	1.6	1.7987	2.0	0.1736	1.2	0.62	1031.7	11.9	1045.0	13.1	1073.0	31.5	1073.0	31.5	96.1
Trial-1_0175	118	103189	1.9	9.1140	2.3	4.8651	3.0	0.3216	2.0	0.64	1797.5	30.7	1796.2	25.6	1794.8	42.5	1794.8	42.5	100.1
Trial-1_0176	314	367966	1.9	9.6000	1.6	3.9745	2.1	0.2767	1.3	0.64	1574.9	18.5	1629.0	16.8	1699.6	29.3	1699.6	29.3	92.7
Trial-1_0177	88	16790	2.0	20.3798	8.0	0.2959	10.0	0.0437	6.0	0.60	275.9	16.1	263.2	23.1	151.1	187.5	275.9	16.1	NA
Trial-1_0178	348	386064	2.1	13.1545	1.9	1.9190	2.4	0.1831	1.5	0.62	1083.8	14.9	1087.7	16.0	1095.6	37.4	1095.6	37.4	98.9
Trial-1_0179	593	138887	1.0	19.6574	3.3	0.2654	4.1	0.0378	2.5	0.60	239.4	5.8	239.0	8.8	235.0	76.1	239.4	5.8	NA
Trial-1_0180	30	14107	1.4	13.5670	6.4	1.8583	8.1	0.1829	5.1	0.62	1082.6	50.3	1066.4	53.6	1033.5	128.5	1033.5	128.5	104.7
Trial-1_0181	173	306108	2.4	9.8528	2.1	3.7662	2.8	0.2691	1.8	0.64	1536.4	24.2	1585.6	22.3	1651.6	39.8	1651.6	39.8	93.0
Trial-1_0182	73	24504	2.9	13.7191	4.2	1.7511	5.4	0.1742	3.3	0.62	1035.4	31.9	1027.6	34.7	1010.9	85.2	1010.9	85.2	102.4
Trial-1_0183	219	53564	3.3	15.9033	3.0	1.0492	3.7	0.1210	2.3	0.61	736.4	16.0	728.5	19.5	704.2	62.9	736.4	16.0	104.6
Trial-1_0184	208	70682	3.0	15.7237	3.1	1.0385	3.9	0.1184	2.4	0.61	721.5	16.3	723.2	20.2	728.3	65.2	721.5	16.3	99.1
Trial-1_0185	32	18017	1.7	13.1518	6.2	1.9157	8.0	0.1827	5.0	0.62	1081.9	49.5	1086.6	53.2	1096.0	124.7	1096.0	124.7	98.7
Trial-1_0186	78	28393	2.1	13.6271	4.0	1.7554	5.2	0.1735	3.2	0.62	1031.3	30.6	1029.2	33.4	1024.5	81.8	1024.5	81.8	100.7
Trial-1_0187	135	374796	1.3	12.5274	3.0	2.0495	3.8	0.1862	2.4	0.63	1100.8	24.2	1132.1	26.1	1192.7	58.9	1192.7	58.9	92.3
Trial-1_0188	48	22165	0.9	9.6286	4.2	3.9580	5.4	0.2764	3.4	0.64	1573.2	47.9	1625.6	43.7	1694.2	76.6	1694.2	76.6	92.9
Trial-1_0189	91	51166	3.7	13.2215	3.7	1.8814	4.8	0.1804	3.0	0.62	1069.2	29.4	1074.6	31.7	1085.4	75.0	1085.4	75.0	98.5
Trial-1_0190	84	46779	1.9	13.3089	3.8	1.9514	4.8	0.1884	3.0	0.62	1112.5	30.9	1098.9	32.6	1072.2	76.2	1072.2	76.2	103.8
Trial-1_0191	81	100461	0.7	18.0598	6.5	0.5491	8.2	0.0719	5.0	0.61	447.7	21.5	444.4	29.5	427.2	145.6	447.7	21.5	104.8
Trial-1_0192	64	213137	0.7	5.3888	2.4	13.2269	3.2	0.5169	2.1	0.66	2686.2	47.1	2696.0	30.5	2703.3	39.8	2703.3	39.8	99.4
Trial-1_0193	161	31558	0.8	19.9778	6.7	0.2407	8.4	0.0349	5.0	0.60	221.0	10.8	219.0	16.5	197.6	155.9	221.0	10.8	NA
Trial-1_0194	18	32632	1.6	13.7120	8.5	1.7413	10.8	0.1732	6.7	0.62	1029.6	63.9	1023.9	69.8	1012.0	171.6	1012.0	171.6	101.7
Trial-1_0195	139	114052	3.2	13.6894	3.1	1.7878	3.9	0.1775	2.4	0.62	1053.3	23.7	1041.0	25.5	1015.3	62.2	1015.3	62.2	103.7
Trial-1_0196	59	17103	0.9	20.5224	12.4	0.1893	15.5	0.0282	9.2	0.60	179.1	16.3	176.0	25.0	134.7	292.4	179.1	16.3	NA
Trial-1_0197	81	249693	1.8	11.4557	3.4	2.9520	4.4	0.2453	2.8	0.63	1414.0	35.0	1395.4	33.1	1366.9	65.1	1366.9	65.1	103.4
Trial-1_0198	82	746465	1.2	5.3752	2.2	12.9517	2.9	0.5049	1.9	0.66	2634.9	41.9	2676.1	27.5	2707.4	36.0	2707.4	36.0	97.3
Trial-1_0199	171	48291	0.8	16.2464	3.6	0.9081	4.6	0.1070	2.8	0.61	655.3	17.5	656.1	22.2	658.6	78.0	655.3	17.5	99.5
Trial-1_0200	22	8012	1.2	19.4078	13.0	0.4774	16.3	0.0672	9.8	0.60	419.2	39.7	396.3	53.5	264.4	300.1	419.2	39.7	158.5
Trial-1_0201	350	759636	1.2	18.2655	3.3	0.4950	4.2	0.0656	2.5	0.60	409.5	10.0	408.3	14.0	401.9	74.2	409.5	10.0	101.9
Trial-1_0202	91	62246	2.0	13.3588	3.8	1.8357	4.9	0.1779	3.0	0.62	1055.2	29.5	1058.3	31.9	1064.7	76.3	1064.7	76.3	99.1
Trial-1_0203	152	14424	0.6	19.7415	6.8	0.2627	8.5	0.0376	5.1	0.60	238.1	11.8	236.9	17.9	225.1	156.8	238.1	11.8	NA
Trial-1_0204	81	121644	1.7	13.5474	4.1	1.7924	5.2	0.1761	3.2	0.62	1045.7	31.2	1042.7	33.9	1036.4	82.1	1036.4	82.1	100.9
Trial-1_0205	94	85799	3.0	11.1988	3.1	3.1272	3.9	0.2540	2.5	0.63	1459.0	32.6	1439.4	30.4	1410.5	58.4	1410.5	58.4	103.4
Trial-1_0206	213	30689	2.8	19.0579	4.5	0.4385	5.6	0.0606	3.4	0.60	379.4	12.4	369.2	17.3	306.0	102.0	379.4	12.4	NA
Trial-1_0207	212	143993	1.2	10.9467	2.1	3.1331	2.7	0.2487	1.7	0.63	1432.0	21.8	1440.8	20.6	1453.9	39.4	1453.9	39.4	98.5
Trial-1_0208	745	432840	10.0	9.6947	1.1	3.6607	1.4	0.2574	0.9	0.64	1476.5	11.7	1562.8	11.1	1681.5	19.7	1681.5	19.7	87.8
Trial-1_0209	40	20037	1.5	11.6752	4.9	2.8151	6.3	0.2384	4.0	0.63	1378.2	49.6	1359.6	47.4	1330.3	95.0	1330.3	95.0	103.6
Trial-1_0210	42	66940	2.3	12.3294	5.2	2.2928	6.6	0.2050	4.2	0.63	1202.3	45.8	1210.1	47.0	1224.0	101.7	1224.0	101.7	98.2
Trial-1_0211	144	156821	2.5	11.6807	2.6	2.7378	3.4	0.2319	2.1	0.63	1344.7	26.0	1338.8	25.2	1329.4	50.9	1329.4	50.9	101.1
Trial-1_0212	103	284964	1.3	10.9833	3.0	3.1676	3.8	0.2523	2.4	0.63	1450.5	31.5	1449.3	29.5	1447.6	56.3	1447.6	56.3	100.2
Trial-1_0213	55	44982	1.1	9.4881	3.6	4.4764	4.7	0.3080	3.0	0.64	1731.0	45.8	1726.6	39.1	1721.2	66.4	1721.2	66.4	100.6
Trial-1_0214	235	83326	1.3	11.9933	2.1	2.5645	2.7	0.2231	1.7	0.63	1298.1	20.0	1290.6	19.8	1278.1	40.9	1278.1	40.9	101.6
Trial-1_0215	174	60965	3.2	12.6269	2.6	2.0980	3.4	0.1921	2.1	0.63	1132.9	22.1	1148.1	23.3	1177.1	52.3	1177.1	52.3	96.2
Trial-1_0216	68	7199	0.6	22.4028	12.0	0.1696	14.9	0.0276	8.8	0.59	175.3	15.2	159.1	21.9	-75.4	294.8	175.3	15.2	NA
Trial-1_0217	95	6863	0.8	20.1437	9.3	0.2022	11.6	0.0295	6.9	0.60	187.7	12.8	187.0	19.9	178.3	218.0	187.7	12.8	NA
Trial-1_0218	108	178946	0.7	5.4922	1.9	12.7633	2.6	0.5084	1.7	0.66	2649.8	36.9	2662.3	24.1	2671.9	31.7	2671.9	31.7	99.2
Trial-1_0219	511	203783	1.6	10.9081	1.4	3.0669	1.8	0.2426	1.1	0.63	1400.4	14.1	1424.5	13.5	1460.6	26.0	1460.6	26.0	95.9
Trial-1_0220	17	15569	0.6	13.1336	8.8	1.9508	11.2	0.1858	7.0	0.62	1098.7	70.9	1098.7	75.6	1098.8	175.9	1098.8	175.9	100.0
Trial-1_0221	169	174068	1.9	13.5609	2.9	1.7526	3.7	0.1724	2.3	0.62	1025.2	21.8	1028.1	23.8	1034.4	58.3	1034.4	58.3	99.1
Trial-1_0222	127	66103	3.6	12.9210	3.1	2.0334	4.0	0.1906	2.5	0.63	1124.4	25.8	1126.8	27.2	1131.4	62.2	1131.4	62.2	99.4
Trial-1_0223	168	197832	2.0	10.9039	2.3	3.1814	2.9	0.2516	1.9	0.64	1446.7	24.2	1452.6	22.6	1461.4	43.0	1461.4	43.0	99.0
Trial-1_0224	898	469478	15.1	13.0915	1.2	1.8410	1.6	0.1748	1.0	0.62	1038.5	9.5	1060.2	10.4	1105.2	24.8	1105.2	24.8	94.0
Trial-1_0225	371	28410	10.2	19.7072	4.0	0.3149	5.0	0.0450	3.0	0.60	283.8	8.3	278.0	12.1	229.1	92.1	283.8	8.3	NA
Trial-1_0226	96	69067	1.2	11.1268	3.1	3.0753	4.0	0.2482	2.5	0.63	1429.1	32.6	1426.6	30.8	1422.8	59.4	1422.8	59.4	100.4
Trial-1_0227	67	24052	0.9	13.5662	4.5	1.8071	5.7	0.1778	3.6	0.62	1055.0	34.6	1048.0	37.4	1033.6	90.4	1033.6	90.4	102.1
Trial-1_0228	112	46271	1.3	19.7175	8.0	0.2569	10.0	0.0367	6.0	0.60	232.5	13.7	232.1	20.8	228.0	185.3	232.5	13.7	NA
Trial-1_0229	148	280630	2.2	13.4714	3.0	1.8373	3.9	0.1795	2.4	0.62	1064.3	23.6	1058.9	25.4	1047.8	60.9	1047.8	60.9	101.6
Trial-1_0230	323	274475	1.9	13.5815	2.1	1.7240	2.7	0.1698	1.7	0.62	1011.1	15.7	1017.5	17.3	1031.3	42.6	1031.3	42.6	98.0
Trial-1_0231	214	75659	1.3	18.3811	4.2	0.5111	5.3	0.0681	3.2	0.60	424.9	13.1	419.2	18.1	387.8	94.3	424.9	13.1	109.6
Trial-1_0232	150	204785	0.6	7.5994	1.9	6.9661	2.5	0.3839	1.7	0.65	2094.7	29.6	2107.1	22.6	2119.3	33.8	2119.3	33.8	98.8
Trial-1_0233	76	22549	1.0	14.0288	4.4	1.6140	5.6	0.1642	3.5	0.62	980.2	31.8	975.7	35.2	965.5	90.0	965.5	90.0	101.5
Trial-1_0234	310	352074	1.0	9.5462	1.6	4.2786	2.0	0.2962	1.3	0.64	1672.6	19.2	1689.3	16.7	1710.0	28.8	1710.0	28.8	97.8
Trial-1_0235	120	594131	0.7	5.3958	1.8	13.1339	2.4	0.5140	1.6	0.66	2673.6	35.2	2689.3	22.8	2701.1	29.9	2701.1	29.9	99.0
Trial-1_0236	77	50582	0.3	10.0316	3.2	3.9385	4.2	0.2866	2.7	0.64	1624.3	38.5	1621.6	34.0	1618.2	60.2	1618.2	60.2	100.4
Trial-1_0237	97	58842	1.7	13.0988	3.7	1.9687	4.7	0.1870	2.9	0.62	1105.3	29.7	1104.9	31.5	1104.1	73.1	1104.1	73.1	100.1
Trial-1_0238	103	65881	1.1	12.7957	3.4	2.1086	4.4	0.1957	2.8	0.63	1152.1	29.0	1151.6	30.3	1150.7	68.2	1150.7	68.2	100.1
Trial-1_0239	52	17889	1.4	11.7161	4.3	2.8024	5.6	0.2381	3.5	0.63	1377.0	43.7	1356.2	41.8	1323.5	84.0	1323.5	84.0	104.0
Trial-1_0240	512	67210	1.6	18.7500	3.1	0.4094	3.8	0.0557	2.3	0.60	349.2	7.8	348.4	11.3	343.0	69.2	349.2	7.8	NA
Trial-1_0241	286	374288	1.1	15.4868	2.6	1.1166	3.3	0.1254	2.1	0.61	761.7	14.8	761.4	17.9	760.4	55.7	761.7	14.8	100.2
Trial-1_0242	110	92540	2.3	11.8782	3.1	2.6070	4.0	0.2246	2.5	0.63	1306.1	29.9	1302.6	29.5	1296.9	60.6	1296.9	60.6	100.7
Trial-1_0243	76	52645	3.5	9.4281	3.0	4.6911	4.0	0.3208	2.5	0.64	1793.5	39.9	1765.6	33.3	1732.8	55.9	1732.8	55.9	103.5
Trial-1_0244	120	41894	1.5	19.5242	7.5	0.2928	9.4	0.0415	5.6	0.60	261.9	14.4	260.8	21.5	250.7	172.4	261.9	14.4	NA
Trial-1_0245	747	444508	1.1	15.5504	1.7	1.0539	2.1	0.1189	1.3	0.61	724.0	9.0	730.8	11.2	751.7	35.8	724.0	9.0	96.3
Trial-1_0246	88	8473	1.7	20.5270	8.7	0.2694	10.9	0.0401	6.5	0.60	253.5	16.2	242.2	23.5	134.2	205.6	253.5	16.2	NA
Trial-1_0247	103	45862	1.6	13.8490	3.7	1.7056	4.8	0.1713	3.0	0.62	1019.3	28.0	1010.6	30.6	991.8	76.2	991.8	76.2	102.8
Trial-1_0248	200	167677	1.2	10.9470	2.2	3.1891	2.8	0.2532	1.8	0.63	1454.9	23.0	1454.5	21.5	1453.9	41.0	1453.9	41.0	100.1
Trial-1_0249	15	69131	2.8	12.8850	9.2	2.0924	11.8	0.1955	7.4	0.62	1151.3	77.7	1146.3	81.3	1136.9	183.9	1136.9	183.9	101.3
Trial-1_0250	101	9744	0.5	21.2115	9.8	0.1839	12.2	0.0283	7.2	0.59	179.9	12.9	171.4	19.2	56.5	234.0	179.9	12.9	NA
Trial-1_0251	34	207983	1.3	13.3910	6.3	1.9299	8.0	0.1874	5.0	0.62	1107.5	51.0	1091.5	53.8	1059.8	126.7	1059.8	126.7	104.5
Trial-1_0252	51	313413	1.1	13.0209	5.0	1.9976	6.5	0.1886	4.0	0.62	1114.1	41.3	1114.7	43.7	1116.0	100.7	1116.0	100.7	99.8
Trial-1_0253	85	613700	2.4	12.0786	3.6	2.4742	4.7	0.2167	2.9	0.63	1264.7	33.8	1264.5	33.8	1264.3	71.1	1264.3	71.1	100.0
Trial-1_0254	184	114037	1.7	8.8267	1.9	5.3343	2.5	0.3415	1.6	0.64	1893.8	26.3	1874.4	21.2	1852.9	34.3	1852.9	34.3	102.2
Trial-1_0255	85	10391	0.8	20.1605	9.1	0.2679	11.3	0.0392	6.8	0.60	247.7	16.5	241.0	24.3	176.4	212.3	247.7	16.5	NA
Trial-1_0256	704	6980	2.8	11.5062	2.8	0.5073	3.6	0.0423	2.2	0.62	267.3	5.9	416.7	12.3	1358.5	54.3	267.3	5.9	NA
Trial-1_0257	235	193652	7.4	13.5550	2.5	1.7872	3.1	0.1757	2.0	0.62	1043.5	18.9	1040.8	20.5	1035.3	49.8	1035.3	49.8	100.8
Trial-1_0258	271	78481	0.9	19.1964	4.3	0.3957	5.3	0.0551	3.2	0.60	345.7	10.8	338.5	15.4	289.5	97.6	345.7	10.8	NA
Trial-1_0259	311	89077	2.4	20.4825	5.3	0.2077	6.7	0.0309	4.0	0.60	195.9	7.7	191.6	11.6	139.3	125.6	195.9	7.7	NA
Trial-1_0260	128	31404	1.0	19.9255	7.1	0.2928	8.8	0.0423	5.3	0.60	267.2	13.8	260.8	20.3	203.6	164.1	267.2	13.8	NA
Trial-1_0261	300	109367	0.5	20.5808	5.7	0.1892	7.1	0.0282	4.2	0.60	179.6	7.5	176.0	11.5	128.1	134.7	179.6	7.5	NA
Trial-1_0262	254	311028	3.7	9.8707	1.8	4.0559	2.3	0.2904	1.5	0.64	1643.3	21.4	1645.5	18.9	1648.2	33.1	1648.2	33.1	99.7
Trial-1_0263	351	161743	2.3	10.9380	1.6	3.1987	2.1	0.2538	1.3	0.63	1457.8	17.5	1456.9	16.4	1455.4	31.1	1455.4	31.1	100.2
Trial-1_0264	225	118336	2.6	19.3890	5.1	0.3313	6.3	0.0466	3.8	0.60	293.5	10.9	290.5	16.0	266.7	116.6	293.5	10.9	NA
Trial-1_0265	35	22169	1.4	11.2037	5.2	3.0957	6.7	0.2516	4.3	0.63	1446.5	55.3	1431.6	51.8	1409.7	100.0	1409.7	100.0	102.6
Trial-1_0266	61	34973	0.7	9.9849	3.5	4.1637	4.6	0.3015	2.9	0.64	1698.9	43.8	1666.9	37.7	1626.9	65.8	1626.9	65.8	104.4
Trial-1_0267	180	5578	2.0	21.2901	10.2	0.0959	12.7	0.0148	7.5	0.59	94.7	7.1	92.9	11.3	47.7	244.5	94.7	7.1	NA
Trial-1_0268	41	41850	2.0	13.6129	6.0	1.7269	7.6	0.1705	4.8	0.62	1014.8	44.6	1018.6	49.2	1026.6	121.2	1026.6	121.2	98.8
Trial-1_0269	686	748441	2.5	12.0527	1.3	2.3519	1.7	0.2056	1.1	0.63	1205.3	11.7	1228.1	12.0	1268.5	25.7	1268.5	25.7	95.0

Trial-1_0270	60	230463	0.8	3.2334	2.1	30.9214	2.9	0.7251	2.0	0.68	3515.2	53.2	3516.6	28.4	3517.3	32.7	3517.3	32.7	99.9
Trial-1_0271	172	212457	1.1	9.6527	2.1	4.2362	2.8	0.2966	1.8	0.64	1674.3	26.0	1681.1	22.7	1689.5	39.2	1689.5	39.2	99.1
Trial-1_0272	41	18601	2.3	13.0471	5.5	2.0960	7.0	0.1983	4.4	0.62	1166.4	46.8	1147.5	48.3	1112.0	109.5	1112.0	109.5	104.9
Trial-1_0273	84	59656	0.8	13.4853	4.1	1.7757	5.3	0.1737	3.3	0.62	1032.3	31.3	1036.6	34.2	1045.7	83.2	1045.7	83.2	98.7
Trial-1_0274	100	97357	1.0	9.3856	2.7	4.5113	3.6	0.3071	2.3	0.64	1726.4	34.6	1733.1	29.6	1741.1	50.2	1741.1	50.2	99.2
Trial-1_0275	112	115344	0.9	10.9974	2.9	3.1711	3.7	0.2529	2.4	0.63	1453.5	30.7	1450.1	28.8	1445.1	55.0	1445.1	55.0	100.6
Trial-1_0276	255	673301	1.5	10.9386	1.9	3.1258	2.5	0.2480	1.6	0.63	1428.1	20.2	1439.1	19.2	1455.3	36.7	1455.3	36.7	98.1
Trial-1_0277	249	983570	2.1	11.9919	2.1	2.4891	2.7	0.2165	1.7	0.63	1263.3	19.5	1268.9	19.6	1278.3	41.0	1278.3	41.0	98.8
Trial-1_0278	114	42469	0.7	12.7471	3.3	2.1634	4.2	0.2000	2.6	0.63	1175.4	28.3	1169.4	29.2	1158.3	65.1	1158.3	65.1	101.5
Trial-1_0279	121	27078	1.0	15.8929	4.2	1.0414	5.3	0.1200	3.3	0.61	730.8	22.6	724.6	27.6	705.6	89.7	730.8	22.6	103.6
Trial-1_0280	156	1058383	1.2	5.3215	1.6	13.0602	2.2	0.5041	1.4	0.66	2631.2	31.0	2684.0	20.4	2724.0	26.7	2724.0	26.7	96.6
Trial-1_0281	61	43985	1.3	11.3650	4.0	2.9101	5.2	0.2399	3.3	0.63	1386.0	41.0	1384.5	39.4	1382.2	77.6	1382.2	77.6	100.3
Trial-1_0282	120	58367	4.5	13.4607	3.4	1.8351	4.3	0.1792	2.7	0.62	1062.4	26.5	1058.1	28.6	1049.4	68.7	1049.4	68.7	101.2
Trial-1_0283	62	18661	2.2	13.5379	4.8	1.8516	6.1	0.1818	3.8	0.62	1076.8	37.5	1064.0	40.0	1037.8	96.0	1037.8	96.0	103.8
Trial-1_0284	72	319023	2.8	9.0652	3.1	4.8567	4.1	0.3193	2.6	0.64	1786.4	41.1	1794.8	34.6	1804.5	57.2	1804.5	57.2	99.0
Trial-1_0285	419	154769	1.5	13.6921	1.9	1.7060	2.4	0.1694	1.5	0.62	1008.9	13.9	1010.8	15.4	1014.9	38.1	1014.9	38.1	99.4
Trial-1_0286	97	27717	0.9	17.9456	6.2	0.5664	7.7	0.0737	4.7	0.60	458.6	20.7	455.7	28.4	441.4	137.3	458.6	20.7	103.9
Trial-1_0287	14	16947	2.2	5.2447	5.3	14.4885	7.0	0.5511	4.7	0.66	2829.8	107.2	2782.3	66.9	2747.9	86.5	2747.9	86.5	103.0
Trial-1_0288	204	164535	2.3	13.5405	2.6	1.7882	3.4	0.1756	2.1	0.62	1042.9	20.2	1041.2	22.0	1037.4	53.4	1037.4	53.4	100.5
Trial-1_0289	54	109165	0.7	10.8062	4.3	3.0493	5.6	0.2390	3.5	0.63	1381.4	43.9	1420.1	42.7	1478.5	81.9	1478.5	81.9	93.4
Trial-1_0290	60	60118	1.6	9.3595	3.5	4.6395	4.6	0.3149	2.9	0.64	1764.9	45.5	1756.4	38.4	1746.2	64.7	1746.2	64.7	101.1
Trial-1_0291	58	534756	1.2	10.8848	4.0	3.2921	5.2	0.2599	3.3	0.63	1489.3	43.5	1479.2	40.2	1464.7	75.9	1464.7	75.9	101.7
Trial-1_0292	93	36959	2.4	13.5333	3.9	1.7805	5.0	0.1748	3.1	0.62	1038.3	29.8	1038.4	32.5	1038.5	79.1	1038.5	79.1	100.0
Trial-1_0293	287	19097	1.5	19.1530	5.2	0.2579	6.4	0.0358	3.9	0.60	226.9	8.6	233.0	13.4	294.7	117.7	226.9	8.6	NA
Trial-1_0294	122	313054	0.9	5.1830	1.8	14.2223	2.4	0.5346	1.6	0.67	2760.9	35.8	2764.6	22.8	2767.4	29.4	2767.4	29.4	99.8
Trial-1_0295	113	214536	6.1	13.7275	3.6	1.7611	4.6	0.1753	2.8	0.62	1041.4	27.2	1031.2	29.5	1009.7	72.4	1009.7	72.4	103.1
Trial-1_0296	60	59432	1.4	9.0994	3.4	4.9369	4.5	0.3258	2.9	0.64	1818.0	45.6	1808.6	37.9	1797.7	62.6	1797.7	62.6	101.1
Trial-1_0297	469	134593	2.9	18.8337	3.2	0.4038	4.1	0.0552	2.4	0.60	346.1	8.2	344.4	11.9	332.9	73.5	346.1	8.2	NA
Trial-1_0298	1576	47109	4.8	9.3731	1.4	1.1393	1.8	0.0774	1.1	0.63	480.9	5.2	772.2	9.6	1743.6	25.2	1743.6	25.2	27.6
Trial-1_0299	178	85299	1.5	11.5151	2.4	2.8240	3.1	0.2358	2.0	0.63	1365.1	24.1	1361.9	23.3	1357.0	46.5	1357.0	46.5	100.6
Trial-1_0300	60	193140	1.6	13.4220	4.8	1.8766	6.2	0.1827	3.8	0.62	1081.6	38.1	1072.9	40.8	1055.2	97.1	1055.2	97.1	102.5
Trial-1_0301	126	297086	2.4	13.3851	3.3	1.8632	4.2	0.1809	2.6	0.62	1071.8	26.1	1068.1	28.1	1060.7	66.9	1060.7	66.9	101.0
Trial-1_0302	175	169856	4.0	13.4451	2.8	1.8238	3.6	0.1778	2.3	0.62	1055.2	22.0	1054.1	23.9	1051.7	57.4	1051.7	57.4	100.3
Trial-1_0303	233	111213	1.2	13.1860	2.4	1.8986	3.1	0.1816	1.9	0.62	1075.5	19.2	1080.6	20.7	1090.8	48.8	1090.8	48.8	98.6
Trial-1_0304	118	723402	1.5	5.4317	1.9	13.0780	2.5	0.5152	1.6	0.66	2678.8	36.2	2685.3	23.5	2690.2	30.8	2690.2	30.8	99.6
Trial-1_0305	108	81327	1.8	12.9329	3.4	2.1155	4.4	0.1984	2.7	0.62	1166.9	29.2	1153.9	30.2	1129.5	68.1	1129.5	68.1	103.3
Trial-1_0306	255	939039	1.2	15.5964	2.9	1.0604	3.7	0.1199	2.3	0.61	730.2	15.7	734.0	19.4	745.5	61.9	730.2	15.7	98.0
Trial-1_0307	297	284517	0.9	13.1585	2.2	1.8699	2.8	0.1785	1.7	0.62	1058.5	16.9	1070.5	18.4	1095.0	43.5	1095.0	43.5	96.7
Trial-1_0308	60	52178	1.3	9.9241	3.7	3.9817	4.8	0.2866	3.1	0.64	1624.5	44.2	1630.5	39.2	1638.2	69.0	1638.2	69.0	99.2
Trial-1_0309	86	8069	1.8	20.4040	9.1	0.2630	11.4	0.0389	6.8	0.60	246.1	16.4	237.1	24.1	148.3	214.5	246.1	16.4	NA
Trial-1_0310	87	18713	0.7	17.9100	6.5	0.5687	8.2	0.0739	4.9	0.60	459.4	21.9	457.2	30.0	445.8	144.6	459.4	21.9	103.1
Trial-1_0311	138	134240	1.7	13.0050	3.1	1.9717	4.0	0.1860	2.5	0.62	1099.5	25.2	1105.9	26.9	1118.4	62.3	1118.4	62.3	98.3
Trial-1_0312	136	98491	1.0	13.0911	3.2	1.9153	4.1	0.1819	2.5	0.62	1077.1	25.2	1086.4	27.1	1105.3	63.6	1105.3	63.6	97.4
Trial-1_0313	103	92623	1.6	9.3984	2.7	4.4862	3.6	0.3058	2.3	0.64	1720.0	34.4	1728.4	29.5	1738.6	50.1	1738.6	50.1	98.9
Trial-1_0314	28	15914	1.6	5.9334	4.0	11.3845	5.3	0.4899	3.5	0.66	2570.3	74.5	2555.2	49.8	2543.2	67.1	2543.2	67.1	101.1
Trial-1_0315	184	11539	2.9	19.7399	6.0	0.3040	7.4	0.0435	4.4	0.59	274.7	11.8	269.6	17.5	225.3	137.9	274.7	11.8	NA
Trial-1_0316	448	1342188	5.0	10.9419	1.5	3.0872	1.9	0.2450	1.2	0.63	1412.6	15.6	1429.5	14.8	1454.8	28.4	1454.8	28.4	97.1
Trial-1_0317	105	269743	5.0	8.9692	2.6	4.8654	3.4	0.3165	2.2	0.64	1772.6	34.3	1796.3	29.0	1823.9	47.9	1823.9	47.9	97.2
Trial-1_0318	112	14507	1.2	19.5309	7.8	0.2837	9.8	0.0402	5.9	0.60	254.0	14.6	253.6	22.0	249.9	180.6	254.0	14.6	NA
Trial-1_0319	110	106791	2.1	13.3584	3.6	1.8139	4.6	0.1757	2.9	0.62	1043.6	27.6	1050.5	30.2	1064.7	72.6	1064.7	72.6	98.0
Trial-1_0320	764	58090	1.2	18.2870	3.0	0.3006	3.7	0.0399	2.3	0.60	252.0	5.6	266.9	8.8	399.3	67.1	252.0	5.6	NA
Trial-1_0321	83	698250	1.6	9.7684	3.2	3.9148	4.2	0.2773	2.7	0.64	1578.0	37.4	1616.7	34.0	1667.5	60.0	1667.5	60.0	94.6
Trial-1_0322	77	74298	1.4	13.0836	4.2	1.9621	5.4	0.1862	3.4	0.62	1100.7	33.9	1102.6	36.2	1106.4	84.1	1106.4	84.1	99.5
Trial-1_0323	51	65408	1.0	5.3592	2.8	13.5389	3.8	0.5262	2.5	0.66	2725.6	56.0	2718.0	35.9	2712.4	46.8	2712.4	46.8	100.5
Trial-1_0324	234	35565	1.1	20.0061	6.3	0.2115	7.8	0.0307	4.6	0.60	194.8	8.9	194.8	13.8	194.3	145.5	194.8	8.9	NA
Trial-1_0325	119	111310	1.1	13.6302	3.6	1.7216	4.5	0.1702	2.8	0.62	1013.2	26.5	1016.6	29.2	1024.1	72.1	1024.1	72.1	98.9
Trial-1_0326	30	138798	0.9	5.4082	3.8	12.7703	5.0	0.5009	3.3	0.66	2617.7	71.9	2662.9	47.5	2697.3	62.3	2697.3	62.3	97.0
Trial-1_0327	221	36715	1.0	19.6329	5.7	0.2719	7.1	0.0387	4.3	0.60	244.9	10.2	244.2	15.5	237.9	132.0	244.9	10.2	NA
Trial-1_0328	131	423532	0.9	10.8858	2.8	2.8981	3.7	0.2288	2.3	0.63	1328.3	27.7	1381.4	27.6	1464.5	53.7	1464.5	53.7	90.7
Trial-1_0329	235	89317	1.4	13.7615	2.5	1.7079	3.2	0.1705	2.0	0.62	1014.6	18.7	1011.5	20.5	1004.7	51.0	1004.7	51.0	101.0
Trial-1_0330	198	34901	1.6	16.7283	3.7	0.8383	4.6	0.1017	2.8	0.61	624.4	16.7	618.2	21.3	595.6	79.2	624.4	16.7	104.8
Trial-1_0331	35	35803	1.1	13.4078	6.3	1.8768	8.1	0.1825	5.0	0.62	1080.6	49.9	1072.9	53.4	1057.3	127.1	1057.3	127.1	102.2
Trial-1_0332	32	48055	2.8	9.9502	5.0	4.1010	6.5	0.2960	4.1	0.64	1671.2	61.0	1654.5	53.1	1633.3	92.9	1633.3	92.9	102.3
Trial-1_0333	53	54837	1.7	9.6178	3.8	4.4301	4.9	0.3090	3.1	0.64	1735.9	47.8	1718.0	40.8	1696.2	69.7	1696.2	69.7	102.3
Trial-1_0334	132	81620	0.9	18.4894	5.7	0.4821	7.1	0.0647	4.3	0.60	403.9	16.7	399.5	23.4	374.6	127.5	403.9	16.7	107.8
Trial-1_0335	71	15114	1.4	19.9086	9.9	0.2785	12.3	0.0402	7.4	0.60	254.1	18.4	249.4	27.2	205.6	229.3	254.1	18.4	NA
Trial-1_0336	209	23127	1.0	18.2127	4.2	0.5408	5.3	0.0714	3.2	0.60	444.8	13.9	438.9	19.0	408.4	95.1	444.8	13.9	108.9
Trial-1_0337	224	227736	2.0	13.0762	2.4	1.9809	3.1	0.1879	1.9	0.62	1109.8	19.8	1109.0	21.0	1107.5	48.7	1107.5	48.7	100.2
Trial-1_0338	212	264987	3.0	11.5745	2.2	2.8119	2.8	0.2360	1.8	0.63	1366.1	22.0	1358.7	21.3	1347.1	42.5	1347.1	42.5	101.4
Trial-1_0339	106	47115	2.9	12.9670	3.5	2.0758	4.4	0.1952	2.8	0.62	1149.6	29.2	1140.9	30.5	1124.3	69.3	1124.3	69.3	102.3
Trial-1_0340	106	19363	1.1	13.6000	3.7	1.7681	4.7	0.1744	2.9	0.62	1036.3	27.9	1033.8	30.4	1028.6	74.3	1028.6	74.3	100.8
Trial-1_0341	192	122502	1.7	12.7262	2.6	2.1329	3.3	0.1969	2.1	0.63	1158.5	21.8	1159.5	22.7	1161.5	50.8	1161.5	50.8	99.7
Trial-1_0342	67	39394	1.1	12.8846	4.4	2.0246	5.6	0.1892	3.5	0.62	1117.0	36.0	1123.8	38.2	1137.0	87.3	1137.0	87.3	98.2
Trial-1_0343	79	73348	1.2	9.0808	3.0	4.9518	3.9	0.3261	2.5	0.64	1819.6	39.6	1811.1	32.8	1801.4	54.1	1801.4	54.1	101.0
Trial-1_0344	53	205742	0.8	11.9361	4.5	2.5658	5.8	0.2221	3.7	0.63	1293.0	43.0	1290.9	42.7	1287.4	88.4	1287.4	88.4	100.4
Trial-1_0345	357	241448	2.4	9.0997	1.4	4.7114	1.9	0.3109	1.2	0.64	1745.3	18.4	1769.3	15.7	1797.6	26.2	1797.6	26.2	97.1
Trial-1_0346	209	1794464	8.2	5.5007	1.5	11.8189	1.9	0.4715	1.3	0.66	2490.2	26.6	2590.2	18.2	2669.3	24.2	2669.3	24.2	93.3
Trial-1_0347	107	148211	2.5	9.8101	2.7	4.1089	3.5	0.2923	2.3	0.64	1653.2	33.0	1656.1	28.9	1659.6	50.5	1659.6	50.5	99.6
Trial-1_0348	46	21419	2.5	12.8745	5.1	2.1799	6.5	0.2035	4.1	0.63	1194.3	44.6	1174.6	45.6	1138.5	101.6	1138.5	101.6	104.9
Trial-1_0349	274	80476	1.4	19.3232	4.9	0.2920	6.1	0.0409	3.7	0.60	258.5	9.3	260.1	14.1	274.4	112.7	258.5	9.3	NA
Trial-1_0350	235	47720	0.9	18.0775	4.1	0.5234	5.1	0.0686	3.1	0.60	427.8	12.8	427.4	17.8	425.1	90.9	427.8	12.8	100.7
Trial-1_0351	13	14672	1.2	5.8190	5.7	11.8777	7.6	0.5013	5.1	0.66	2619.3	108.8	2594.8	71.7	2575.8	96.0	2575.8	96.0	101.7
Trial-1_0352	67	27903	1.4	13.4228	4.6	1.8412	5.9	0.1792	3.6	0.62	1062.8	35.7	1060.3	38.6	1055.1	92.5	1055.1	92.5	100.7
Trial-1_0353	93	35199	1.1	16.0490	5.4	0.8192	6.9	0.0954	4.2	0.61	587.1	23.5	607.6	31.4	684.8	115.9	587.1	23.5	85.7
Trial-1_0354	51	102754	1.3	9.5963	3.9	4.3003	5.0	0.2993	3.2	0.64	1687.8	47.9	1693.4	41.5	1700.3	71.4	1700.3	71.4	99.3
Trial-1_0355	129	266324	1.7	13.5448	3.3	1.7689	4.2	0.1738	2.6	0.62	1032.9	25.2	1034.1	27.6	1036.8	67.2	1036.8	67.2	99.6
Trial-1_0356	131	63889	2.3	13.4650	3.3	1.7803	4.2	0.1739	2.6	0.62	1033.3	25.1	1038.3	27.4	1048.7	66.6	1048.7	66.6	98.5
Trial-1_0357	63	153290	0.9	17.7113	7.4	0.6045	9.3	0.0776	5.6	0.61	482.1	26.2	480.1	35.6	470.6	164.2	482.1	26.2	102.4
Trial-1_0358	245	35417	1.8	20.5413	8.5	0.1051	10.5	0.0157	6.2	0.59	100.2	6.2	101.5	10.2	132.5	199.4	100.2	6.2	NA
Trial-1_0359	48	12301	1.4	13.8236	5.4	1.7498	6.9	0.1754	4.3	0.62	1042.0	41.6	1027.1	44.9	995.5	110.6	995.5	110.6	104.7
Trial-1_0360	184	296788	0.7	9.0387	2.0	4.8983	2.6	0.3211	1.7	0.64	1795.2	25.9	1802.0	21.7	1809.9	35.8	1809.9	35.8	99.2
Trial-1_0361	630	245957	0.7	18.8034	2.8	0.3939	3.5	0.0537	2.1	0.60	337.3	7.0	337.2	10.1	336.6	64.0	337.3	7.0	NA
Trial-1_0362	32	14430	1.2	13.2121	6.1	2.0080	7.9	0.1924	4.9	0.62	1134.4	51.1	1118.2	53.3	1086.8	123.2	1086.8	123.2	104.4
Trial-1_0363	36	52678	0.8	5.4008	3.2	13.2653	4.3	0.5196	2.9	0.66	2697.5	62.9	2698.7	40.6	2699.6	53.1	2699.6	53.1	99.9
Trial-1_0364	84	115029	1.5	8.0602	2.6	6.2544	3.4	0.3656	2.2	0.65	2008.8	38.3	2012.1	30.0	2015.5	46.3	2015.5	46.3	99.7
Trial-1_0365	48	54312	0.4	5.3529	2.8	13.4230	3.7	0.5211	2.5	0.66	2703.9	54.6	2709.9	35.2	2714.3	45.9	2714.3	45.9	99.6
Trial-1_0366	24	31678	3.1	13.1641	7.0	2.0200	9.0	0.1929	5.6	0.62	1136.9	58.4	1122.3	61.1	1094.1	140.8	1094.1	140.8	103.9
Trial-1_0367	61	109787	0.7	5.2193	2.4	14.1952	3.3	0.5373	2.2	0.67	2772.3	48.9	2762.8	30.9	2755.9	40.0	2755.9	40.0	100.6
Trial-1_0368	25	36241	0.9	13.2555	7.2	1.8838	9.2	0.1811	5.7	0.62	1073.0	56.4	1075.4	60.8	1080.3	143.7	1080.3	143.7	99.3
Trial-1_0369	55	62168	2.1	13.4935	5.0	1.7586	6.4	0.1721	4.0	0.62	1023.7	37.5	1030.3	41.3	1044.4	100.8	1044.4	100.8	98.0
Trial-1_0370	68	52714	1.6	9.1132	3.1	4.8800	4.1	0.3225	2.6	0.64	1802.1	41.4	1798.8	34.5	1794.9	57.2	1794.9	57.2	100.4
Trial-1_0371	201	45954	1.5	13.7920	2.6	1.7291	3.3	0.1730	2.1	0.62	1028.4	19.6	1019.4	21.3	1000.2	52.7	1000.2	52.7	102.8
Trial-1_0372	46	64998	1.2	5.8074	3.0	11.6626	3.9	0.4912	2.6	0.66	2576.0	55.3	2577.7	36.9	2579.1	49.4	2579.1	49.4	99.9
Trial-1_0373	110	161463	2.2	11.0102	2.9	3.0736	3.7	0.2454	2.3	0.63	1414.9	29.8	1426.1	28.4	1442.9	54.7	1442.9	54.7	98.1
Trial-1_0374	28	49110	1.2	8.7108	4.7	5.3195	6.2	0.3361	4.0	0.64	1867.7	64.9	1872.0	53.1	1876.7	85.6	1876.7	85.6	99.5
Trial-1_0375	144	249886	1.5	13.1129	3.0	1.9739	3.8	0.1877	2.4	0.62	1109.0	24.0	1106.6	25.5	1101.9	59.0	1101.9	59.0	100.6
Trial-1_0376	186	33444	2.0	19.8073	6.1	0.2587	7.6	0.0372	4.6	0.60	235.2	10.5	233.6	15.9	217.4	141.7	235.2	10.5	NA
Trial-1_0377	152	272968	2.2	18.8570	5.2	0.4495	6.5	0.0615	3.9	0.60	384.6	14.7	377.0	20.6	330.1	118.7	384.6	14.7	NA
Trial-1_0378	105	11995	1.7	18.3715	5.8	0.5496	7.2	0.0732	4.4	0.60	455.6	19.2	444.7	26.0	389.0	129.3	455.6	19.2	117.1
Trial-1_0379	12	14934	1.2	12.3666	9.4	2.3339	12.1	0.2093	7.6	0.63	1225.2	84.5	1222.7	86.0	1218.1	185.1	1218.1	185.1	100.6
Trial-1_0380	38	98593	0.6	8.9347	4.1	5.1572	5.4	0.3342	3.5	0.64	1858.6	56.1	1845.6	45.9	1830.9	74.9	1830.9	74.9	101.5
Trial-1_0381	336	16350	1.0	22.1839	7.3	0.0929	9.1	0.0149	5.3	0.59	95.6	5.1	90.2	7.8	-51.4	178.3	95.6	5.1	NA
Trial-1_0382	156	92081	1.5	17.3437	4.3	0.6894	5.4	0.0867	3.3	0.61	536.1	16.9	532.4	22.5	516.8	94.7	536.1	16.9	103.7
Trial-1_0383	67	193522	0.6	4.9065	2.3	15.5842	3.1	0.5546	2.1	0.67	2844.2	47.4	2851.7	29.5	2856.9	37.6	2856.9	37.6	99.6
Trial-1_0384	42	18855	0.7	10.8104	4.5	3.1417	5.8	0.2463	3.7	0.63	1419.5	46.9	1443.0	44.7	1477.7	85.0	1477.7	85.0	96.1
Trial-1_0385	28	47724	1.1	5.3448	3.7	13.8701	4.9	0.5377	3.3	0.66	2773.7	73.3	2740.9	46.4	2716.8	60.3	2716.8	60.3	102.1
Trial-1_0386	124	20062	2.2	19.2045	6.0	0.4127	7.5	0.0575	4.5	0.60	360.3	15.9	350.8	22.4	288.5	137.8	360.3	15.9	NA
Trial-1_0387	118	326768	1.1	5.2300	1.8	14.0200	2.4	0.5318	1.6	0.66	2749.0	35.3	2751.1	22.5	2752.5	29.2	2752.5	29.2	99.9
Trial-1_0388	166	33310	0.9	18.8926	5.1	0.4335	6.4	0.0594	3.9	0.60	372.0	14.0	365.7	19.7	325.8	116.4	372.0	14.0	NA
Trial-1_0389	73	50555	1.0	10.0225	3.3	3.9258	4.2	0.2854	2.7	0.64	1618.3	38.8	1619.0	34.4	1619.9	60.9	1619.9	60.9	99.9
Trial-1_0390	524	121642	12.5	19.4401	3.4	0.3141	4.2	0.0443	2.5	0.60	279.4	6.9	277.4	10.2	260.6	77.2	279.4	6.9	NA
Trial-1_0391	93	19380	1.6	18.3279	6.0	0.5734	7.5	0.0762	4.6	0.60	473.5	20.8	460.2	27.9	394.3	135.0	473.5	20.8	120.1
Trial-1_0392	388	465074	1.0	13.2812	1.8	1.8741	2.4	0.1805	1.5	0.62	1069.8	14.5	1072.0	15.6	1076.4	37.0	1076.4	37.0	99.4
Trial-1_0393	78	237261	1.6	5.4013	2.2	13.3578	3.0	0.5233	2.0	0.66	2713.1	43.4	2705.3	27.9	2699.4	36.5	2699.4	36.5	100.5
Trial-1_0394	81	41299	1.1	13.0904	3.9	2.0160	5.0	0.1914	3.1	0.62	1128.9	32.5	1120.9	34.1	1105.4	78.6	1105.4	78.6	102.1
Trial-1_0395	236	468435	0.8	9.6439	1.8	3.9945	2.4	0.2794	1.5	0.64	1588.3	21.5	1633.1	19.4	1691.2	33.9	1691.2	33.9	93.9
Trial-1_0396	30	100169	1.3	5.2276	3.5	14.0367	4.7	0.5322	3.1	0.67	2750.7	69.9	2752.2	44.5	2753.3	57.6	2753.3	57.6	99.9
Trial-1_0397	53	67727	1.5	12.3151	4.6	2.3863	5.8	0.2131	3.7	0.63	1245.5	41.5	1238.5	41.8	1226.3	89.4	1226.3	89.4	101.6
Trial-1_0398	40	13753	1.0	12.6638	5.3	2.2764	6.8	0.2091	4.3	0.63	1223.9	47.8	1205.0	48.3	1171.3	105.5	1171.3	105.5	104.5
Trial-1_0399	32	19259	1.2	13.2378	6.3	1.9722	8.1	0.1894	5.0	0.62	1117.9	51.7	1106.1	54.5	1082.9	127.0	1082.9	127.0	103.2
Trial-1_0400	74	74269	2.0	13.4683	4.2	1.8861	5.4	0.1842	3.4	0.62	1090.1	33.7	1076.2	35.8	1048.3	85.1	1048.3	85.1	104.0
Trial-1_0401	59	158053	0.2	3.8626	2.2	23.2660	3.0	0.6518	2.0	0.67	3235.1	51.7	3238.1	29.3	3240.0	35.0	3240.0	35.0	99.9
Trial-1_0402	77	11507	1.6	19.0390	7.1	0.4797	8.9	0.0662	5.4	0.60	413.4	21.6	397.9	29.4	308.3	162.7	413.4	21.6	134.1
Trial-1_0403	52	29900	2.0	13.2226	5.0	1.9121	6.4	0.1834	4.0	0.62	1085.3	39.9	1085.3	42.8	1085.2	100.5	1085.2	100.5	100.0
Trial-1_0404	67	55048	0.9	11.1556	3.7	3.1349	4.8	0.2536	3.0	0.63	1457.2	39.6	1441.3	36.9	1417.9	71.0	1417.9	71.0	102.8
Trial-1_0405	87	54569	2.6	12.7789	3.6	2.2034	4.7	0.2042	2.9	0.63	1197.9	31.9	1182.1	32.6	1153.3	72.2	1153.3	72.2	103.9
Trial-1_0406	26	8012	3.0	13.6304	7.2	1.7914	9.2	0.1771	5.8	0.62	1051.1	55.9	1042.3	60.3	1024.0	146.6	1024.0	146.6	102.6
Trial-1_0407	145	34276	1.4	17.9979	5.1	0.5183	6.4	0.0677	3.9	0.60	422.0	15.9	424.0	22.3	434.9	114.4	422.0	15.9	97.0

Trial-1_0408	9	43380	1.9	9.8600	9.4	3.9255	12.2	0.2807	7.8	0.64	1595.0	109.8	1618.9	98.8	1650.3	173.9	1650.3	173.9	96.6
Trial-1_0409	185	430345	2.1	13.2625	2.6	1.9749	3.3	0.1900	2.1	0.62	1121.2	21.4	1107.0	22.5	1079.2	52.4	1079.2	52.4	103.9
Trial-1_0410	171	163174	4.1	13.1125	2.7	2.0159	3.5	0.1917	2.2	0.62	1130.7	22.4	1120.9	23.5	1102.0	54.0	1102.0	54.0	102.6
Trial-1_0411	106	74542	1.3	12.3110	3.2	2.4352	4.1	0.2174	2.6	0.63	1268.3	29.7	1253.1	29.6	1227.0	62.9	1227.0	62.9	103.4
Trial-1_0412	77	90134	2.2	13.0813	4.0	2.0365	5.1	0.1932	3.2	0.62	1138.7	33.5	1127.8	35.0	1106.8	80.3	1106.8	80.3	102.9
Trial-1_0413	131	228870	1.3	13.1320	3.1	1.9860	4.0	0.1892	2.5	0.62	1116.8	25.5	1110.8	26.9	1099.0	62.4	1099.0	62.4	101.6
Trial-1_0414	112	25594	1.6	17.0670	5.0	0.7325	6.3	0.0907	3.9	0.61	559.5	20.6	558.0	27.2	552.0	109.8	559.5	20.6	101.4
Trial-1_0415	54	82049	0.6	10.7505	4.0	3.2282	5.2	0.2517	3.3	0.63	1447.3	42.6	1464.0	40.1	1488.3	75.8	1488.3	75.8	97.2
Trial-1_0416	142	101592	4.0	13.8479	3.2	1.6705	4.1	0.1678	2.5	0.62	999.8	23.4	997.4	25.9	992.0	64.9	992.0	64.9	100.8
Trial-1_0417	127	12876	0.3	19.5983	6.2	0.3771	7.8	0.0536	4.6	0.60	336.6	15.2	324.9	21.6	241.9	143.3	336.6	15.2	NA
Trial-1_0418	164	22356	1.3	18.8302	5.4	0.4008	6.7	0.0547	4.0	0.60	343.5	13.5	342.2	19.6	333.3	122.2	343.5	13.5	NA
Trial-1_0419	157	16199	1.4	19.7720	6.7	0.2778	8.4	0.0398	5.0	0.60	251.8	12.3	248.9	18.5	221.6	155.4	251.8	12.3	NA
Trial-1_0420	90	15553	0.8	18.7503	6.4	0.5206	8.0	0.0708	4.8	0.60	440.9	20.6	425.5	27.9	343.0	145.2	440.9	20.6	128.6
Trial-1_0421	80	111525	1.0	8.7452	2.8	5.4945	3.7	0.3485	2.4	0.64	1927.4	39.5	1899.7	31.6	1869.6	50.7	1869.6	50.7	103.1
Trial-1_0422	72	144306	1.2	12.6920	4.0	2.2089	5.2	0.2033	3.2	0.63	1193.2	35.2	1183.9	36.1	1166.8	79.8	1166.8	79.8	102.3
Trial-1_0423	66	146012	2.1	12.8535	4.3	2.1017	5.5	0.1959	3.4	0.62	1153.4	36.4	1149.4	38.0	1141.8	85.9	1141.8	85.9	101.0
Trial-1_0424	24	11044	0.5	12.7390	7.2	2.1205	9.2	0.1959	5.8	0.62	1153.4	60.9	1155.5	63.7	1159.5	143.0	1159.5	143.0	99.5
Trial-1_0425	110	15771	1.2	19.3294	7.7	0.2918	9.6	0.0409	5.7	0.60	258.4	14.5	260.0	22.0	273.7	176.2	258.4	14.5	NA
Trial-1_0426	19	20415	1.3	4.8958	4.5	15.4593	6.0	0.5489	4.0	0.66	2820.7	91.1	2844.0	57.3	2860.5	73.0	2860.5	73.0	98.6
Trial-1_0427	91	94106	0.5	8.7936	2.7	5.3024	3.5	0.3382	2.3	0.64	1877.9	36.7	1869.3	29.9	1859.7	48.4	1859.7	48.4	101.0
Trial-1_0428	144	625958	1.9	12.3471	2.8	2.3603	3.6	0.2114	2.2	0.63	1236.1	25.3	1230.7	25.5	1221.2	54.8	1221.2	54.8	101.2
Trial-1_0429	155	74633	1.5	12.7980	2.8	2.1729	3.5	0.2017	2.2	0.63	1184.4	23.9	1172.4	24.6	1150.4	54.8	1150.4	54.8	103.0
Trial-1_0430	175	19942	2.2	18.3717	4.6	0.5369	5.7	0.0715	3.5	0.60	445.4	14.9	436.4	20.4	388.9	102.9	445.4	14.9	114.5
Trial-1_0431	588	381437	5.9	13.3166	1.5	1.8210	2.0	0.1759	1.2	0.62	1044.4	11.7	1053.1	12.8	1071.0	30.7	1071.0	30.7	97.5
Trial-1_0432	53	27854	1.9	13.0475	4.9	1.9715	6.3	0.1866	3.9	0.62	1102.7	39.9	1105.8	42.6	1111.9	98.7	1111.9	98.7	99.2
Trial-1_0433	295	182999	1.5	13.3199	2.1	1.8811	2.7	0.1817	1.7	0.62	1076.4	16.7	1074.4	17.9	1070.5	42.3	1070.5	42.3	100.5
Trial-1_0434	122	80941	1.9	9.9597	2.5	4.0854	3.3	0.2951	2.1	0.64	1667.0	30.5	1651.4	26.5	1631.6	46.5	1631.6	46.5	102.2
Trial-1_0435	130	83538	0.9	13.5215	3.2	1.8356	4.1	0.1800	2.6	0.62	1067.1	25.1	1058.3	26.9	1040.3	64.8	1040.3	64.8	102.6
Trial-1_0436	82	931084	2.6	9.2891	2.9	4.7390	3.8	0.3193	2.4	0.64	1786.2	37.9	1774.2	31.7	1760.0	53.1	1760.0	53.1	101.5
Trial-1_0437	161	74955	3.6	12.6226	2.7	2.2095	3.5	0.2023	2.2	0.63	1187.5	23.5	1184.1	24.2	1177.7	53.3	1177.7	53.3	100.8
Trial-1_0438	484	556301	1.2	12.4047	1.5	2.2625	2.0	0.2036	1.2	0.63	1194.4	13.6	1200.7	14.0	1212.1	30.4	1212.1	30.4	98.5
Trial-1_0439	61	9886	0.5	19.0950	7.9	0.5006	9.9	0.0693	5.9	0.60	432.1	24.8	412.1	33.5	301.6	180.3	432.1	24.8	143.3
Trial-1_0440	314	100579	1.5	18.5947	3.6	0.4643	4.6	0.0626	2.8	0.60	391.5	10.4	387.2	14.7	361.8	82.1	391.5	10.4	NA
Trial-1_0441	222	324335	3.2	5.3698	1.3	13.4076	1.7	0.5222	1.2	0.66	2708.4	25.6	2708.8	16.5	2709.1	21.5	2709.1	21.5	100.0
Trial-1_0442	141	96011	1.4	11.0173	2.5	3.1956	3.3	0.2553	2.1	0.63	1466.0	27.2	1456.1	25.3	1441.7	48.3	1441.7	48.3	101.7
Trial-1_0443	199	1108473	1.2	13.5496	2.6	1.8198	3.3	0.1788	2.1	0.62	1060.6	20.2	1052.6	21.8	1036.1	52.6	1036.1	52.6	102.4
Trial-1_0444	234	145080	3.8	12.2558	2.2	2.4262	2.8	0.2157	1.7	0.63	1258.9	19.9	1250.4	20.0	1235.8	42.4	1235.8	42.4	101.9
Trial-1_0445	86	100352	2.7	13.5132	3.9	1.7804	4.9	0.1745	3.1	0.62	1036.8	29.5	1038.3	32.2	1041.5	78.2	1041.5	78.2	99.6
Trial-1_0446	72	12634	3.4	19.1086	8.5	0.3314	10.6	0.0459	6.4	0.60	289.5	18.0	290.6	26.8	299.9	193.3	289.5	18.0	NA
Trial-1_0447	37	17090	1.8	13.3184	5.9	1.8377	7.5	0.1775	4.7	0.62	1053.3	45.2	1059.0	49.2	1070.8	117.8	1070.8	117.8	98.4
Trial-1_0448	95	96825	2.3	13.5177	3.7	1.7612	4.7	0.1727	2.9	0.62	1026.8	28.0	1031.3	30.7	1040.8	74.9	1040.8	74.9	98.7
Trial-1_0449	201	44042	2.7	12.1763	2.4	2.1138	3.1	0.1867	2.0	0.63	1103.3	19.9	1153.3	21.6	1248.5	47.8	1248.5	47.8	88.4
Trial-1_0450	303	306427	1.3	12.5443	1.9	2.2086	2.5	0.2009	1.5	0.63	1180.4	16.6	1183.8	17.2	1190.0	37.9	1190.0	37.9	99.2
Trial-1_0451	376	256579	3.4	12.5165	1.7	2.2735	2.2	0.2064	1.4	0.63	1209.5	15.1	1204.1	15.4	1194.4	33.6	1194.4	33.6	101.3
Trial-1_0452	210	109467	1.3	11.9584	2.2	2.5082	2.8	0.2175	1.8	0.63	1268.9	20.6	1274.4	20.7	1283.8	43.2	1283.8	43.2	98.8
Trial-1_0453	210	328794	1.0	5.1354	1.3	13.5089	1.8	0.5031	1.2	0.67	2627.3	25.8	2715.9	17.0	2782.5	22.0	2782.5	22.0	94.4
Trial-1_0454	50	22535	1.6	13.5478	5.2	1.7532	6.6	0.1723	4.1	0.62	1024.6	39.0	1028.4	43.0	1036.3	105.3	1036.3	105.3	98.9
Trial-1_0455	620	50850	0.9	20.0588	3.8	0.1738	4.8	0.0253	3.0	0.63	160.9	4.8	162.7	7.2	188.1	87.5	160.9	4.8	NA
Trial-1_0456	77	43551	1.8	8.7635	2.9	5.3298	3.7	0.3388	2.4	0.64	1880.7	39.1	1873.7	31.9	1865.9	51.5	1865.9	51.5	100.8
Trial-1_0457	265	294982	3.8	12.7200	2.1	2.1626	2.7	0.1995	1.7	0.63	1172.7	18.0	1169.1	18.6	1162.5	41.4	1162.5	41.4	100.9
Trial-1_0458	132	447153	0.8	10.9958	2.6	3.1626	3.3	0.2522	2.1	0.63	1449.9	27.5	1448.1	25.8	1445.4	49.4	1445.4	49.4	100.3
Trial-1_0459	34	24217	1.0	13.1051	6.1	1.9609	7.8	0.1864	4.8	0.62	1101.7	48.9	1102.2	52.2	1103.1	121.4	1103.1	121.4	99.9
Trial-1_0460	54	52303	1.3	8.7127	3.4	5.1738	4.5	0.3269	2.9	0.64	1823.5	46.0	1848.3	38.3	1876.3	62.1	1876.3	62.1	97.2
Trial-1_0461	93	34294	1.8	13.6777	3.9	1.6995	4.9	0.1686	3.1	0.62	1004.3	28.6	1008.3	31.6	1017.0	78.5	1017.0	78.5	98.8
Trial-1_0462	82	169982	1.9	12.9202	3.8	2.0662	4.9	0.1936	3.0	0.62	1140.9	31.8	1137.7	33.3	1131.5	75.8	1131.5	75.8	100.8
Trial-1_0463	29	46714	1.5	7.5717	4.3	7.2800	5.6	0.3998	3.7	0.65	2168.0	67.4	2146.4	50.3	2125.7	74.9	2125.7	74.9	102.0
Trial-1_0464	256	115634	0.7	17.8581	3.9	0.5179	4.9	0.0671	3.0	0.60	418.6	12.0	423.8	17.0	452.2	87.0	418.6	12.0	92.6
Trial-1_0465	39	32077	0.6	13.1344	5.6	1.9642	7.2	0.1871	4.5	0.62	1105.7	45.7	1103.3	48.6	1098.7	112.9	1098.7	112.9	100.6
Trial-1_0466	140	343503	2.5	12.5562	2.8	2.2492	3.6	0.2048	2.3	0.63	1201.2	25.0	1196.5	25.6	1188.2	56.1	1188.2	56.1	101.1
Trial-1_0467	59	38690	4.2	13.0852	4.5	2.0502	5.8	0.1946	3.6	0.62	1146.1	38.0	1132.4	39.6	1106.2	90.7	1106.2	90.7	103.6
Trial-1_0468	29	103175	1.4	13.6045	6.8	1.7510	8.7	0.1728	5.4	0.62	1027.3	51.5	1027.5	56.5	1027.9	138.3	1027.9	138.3	99.9
Trial-1_0469	54	191454	1.3	12.8944	4.7	2.1245	6.1	0.1987	3.8	0.62	1168.2	40.5	1156.8	42.0	1135.4	94.6	1135.4	94.6	102.9
Trial-1_0470	72	911199	1.8	13.1477	4.2	1.9283	5.3	0.1839	3.3	0.62	1088.1	33.2	1090.9	35.6	1096.6	83.4	1096.6	83.4	99.2
Trial-1_0471	94	123867	0.5	8.6853	2.6	5.3295	3.4	0.3357	2.2	0.64	1866.0	35.4	1873.6	29.0	1882.0	46.7	1882.0	46.7	99.2
Trial-1_0472	109	183005	1.6	5.3812	1.9	12.8643	2.5	0.5021	1.7	0.66	2622.7	36.2	2669.8	23.9	2705.6	31.3	2705.6	31.3	96.9
Trial-1_0473	58	11584	1.6	17.7170	7.2	0.6592	9.0	0.0847	5.5	0.61	524.2	27.5	514.2	36.4	469.8	158.8	524.2	27.5	111.6
Trial-1_0474	200	255940	3.5	8.9721	1.8	4.9722	2.4	0.3235	1.5	0.64	1807.0	24.2	1814.6	20.2	1823.3	33.2	1823.3	33.2	99.1
Trial-1_0475	99	169743	3.6	9.0891	2.6	4.9999	3.4	0.3296	2.2	0.64	1836.4	34.6	1819.3	28.5	1799.8	46.9	1799.8	46.9	102.0
Trial-1_0476	198	49033	0.6	17.8928	4.4	0.5015	5.5	0.0651	3.4	0.60	406.4	13.2	412.7	18.8	447.9	98.3	406.4	13.2	90.7
Trial-1_0477	147	57631	1.7	13.5400	3.0	1.7628	3.9	0.1731	2.4	0.62	1029.2	23.0	1031.9	25.1	1037.5	61.4	1037.5	61.4	99.2
Trial-1_0478	203	40303	3.2	18.2182	4.2	0.5247	5.3	0.0693	3.2	0.60	432.1	13.4	428.3	18.6	407.7	95.1	432.1	13.4	106.0
Trial-1_0479	48	994678	1.8	5.2765	2.8	13.8952	3.7	0.5317	2.5	0.66	2748.8	55.3	2742.6	35.3	2738.0	45.8	2738.0	45.8	100.4
Trial-1_0480	149	305768	1.8	12.9711	2.8	2.0441	3.6	0.1923	2.3	0.62	1133.8	23.7	1130.3	24.9	1123.6	56.8	1123.6	56.8	100.9
Trial-1_0481	149	137577	1.1	10.9304	2.5	3.2289	3.2	0.2560	2.0	0.63	1469.2	26.4	1464.1	24.6	1456.8	46.8	1456.8	46.8	100.9
Trial-1_0482	137	480991	0.7	5.6643	1.7	12.1764	2.3	0.5002	1.5	0.66	2614.8	32.5	2618.1	21.4	2620.7	28.5	2620.7	28.5	99.8
Trial-1_0483	76	88466	2.5	10.9259	3.4	3.1783	4.4	0.2519	2.8	0.64	1448.0	36.0	1451.9	33.7	1457.5	64.1	1457.5	64.1	99.3
Trial-1_0484	592	131558	1.2	19.2109	3.2	0.3077	4.0	0.0429	2.4	0.60	270.6	6.3	272.4	9.5	287.8	72.8	270.6	6.3	NA
Trial-1_0485	105	49979	2.1	13.4729	3.5	1.8924	4.5	0.1849	2.8	0.62	1093.8	28.3	1078.4	30.0	1047.6	71.3	1047.6	71.3	104.4
Trial-1_0486	39	90477	3.8	11.4021	5.0	2.8395	6.5	0.2348	4.1	0.63	1359.7	49.9	1366.0	48.5	1376.0	96.4	1376.0	96.4	98.8
Trial-1_0487	31	33288	1.2	4.4815	3.2	18.5138	4.3	0.6017	2.9	0.67	3036.9	70.4	3016.8	41.8	3003.5	51.8	3003.5	51.8	101.1
Trial-1_0488	26	22801	2.4	7.6773	4.6	6.8979	6.1	0.3841	3.9	0.65	2095.3	70.5	2098.4	53.9	2101.4	81.0	2101.4	81.0	99.7
Trial-1_0489	45	28182	1.7	13.7027	5.6	1.7415	7.2	0.1731	4.4	0.62	1029.0	42.3	1024.0	46.2	1013.3	113.7	1013.3	113.7	101.5
Trial-1_0490	98	419549	1.1	9.3163	2.7	4.6312	3.5	0.3129	2.2	0.64	1755.1	34.5	1754.9	29.3	1754.7	49.2	1754.7	49.2	100.0
Trial-1_0491	166	228458	1.4	12.4615	2.7	2.1397	3.5	0.1934	2.2	0.63	1139.7	22.7	1161.7	24.0	1203.1	53.2	1203.1	53.2	94.7
Trial-1_0492	134	160311	1.2	8.8292	2.2	5.2307	2.9	0.3349	1.9	0.64	1862.3	30.2	1857.6	24.8	1852.4	40.2	1852.4	40.2	100.5
Trial-1_0493	159	101664	1.6	18.4109	5.1	0.4667	6.4	0.0623	3.9	0.60	389.7	14.6	388.9	20.7	384.1	115.0	389.7	14.6	NA
Trial-1_0494	154	59561	5.8	13.3691	2.9	1.8550	3.8	0.1799	2.3	0.62	1066.2	23.1	1065.2	24.9	1063.1	59.3	1063.1	59.3	100.3
Trial-1_0495	207	485952	1.0	10.8633	2.1	3.1440	2.7	0.2477	1.7	0.63	1426.7	21.7	1443.5	20.6	1468.5	39.3	1468.5	39.3	97.2
Trial-1_0496	348	122459	0.9	16.4711	2.7	0.8549	3.4	0.1021	2.0	0.61	626.8	12.2	627.3	15.7	629.0	57.3	626.8	12.2	99.7
Trial-1_0497	76	32095	3.5	12.4395	3.9	2.2706	5.0	0.2049	3.2	0.63	1201.4	34.6	1203.2	35.5	1206.6	77.4	1206.6	77.4	99.6
Trial-1_0498	59	126962	3.4	9.2791	3.4	4.7758	4.5	0.3214	2.9	0.64	1796.6	45.0	1780.6	37.6	1762.0	62.7	1762.0	62.7	102.0
Trial-1_0499	39	97963	1.3	13.0135	5.8	1.9970	7.4	0.1885	4.6	0.62	1113.2	47.1	1114.5	50.0	1117.2	115.2	1117.2	115.2	99.6
Trial-1_0500	103	15534	1.6	20.1964	7.9	0.2852	9.8	0.0418	5.9	0.60	263.9	15.2	254.8	22.2	172.2	184.8	263.9	15.2	NA
Trial-1_0501	75	65728	1.2	11.0200	3.5	3.2547	4.5	0.2601	2.8	0.63	1490.5	37.7	1470.3	34.8	1441.2	66.1	1441.2	66.1	103.4
Trial-1_0502	48	267199	2.2	9.3457	3.8	4.6936	5.0	0.3181	3.2	0.64	1780.6	49.9	1766.1	41.9	1748.9	70.3	1748.9	70.3	101.8
Trial-1_0503	196	19297	1.2	20.4286	7.1	0.1819	8.9	0.0270	5.3	0.60	171.5	9.0	169.7	13.9	145.5	167.5	171.5	9.0	NA
Trial-1_0504	51	40605	1.5	12.9963	5.0	2.0025	6.4	0.1888	4.0	0.62	1114.6	40.8	1116.4	43.3	1119.8	99.7	1119.8	99.7	99.5
Trial-1_0505	461	196222	3.0	13.4236	1.7	1.8292	2.2	0.1781	1.4	0.62	1056.5	13.4	1056.0	14.5	1054.9	34.8	1054.9	34.8	100.1
Trial-1_0506	62	35739	1.8	11.9786	4.1	2.5869	5.3	0.2247	3.3	0.63	1306.9	39.3	1296.9	38.8	1280.5	80.3	1280.5	80.3	102.1
Trial-1_0507	107	17765	0.7	20.5440	9.2	0.1989	11.5	0.0296	6.8	0.60	188.3	12.7	184.2	19.3	132.3	216.8	188.3	12.7	NA
Trial-1_0508	202	7130	1.6	21.2565	9.4	0.1004	11.6	0.0155	6.8	0.59	99.1	6.7	97.2	10.7	51.5	223.9	99.1	6.7	NA
Trial-1_0509	98	70649	0.3	13.5041	3.6	1.8895	4.7	0.1851	2.9	0.62	1094.5	29.1	1077.4	30.9	1042.9	73.5	1042.9	73.5	105.0
Trial-1_0510	67	128749	1.8	9.9170	3.4	4.0181	4.4	0.2890	2.8	0.64	1636.5	40.7	1637.9	36.0	1639.5	63.3	1639.5	63.3	99.8
Trial-1_0511	31	70184	0.6	10.8665	5.3	3.3975	6.8	0.2678	4.3	0.63	1529.4	58.6	1503.8	53.4	1467.9	100.0	1467.9	100.0	104.2
Trial-1_0512	130	33891	86.8	18.3766	5.3	0.5434	6.6	0.0724	4.0	0.60	450.7	17.4	440.6	23.7	388.3	118.7	450.7	17.4	116.1
Trial-1_0513	59	19819	2.3	12.4773	4.5	2.2894	5.7	0.2072	3.6	0.63	1213.8	39.5	1209.0	40.4	1200.6	87.8	1200.6	87.8	101.1
Trial-1_0514	42	13112	2.1	13.1507	5.6	1.9259	7.1	0.1837	4.4	0.62	1087.1	44.4	1090.1	47.7	1096.2	111.8	1096.2	111.8	99.2
Trial-1_0515	84	188260	2.7	13.5300	4.0	1.8029	5.1	0.1769	3.2	0.62	1050.1	31.0	1046.5	33.6	1039.0	81.3	1039.0	81.3	101.1
Trial-1_0516	120	67200	2.0	12.9680	3.3	1.9856	4.2	0.1868	2.6	0.62	1103.8	26.4	1110.6	28.2	1124.1	65.2	1124.1	65.2	98.2
Trial-1_0517	137	61174	1.4	19.8106	6.6	0.3118	8.2	0.0448	4.9	0.60	282.5	13.6	275.6	19.8	217.0	152.7	282.5	13.6	NA
Trial-1_0518	58	34209	1.5	13.2173	4.7	1.9607	6.1	0.1879	3.8	0.62	1110.3	38.4	1102.1	40.7	1086.0	95.0	1086.0	95.0	102.2
Trial-1_0519	251	100875	3.0	9.1492	1.7	4.8076	2.2	0.3190	1.4	0.64	1784.9	21.9	1786.2	18.4	1787.8	30.5	1787.8	30.5	99.8
Trial-1_0520	105	93843	0.8	9.8465	2.7	4.1753	3.5	0.2982	2.2	0.64	1682.2	33.0	1669.2	28.6	1652.8	49.8	1652.8	49.8	101.8
Trial-1_0521	174	93553	3.2	12.6502	2.6	2.2075	3.3	0.2025	2.1	0.63	1188.9	22.7	1183.4	23.4	1173.4	51.6	1173.4	51.6	101.3
Trial-1_0522	209	331344	3.1	4.8247	1.3	15.9536	1.8	0.5583	1.2	0.67	2859.4	27.1	2874.0	16.8	2884.3	21.3	2884.3	21.3	99.1
Trial-1_0523	83	141881	0.9	4.8500	2.1	16.1432	2.8	0.5678	1.8	0.67	2898.9	43.1	2885.3	26.5	2875.8	33.5	2875.8	33.5	100.8
Trial-1_0524	140	145117	2.3	12.9789	3.0	2.0478	3.9	0.1928	2.4	0.62	1136.3	25.0	1131.6	26.3	1122.4	60.2	1122.4	60.2	101.2
Trial-1_0525	31	49138	3.0	12.3458	5.9	2.4104	7.6	0.2158	4.8	0.63	1259.8	54.8	1245.7	54.9	1221.4	117.0	1221.4	117.0	103.1
Trial-1_0526	118	101040	3.2	9.2931	2.4	4.6155	3.1	0.3111	2.0	0.64	1746.0	30.5	1752.1	26.0	1759.3	43.6	1759.3	43.6	99.2
Trial-1_0527	120	126911	1.1	5.3999	1.8	13.4855	2.4	0.5281	1.6	0.66	2733.6	35.5	2714.3	22.7	2699.9	29.7	2699.9	29.7	101.2
Trial-1_0528	140	269359	14.9	13.3439	3.1	1.8897	3.9	0.1829	2.4	0.62	1082.7	24.4	1077.5	26.1	1066.9	61.8	1066.9	61.8	101.5
Trial-1_0529	218	117891	1.8	8.4791	1.7	5.6708	2.2	0.3487	1.4	0.65	1928.5	23.9	1926.9	19.2	1925.2	30.4	1925.2	30.4	100.2
Trial-1_0530	83	31662	1.3	19.3746	8.1	0.3636	10.1	0.0511	6.0	0.60	321.2	18.9	314.9	27.3	268.3	185.7	321.2	18.9	NA
Trial-1_0531	19	45855	1.5	12.7620	8.5	1.9279	10.9	0.1784	6.8	0.62	1058.5	66.6	1090.8	73.2	1155.9	169.8	1155.9	169.8	91.6
Trial-1_0532	31	20264	1.4	11.0400	5.4	3.1321	7.0	0.2508	4.4	0.63	1442.5	57.3	1440.6	54.0	1437.8	103.6	1437.8	103.6	100.3
Trial-1_0533	47	31967	1.1	13.4803	5.4	1.8626	6.8	0.1821	4.3	0.62	1078.4	42.3	1067.9	45.2	1046.5	108.0	1046.5	108.0	103.1
Trial-1_0534	125	767903	3.4	13.3339	3.3	1.8457	4.2	0.1785	2.6	0.62	1058.7	25.5	1061.9	27.6	1068.4	65.9	1068.4	65.9	99.1
Trial-1_0535	211	62708	2.0	12.8489	2.4	2.1626	3.0	0.2015	1.9	0.62	1183.5	20.4	1169.1	21.0	1142.5	47.0	1142.5	47.0	103.6
Trial-1_0536	477	123940	4.4	16.4631	2.4	0.7432	3.0	0.0887	1.8	0.61	548.1	9.7	564.3	13.1	630.1	51.7	548.1	9.7	87.0
Trial-1_0537	154	26146	21.8	17.2885	4.5	0.6477	5.6	0.0812	3.4	0.60	503.4	16.4	507.1	22.3	523.8	97.8	503.4	16.4	96.1
Trial-1_0538	47	27141	1.2	18.6115	11.0	0.3352	13.7	0.0452	8.2	0.60	285.2	23.0	293.5	35.0	359.7	248.6	285.2	23.0	NA
Trial-1_0539	79	298625	0.7	3.1358	1.8	31.8516	2.4	0.7244	1.7	0.68	3512.5	44.8	3545.7	23.9	3564.5	27.4	3564.5	27.4	98.5
Trial-1_0540	113	13104	1.7	16.1648	4.8	0.8271	6.0	0.0970	3.7	0.61	596.6	20.8	612.0	27.6	669.4	101.9	596.6	20.8	89.1
Trial-1_0541	99	62623	3.1	12.6934	3.5	2.0454	4.5	0.1883	2.8	0.62	1112.2	28.6	1130.8	30.6	1166.6	69.6	1166.6	69.6	95.3
Trial-1_0542	103	974678	1.3	5.4032	1.9	13.2246	2.6	0.5182	1.7	0.66	2691.7	37.3	2695.8	24.1	2698.9	31.6	2698.9	31.6	99.7
Trial-1_0543	79	31776	1.7	16.9801	6.0	0.6971	7.6	0.0858	4.6	0.61	530.9	23.5	537.0	31.7	563.1	131.7	530.9	23.5	94.3
Trial-1_0544	38	65790	2.1	4.5260	2.9	18.1367	3.9	0.5953	2.6	0.67	3011.1	62.7	2997.0	37.5	2987.6	46.5	2987.6	46.5	100.8
Trial-1_0545	23	55566	1.4	13.0138	7.3	2.0120	9.3	0.1899	5.8	0.62	1120.8	59.7	1119.6	63.3	1117.1	145.9	1117.1	145.9	100.3

Trial-1_0546	36	68619	1.6	16.4149	9.1	0.7011	11.4	0.0835	7.0	0.61	516.8	34.5	539.4	48.0	636.4	196.1	516.8	34.5	81.2
Trial-1_0547	162	192094	2.1	5.5685	1.7	10.6910	2.2	0.4318	1.5	0.66	2313.7	28.4	2496.6	20.5	2649.0	27.6	2649.0	27.6	87.3
Trial-1_0548	68	104414	0.6	5.2178	2.3	13.8732	3.1	0.5250	2.1	0.66	2720.4	46.1	2741.1	29.6	2756.4	38.4	2756.4	38.4	98.7
Trial-1_0549	51	12177	1.2	16.3793	7.5	0.7256	9.4	0.0862	5.7	0.61	533.0	29.4	553.9	40.3	641.1	161.4	533.0	29.4	83.1
Trial-1_0550	174	221545	2.1	10.9531	2.3	3.0488	3.0	0.2422	1.9	0.63	1398.1	23.8	1419.9	22.9	1452.8	44.2	1452.8	44.2	96.2
Trial-1_0551	110	105707	1.7	9.0959	2.5	4.8732	3.2	0.3215	2.1	0.64	1797.0	32.5	1797.6	27.2	1798.4	45.1	1798.4	45.1	99.9
Trial-1_0552	32	158583	1.6	5.0309	3.3	15.5818	4.4	0.5685	2.9	0.67	2901.8	68.2	2851.5	41.8	2816.1	53.3	2816.1	53.3	103.0
Trial-1_0553	49	40579	1.4	8.8239	3.6	5.3457	4.7	0.3421	3.0	0.64	1896.8	49.3	1876.2	39.9	1853.4	64.6	1853.4	64.6	102.3
Trial-1_0554	831	687318	19.9	11.9867	1.2	2.3224	1.5	0.2019	0.9	0.63	1185.5	10.2	1219.1	10.7	1279.2	22.9	1279.2	22.9	92.7
Trial-1_0555	40	13993	1.1	12.5977	5.6	2.0554	7.2	0.1878	4.5	0.62	1109.4	45.7	1134.1	49.0	1181.6	110.8	1181.6	110.8	93.9
Trial-1_0556	291	24997	1.7	18.7200	4.5	0.3233	5.7	0.0439	3.4	0.60	276.9	9.2	284.4	14.0	346.6	102.4	276.9	9.2	NA
Trial-1_0557	59	42548	1.3	13.0793	4.6	1.9060	5.9	0.1808	3.7	0.62	1071.4	36.3	1083.2	39.4	1107.0	92.5	1107.0	92.5	96.8
Trial-1_0558	57	160050	1.7	13.0317	4.8	1.9673	6.1	0.1859	3.8	0.62	1099.3	38.6	1104.4	41.3	1114.3	95.8	1114.3	95.8	98.7
Trial-1_0559	19	12914	2.2	12.8977	8.1	2.0262	10.4	0.1895	6.5	0.62	1118.9	66.5	1124.3	70.7	1134.9	161.7	1134.9	161.7	98.6
Trial-1_0560	448	12928	2.7	20.6122	6.0	0.1099	7.5	0.0164	4.4	0.59	105.0	4.6	105.9	7.5	124.4	141.9	105.0	4.6	NA
Trial-1_0561	66	86772	0.4	9.6243	3.4	4.1318	4.5	0.2884	2.9	0.64	1633.6	41.1	1660.6	36.5	1695.0	63.3	1695.0	63.3	96.4
Trial-1_0562	18	3148	1.5	17.5533	12.7	0.6959	16.0	0.0886	9.7	0.60	547.2	50.7	536.3	66.7	490.3	282.1	547.2	50.7	111.6
Trial-1_0563	46	60114	1.4	5.2600	2.8	14.6788	3.7	0.5600	2.5	0.66	2866.6	57.5	2794.7	35.6	2743.1	45.9	2743.1	45.9	104.5
Trial-1_0564	40	89203	1.8	13.3749	5.7	1.7712	7.3	0.1718	4.5	0.62	1022.1	42.8	1035.0	47.4	1062.3	115.0	1062.3	115.0	96.2
Trial-1_0565	30	13379	1.2	13.2170	6.6	1.9030	8.4	0.1824	5.3	0.62	1080.2	52.3	1082.1	56.3	1086.1	132.6	1086.1	132.6	99.5
Trial-1_0566	81	8895	2.0	18.3010	7.2	0.5048	9.0	0.0670	5.4	0.60	418.1	21.8	414.9	30.5	397.6	160.6	418.1	21.8	105.2
Trial-1_0567	85	56346	2.9	12.6813	3.7	2.0905	4.8	0.1923	3.0	0.62	1133.6	31.0	1145.7	32.8	1168.5	73.7	1168.5	73.7	97.0
Trial-1_0568	115	455251	1.5	8.8061	2.4	5.2860	3.1	0.3376	2.0	0.64	1875.1	32.8	1866.6	26.8	1857.1	43.4	1857.1	43.4	101.0
Trial-1_0569	36	31535	1.9	12.5997	5.7	2.1290	7.3	0.1946	4.6	0.63	1146.0	48.2	1158.3	50.7	1181.3	113.2	1181.3	113.2	97.0
Trial-1_0570	66	60537	0.1	12.5740	4.4	2.0818	5.6	0.1898	3.5	0.63	1120.6	36.2	1142.8	38.6	1185.3	86.6	1185.3	86.6	94.5
Trial-1_0571	67	60592	1.5	11.0352	3.7	3.1278	4.7	0.2503	3.0	0.63	1440.2	38.6	1439.5	36.4	1438.6	69.9	1438.6	69.9	100.1
Trial-1_0572	16	35834	0.7	13.8475	9.9	1.5873	12.6	0.1594	7.8	0.62	953.5	69.0	965.2	78.4	992.0	200.8	992.0	200.8	96.1
Trial-1_0573	14	35637	1.4	7.4901	6.4	7.3101	8.4	0.3971	5.4	0.65	2155.7	99.7	2150.0	75.0	2144.6	111.6	2144.6	111.6	100.5
Trial-1_0574	118	60019	1.9	13.3625	3.4	1.8017	4.3	0.1746	2.7	0.62	1037.5	25.7	1046.1	28.2	1064.1	68.1	1064.1	68.1	97.5
Trial-1_0575	36	49177	2.7	9.5806	4.4	4.1147	5.7	0.2859	3.7	0.64	1621.1	52.4	1657.2	46.7	1703.4	80.8	1703.4	80.8	95.2
Trial-1_0576	18	11857	1.0	13.3397	8.3	1.8297	10.6	0.1770	6.6	0.62	1050.7	64.1	1056.2	69.9	1067.6	167.5	1067.6	167.5	98.4
Trial-1_0577	35	3793	1.9	21.4724	14.0	0.2609	14.2	0.0406	2.3	0.16	256.7	5.8	235.4	29.9	27.3	337.6	256.7	5.8	NA
Trial-1_0578	191	73644	2.1	12.8507	2.6	2.0405	3.3	0.1902	2.1	0.62	1122.3	21.5	1129.1	22.8	1142.2	52.0	1142.2	52.0	98.3
Trial-1_0579	265	297551	3.5	9.4733	1.8	3.9281	2.3	0.2699	1.5	0.64	1540.2	20.4	1619.5	18.9	1724.1	33.0	1724.1	33.0	89.3
Trial-1_0580	202	95482	2.5	13.1886	2.5	2.0162	3.2	0.1929	2.0	0.62	1136.8	20.9	1121.0	21.9	1090.4	50.6	1090.4	50.6	104.3
Trial-1_0581	289	389361	8.1	12.9570	2.1	1.9464	2.7	0.1829	1.7	0.62	1082.8	16.7	1097.2	18.1	1125.8	41.9	1125.8	41.9	96.2
Trial-1_0582	110	77500	2.9	9.5623	2.6	4.3478	3.4	0.3015	2.2	0.64	1698.9	32.8	1702.5	28.4	1706.9	48.7	1706.9	48.7	99.5
Trial-1_0583	144	127780	1.9	17.0125	4.5	0.7547	5.6	0.0931	3.4	0.61	574.0	18.7	571.0	24.5	558.9	97.3	574.0	18.7	102.7
Trial-1_0584	93	100076	2.1	9.2721	2.7	4.6149	3.5	0.3103	2.3	0.64	1742.4	34.4	1752.0	29.4	1763.4	49.4	1763.4	49.4	98.8
Trial-1_0585	133	21822	4.4	17.7430	5.0	0.6147	6.3	0.0791	3.8	0.60	490.8	18.0	486.5	24.5	466.6	111.9	490.8	18.0	105.2
Trial-1_0586	165	100160	2.7	12.4860	2.6	2.2575	3.4	0.2044	2.1	0.63	1199.1	23.2	1199.1	23.9	1199.2	52.2	1199.2	52.2	100.0
Trial-1_0587	28	78052	28.5	8.7322	4.8	5.1425	6.2	0.3257	4.0	0.64	1817.4	63.5	1843.2	53.0	1872.3	85.9	1872.3	85.9	97.1
Trial-1_0588	38	27780	2.5	13.4547	5.8	1.8219	7.3	0.1778	4.6	0.62	1054.8	44.5	1053.4	48.2	1050.3	116.1	1050.3	116.1	100.4
Trial-1_0589	59	29424	0.8	12.9680	4.6	2.0406	5.8	0.1919	3.6	0.62	1131.8	37.8	1129.2	39.8	1124.1	91.1	1124.1	91.1	100.7
Trial-1_0590	55	104862	2.4	8.6600	3.4	5.7129	4.5	0.3588	2.9	0.64	1976.5	49.2	1933.3	38.8	1887.3	61.8	1887.3	61.8	104.7
Trial-1_0591	19	48374	0.7	4.8436	4.2	16.1577	5.7	0.5676	3.8	0.67	2898.0	88.2	2886.2	54.2	2877.9	68.5	2877.9	68.5	100.7
Trial-1_0592	59	197142	0.9	13.0231	4.7	2.0915	6.0	0.1975	3.7	0.62	1162.1	39.5	1146.0	40.9	1115.6	93.0	1115.6	93.0	104.2
Trial-1_0593	40	39640	3.0	10.9957	5.1	3.0919	6.5	0.2466	4.1	0.63	1420.8	52.2	1430.7	50.0	1445.4	96.4	1445.4	96.4	98.3
Trial-1_0594	92	65969	1.8	13.3459	3.9	1.8698	5.0	0.1810	3.1	0.62	1072.3	30.7	1070.4	33.1	1066.6	78.9	1066.6	78.9	100.5
Trial-1_0595	129	89980	4.6	9.3372	2.9	4.4311	3.7	0.3001	2.4	0.64	1691.6	35.4	1718.2	30.9	1750.6	52.6	1750.6	52.6	96.6
Trial-1_0596	45	89992	1.7	9.2178	3.9	4.3671	5.1	0.2920	3.3	0.64	1651.3	48.0	1706.1	42.4	1774.1	71.9	1774.1	71.9	93.1
Trial-1_0597	31	15830	1.4	11.0340	5.5	3.1120	7.1	0.2490	4.5	0.63	1433.5	57.5	1435.7	54.6	1438.8	105.1	1438.8	105.1	99.6
Trial-1_0598	134	54179	1.4	9.7808	2.6	4.0353	3.4	0.2863	2.2	0.63	1622.8	30.9	1641.3	27.6	1665.2	48.6	1665.2	48.6	97.5
Trial-1_0599	160	196555	1.2	19.3503	6.5	0.2977	8.1	0.0418	4.8	0.60	263.8	12.5	264.6	18.9	271.2	149.1	263.8	12.5	NA
Trial-1_0600	141	138027	1.9	13.1260	3.1	1.9677	4.0	0.1873	2.5	0.62	1106.8	25.1	1104.5	26.7	1099.9	62.0	1099.9	62.0	100.6
Trial-1_0601	216	224346	2.9	12.9165	2.4	1.9115	3.1	0.1791	1.9	0.62	1061.9	18.7	1085.1	20.5	1132.0	47.8	1132.0	47.8	93.8
Trial-1_0602	355	210356	3.8	14.0605	2.1	1.6309	2.6	0.1663	1.6	0.62	991.7	15.0	982.2	16.6	960.9	42.4	960.9	42.4	103.2
Trial-1_0603	571	719009	11.8	13.0846	1.5	1.9899	2.0	0.1888	1.2	0.62	1115.1	12.5	1112.1	13.2	1106.2	30.5	1106.2	30.5	100.8
Trial-1_0604	44	152188	1.3	12.9957	5.5	2.0548	7.0	0.1937	4.3	0.62	1141.2	45.5	1133.9	47.7	1119.9	109.0	1119.9	109.0	101.9
Trial-1_0605	83	15473	2.2	16.4919	5.9	0.8505	7.3	0.1017	4.4	0.60	624.5	25.9	624.9	34.0	626.3	126.2	624.5	25.9	99.7
Trial-1_0606	17	22420	0.4	17.2388	14.3	0.6060	14.8	0.0758	3.9	0.26	470.8	17.6	481.0	56.9	530.1	315.5	470.8	17.6	88.8
Trial-1_0607	58	51791	1.4	8.4270	3.2	5.5145	4.2	0.3370	2.7	0.64	1872.4	44.5	1902.9	36.5	1936.2	58.1	1936.2	58.1	96.7
Trial-1_0608	52	192586	1.0	5.3570	2.7	13.4434	3.6	0.5223	2.4	0.66	2709.0	53.4	2711.3	34.4	2713.1	44.9	2713.1	44.9	99.8
Trial-1_0609	193	109544	4.5	13.1983	2.6	1.8525	3.3	0.1773	2.0	0.62	1052.4	19.7	1064.3	21.5	1088.9	51.2	1088.9	51.2	96.6
Trial-1_0610	139	39886	1.9	18.1977	5.6	0.4853	7.0	0.0641	4.2	0.60	400.2	16.3	401.7	23.2	410.2	125.3	400.2	16.3	97.6
Trial-1_0611	25	61181	2.3	12.7762	6.9	2.0359	8.9	0.1887	5.5	0.62	1114.1	56.7	1127.6	60.5	1153.7	137.8	1153.7	137.8	96.6
Trial-1_0612	50	77216	1.8	5.3539	2.7	13.1446	3.6	0.5104	2.4	0.66	2658.4	52.1	2690.1	34.0	2714.0	44.4	2714.0	44.4	97.9
Trial-1_0613	24	96022	4.2	13.1513	7.5	1.9847	9.6	0.1893	6.0	0.62	1117.6	61.3	1110.3	64.9	1096.1	150.7	1096.1	150.7	102.0
Trial-1_0614	57	3961	1.0	22.0298	13.5	0.1886	16.5	0.0301	9.5	0.57	191.4	17.9	175.5	26.7	-34.5	330.0	191.4	17.9	NA
Trial-1_0615	83	78020	1.9	13.1434	3.9	1.8484	5.0	0.1762	3.1	0.62	1046.2	30.2	1062.8	33.1	1097.3	78.8	1097.3	78.8	95.3
Trial-1_0616	24	6160	1.6	13.4615	7.9	1.7462	10.1	0.1705	6.2	0.62	1014.8	58.3	1025.8	65.1	1049.3	160.0	1049.3	160.0	96.7
Trial-1_0617	314	277763	4.0	13.0274	2.0	2.0962	2.6	0.1981	1.6	0.62	1164.9	17.2	1147.6	17.8	1115.0	40.4	1115.0	40.4	104.5
Trial-1_0618	21	15676	0.7	8.3637	5.3	5.6902	7.0	0.3452	4.5	0.65	1911.5	74.4	1929.9	60.3	1949.7	95.2	1949.7	95.2	98.0
Trial-1_0619	123	134286	1.6	13.3231	3.3	1.7667	4.2	0.1707	2.6	0.62	1016.0	24.4	1033.3	27.1	1070.1	65.7	1070.1	65.7	95.0
Trial-1_0620	36	93443	1.5	12.4151	5.5	2.2522	7.1	0.2028	4.4	0.63	1190.3	48.1	1197.5	49.8	1210.4	108.7	1210.4	108.7	98.3
Trial-1_0621	23	57075	1.7	8.8977	5.3	4.9204	6.9	0.3175	4.4	0.64	1777.6	68.7	1805.8	58.1	1838.4	95.3	1838.4	95.3	96.7
Trial-1_0622	114	122336	1.5	4.7435	1.7	16.4568	2.3	0.5662	1.6	0.67	2892.1	36.2	2903.7	22.3	2911.8	28.0	2911.8	28.0	99.3
Trial-1_0623	83	44690	1.4	12.6704	3.8	2.1551	4.9	0.1980	3.1	0.62	1164.8	32.7	1166.7	34.0	1170.2	75.9	1170.2	75.9	99.5
Trial-1_0624	89	182449	2.1	12.3825	3.5	2.1978	4.5	0.1974	2.8	0.63	1161.2	29.8	1180.3	31.2	1215.6	68.6	1215.6	68.6	95.5
Trial-1_0625	186	70309	-949.7	13.3864	2.7	1.8548	3.4	0.1801	2.1	0.62	1067.4	21.0	1065.1	22.6	1060.5	54.1	1060.5	54.1	100.6
Trial-1_0626	61	85368	1.7	8.9918	3.3	4.7035	4.3	0.3067	2.7	0.64	1724.7	41.4	1767.9	35.6	1819.3	59.2	1819.3	59.2	94.8
Trial-1_0627	53	48064	3.5	10.8889	4.3	3.2836	5.5	0.2593	3.5	0.62	1486.3	45.9	1477.2	43.1	1464.0	82.3	1464.0	82.3	101.5
Trial-1_0628	105	61503	2.2	13.2066	3.5	1.9015	4.4	0.1821	2.8	0.62	1078.6	27.3	1081.6	29.4	1087.7	69.3	1087.7	69.3	99.2
Trial-1_0629	58	53857	2.6	11.0640	3.7	3.0946	4.8	0.2483	3.1	0.63	1429.8	39.2	1431.3	37.1	1433.6	71.4	1433.6	71.4	99.7
Trial-1_0630	416	199187	6.1	13.3472	1.7	1.8332	2.2	0.1775	1.4	0.62	1053.1	13.3	1057.4	14.4	1066.4	34.5	1066.4	34.5	98.8
Trial-1_0631	36	31477	0.8	8.0063	4.0	6.2482	5.2	0.3628	3.4	0.65	1995.5	58.1	2011.2	45.9	2027.4	70.8	2027.4	70.8	98.4
Trial-1_0632	153	77372	1.6	12.9281	3.0	2.0968	3.8	0.1966	2.3	0.62	1157.0	24.9	1147.8	26.0	1130.3	59.0	1130.3	59.0	102.4
Trial-1_0633	110	699247	1.6	10.9099	3.3	2.9650	4.2	0.2346	2.7	0.63	1358.6	32.8	1398.7	32.2	1460.3	62.6	1460.3	62.6	93.0
Trial-1_0634	128	61790	4.7	12.7623	3.2	2.1097	4.0	0.1953	2.5	0.62	1149.9	26.4	1152.0	27.8	1155.9	62.6	1155.9	62.6	99.5
Trial-1_0635	34	116827	1.2	11.0530	3.7	2.9281	5.9	0.2347	4.6	0.78	1359.2	56.2	1389.2	44.6	1435.5	70.6	1435.5	70.6	94.7
Trial-1_0636	120	39456	1.6	18.2044	7.2	0.3653	9.0	0.0482	5.4	0.60	303.7	15.9	316.2	24.3	409.4	160.8	303.7	15.9	NA
Trial-1_0637	65	133491	2.2	9.3434	3.5	4.3229	4.5	0.2929	2.9	0.64	1656.2	42.0	1697.7	37.1	1749.4	63.2	1749.4	63.2	94.7
Trial-1_0638	80	55029	2.5	12.8766	4.1	2.0010	5.2	0.1869	3.2	0.62	1104.4	32.7	1115.8	35.2	1138.2	80.9	1138.2	80.9	97.0
Trial-1_0639	147	48288	3.0	17.4177	5.4	0.5401	6.7	0.0682	4.0	0.60	425.5	16.7	438.5	23.9	507.4	118.1	425.5	16.7	83.9
Trial-1_0640	30	37767	0.9	5.3396	3.8	13.2260	5.0	0.5122	3.3	0.66	2666.0	72.8	2695.9	47.7	2718.4	62.5	2718.4	62.5	98.1
Trial-1_0641	342	109580	0.5	17.7025	3.9	0.4282	4.9	0.0550	3.0	0.60	345.0	9.9	361.9	15.0	471.6	87.1	345.0	9.9	NA
Trial-1_0642	250	48126	2.3	16.8052	3.7	0.6751	4.6	0.0823	2.8	0.60	509.7	13.7	523.8	18.9	585.7	79.9	509.7	13.7	87.0
Trial-1_0643	145	100689	1.0	9.7391	2.4	4.0407	3.2	0.2854	2.0	0.63	1618.6	28.7	1642.4	25.8	1673.1	45.3	1673.1	45.3	96.7
Trial-1_0644	35	22167	1.2	8.9274	4.3	4.9304	5.7	0.3192	3.6	0.64	1786.0	56.6	1807.5	47.8	1832.3	78.8	1832.3	78.8	97.5
Trial-1_0645	24	48588	2.0	3.6513	3.5	24.6081	4.7	0.6517	3.2	0.68	3234.7	81.0	3292.8	46.0	3328.3	54.4	3328.3	54.4	97.2
Trial-1_0646	81	13864	0.7	18.6787	9.2	0.3108	11.4	0.0421	6.8	0.60	265.8	17.7	274.8	27.5	351.6	207.7	265.8	17.7	NA
Trial-1_0647	28	22184	0.4	9.8652	5.5	4.1218	7.1	0.2949	4.5	0.63	1666.0	65.9	1658.6	57.9	1649.3	101.4	1649.3	101.4	101.0
Trial-1_0648	49	53139	2.6	13.5336	5.0	1.7038	6.4	0.1672	4.0	0.63	996.9	36.7	1010.0	40.7	1038.5	100.3	1038.5	100.3	96.0
Trial-1_0649	74	18081	1.5	15.7072	5.8	0.9621	7.3	0.1096	4.5	0.61	670.4	28.4	684.4	36.5	730.5	123.1	670.4	28.4	91.8
Trial-1_0650	59	362267	2.0	13.1733	4.7	1.8417	6.0	0.1760	3.7	0.62	1044.8	35.9	1060.5	39.2	1092.7	93.3	1092.7	93.3	95.6
Trial-1_0651	81	7774	1.4	18.9543	9.3	0.3020	11.6	0.0415	6.9	0.60	262.3	17.7	268.0	27.3	318.4	211.7	262.3	17.7	NA
Trial-1_0652	32	33499	2.7	9.3259	4.8	4.3237	6.3	0.2924	4.0	0.64	1653.7	58.4	1697.9	51.8	1752.8	88.4	1752.8	88.4	94.3
Trial-1_0653	67	12121	1.7	18.7322	10.5	0.2799	13.1	0.0380	7.8	0.60	240.6	18.4	250.6	29.1	345.1	238.5	240.6	18.4	NA
Trial-1_0654	23	82605	1.1	4.9612	4.1	14.8758	5.4	0.5353	3.6	0.67	2763.6	81.4	2807.3	51.8	2838.9	66.3	2838.9	66.3	97.3
Trial-1_0655	25	8716	1.8	11.0539	3.6	2.8902	6.4	0.2317	5.3	0.83	1343.5	64.1	1379.4	48.1	1435.4	68.1	1435.4	68.1	93.6
Trial-1_0656	179	264478	45.1	12.7903	2.5	2.0008	3.2	0.1856	2.0	0.63	1097.5	20.5	1115.8	22.0	1151.5	50.2	1151.5	50.2	95.3
Trial-1_0657	61	44164	1.7	13.6361	4.8	1.7205	6.2	0.1702	3.8	0.62	1013.0	35.9	1016.2	39.6	1023.2	97.8	1023.2	97.8	99.0
Trial-1_0658	91	1085420	1.5	12.5579	3.9	2.0503	5.0	0.1867	3.1	0.62	1103.7	31.4	1132.4	34.0	1187.9	77.0	1187.9	77.0	92.9
Trial-1_0659	691	499778	4.3	12.9816	1.4	1.9729	1.8	0.1858	1.1	0.62	1098.3	11.1	1106.3	11.8	1122.0	27.4	1122.0	27.4	97.9
Trial-1_0660	135	2673346	1.5	8.9439	2.2	4.7868	2.9	0.3105	1.9	0.64	1743.2	28.5	1782.6	24.4	1829.0	40.4	1829.0	40.4	95.3
Trial-1_0661	46	24961	2.2	13.0648	5.6	1.9139	7.1	0.1813	4.4	0.62	1074.3	43.9	1085.9	47.6	1109.3	111.8	1109.3	111.8	96.9
Trial-1_0662	30	7047	1.6	16.6079	10.0	0.7724	12.5	0.0930	7.6	0.61	573.5	41.7	581.1	55.6	611.2	216.2	573.5	41.7	93.8
Trial-1_0663	41	40849	0.9	12.3952	5.6	2.2522	7.2	0.2025	4.5	0.62	1188.6	48.8	1197.5	50.7	1213.6	111.0	1213.6	111.0	97.9
Trial-1_0664	144	51208	1.5	12.7677	3.1	2.0681	3.9	0.1915	2.4	0.62	1129.5	25.3	1138.3	26.9	1155.1	61.1	1155.1	61.1	97.8
Trial-1_0665	33	22736	0.7	9.8634	5.1	3.9976	6.7	0.2860	4.2	0.64	1621.4	60.7	1633.7	54.1	1649.6	95.3	1649.6	95.3	98.3
Trial-1_0666	152	63532	5.5	12.7184	3.0	2.0230	3.8	0.1866	2.4	0.62	1102.9	24.3	1123.3	26.2	1162.7	59.8	1162.7	59.8	94.9
Trial-1_0667	77	164548	2.0	8.7844	3.1	5.1544	4.0	0.3284	2.6	0.64	1830.6	40.8	1845.1	33.9	1861.6	55.3	1861.6	55.3	98.3
Trial-1_0668	57	33268	5.0	12.0286	4.6	2.4285	5.9	0.2119	3.7	0.63	1238.7	41.6	1251.1	42.4	1272.4	89.8	1272.4	89.8	97.4
Trial-1_0669	54	339099	1.2	8.3003	3.5	5.8933	4.6	0.3548	3.0	0.64	1957.4	49.9	1960.2	39.8	1963.3	62.6	1963.3	62.6	99.7
Trial-1_0670	46	50126	0.6	9.1367	4.0	4.4108	5.2	0.2923	3.3	0.64	1652.9	48.4	1714.4	42.9	1790.2	72.6	1790.2	72.6	92.3
Trial-1_0671	330	390970	3.7	9.8268	1.6	3.8772	2.1	0.2763	1.3	0.63	1572.9	18.8	1609.0	17.2	1656.5	30.5	1656.5	30.5	95.0
Trial-1_0672	89	92715	1.4	10.8763	3.3	3.2409	4.3	0.2556	2.7	0.63	1467.5	35.5	1467.0	33.2	1466.2	63.1	1466.2	63.1	100.1
Trial-1_0673	64	74195	1.3	9.2697	3.2	4.3970	4.1	0.2956	2.7	0.64	1669.5	39.2	1711.8	34.3	1763.9	58.0	1763.9	58.0	94.6
Trial-1_0674	213	44373	2.3	15.6379	3.3	1.0327	4.2	0.1171	2.5	0.61	714.0	17.1	720.3	21.4	739.9	69.7	714.0	17.1	96.5
Trial-1_0675	26	34499	1.4	13.2137	7.5	1.8741	9.6	0.1796	6.0	0.62	1064.8	58.4	1072.0	63.6	1086.6	151.1	1086.6	151.1	98.0
Trial-1_0676	20	33535	1.3	4.5121	4.1	17.6905	5.6	0.5789	3.7	0.67	2944.3	88.2	2973.1	53.7	2992.5	66.8	2992.5	66.8	98.4
Trial-1_0677	20	19630	0.9	7.6236	5.5	6.9216	7.2	0.3827	4.7	0.65	2088.9	83.4	2101.4	64.3	2113.7	97.1	2113.7	97.1	98.8
Trial-1_0678	150	144815	1.5	12.6551	2.9	2.0951	3.8	0.1923	2.3	0.62	1133.8	24.4	1147.2	25.9	1172.7	58.4	1172.7	58.4	96.7
Trial-1_0679	69	22992	0.8	15.5815	5.9	1.0591	7.4	0.1197	4.5	0.61	728.8	31.1	733.4	38.7	747.5	124.1	728.8	31.1	97.5
Trial-1_0680	80	62637	0.7	12.7025	4.1	2.1548	5.2	0.1985	3.2	0.62	1167.4	34.7	1166.6	36.2	1165.2	81.0	1165.2	81.0	100.2
Trial-1_0681	46	19453	2.4	13.3677	5.8	1.7978	7.4	0.1743	4.6	0.62	1035.8	43.7	1044.7	48.1	1063.3	116.5	1063.3	116.5	97.4
Trial-1_0682	82	169655	2.1	12.8682	4.0	1.9431	5.1	0.1813	3.2	0.62	1074.3	31.5	1096.1	34.3	1139.5	79.6	1139.5	79.6	94.3
Trial-1_0683	141	82380	3.1	13.2989	2.9	1.7826	3.7	0.1719	2.3	0.62	1022.8	22.1	1039.1	24.3	1073.7	58.8	1073.7	58.8	95.3

Trial-1_0684	225	235654	0.9	8.8595	1.7	4.8942	2.3	0.3145	1.5	0.64	1762.7	22.5	1801.3	19.1	1846.2	31.4	1846.2	31.4	95.5
Trial-1_0685	84	34123	0.9	12.8473	4.2	1.9592	5.3	0.1825	3.3	0.62	1080.9	33.0	1101.6	35.8	1142.7	83.0	1142.7	83.0	94.6
Trial-1_0686	239	136462	3.3	10.0021	2.0	3.6160	2.6	0.2623	1.6	0.63	1501.7	22.0	1553.1	20.6	1623.7	37.3	1623.7	37.3	92.5
Trial-1_0687	51	52985	1.5	16.1256	7.8	0.7940	9.8	0.0929	5.9	0.61	572.4	32.5	593.4	44.0	674.6	166.5	572.4	32.5	84.9
Trial-1_0688	33	102479	2.1	10.8934	5.4	3.0450	7.0	0.2406	4.4	0.63	1389.7	55.0	1419.0	53.3	1463.2	102.9	1463.2	102.9	95.0
Trial-1_0689	78	470670	1.8	5.0627	2.2	14.7558	3.0	0.5418	2.0	0.66	2791.0	44.6	2799.6	28.2	2805.8	36.2	2805.8	36.2	99.5
Trial-1_0690	65	51746	1.5	12.5636	4.4	2.1453	5.7	0.1955	3.5	0.62	1151.0	37.4	1163.5	39.4	1187.0	87.9	1187.0	87.9	97.0
Trial-1_0691	43	17592	3.0	13.4022	5.3	1.7430	6.8	0.1694	4.3	0.62	1008.9	39.7	1024.6	43.9	1058.2	107.1	1058.2	107.1	95.3
Trial-1_0692	69	50180	1.8	8.3971	3.0	5.5976	3.9	0.3409	2.5	0.64	1891.0	41.4	1915.7	33.8	1942.6	53.7	1942.6	53.7	97.3
Trial-1_0693	108	47728	1.6	9.6819	2.9	4.0784	3.7	0.2864	2.3	0.63	1623.4	33.5	1650.0	30.1	1684.0	52.7	1684.0	52.7	96.4
Trial-1_0694	31	11293	1.0	12.7651	6.2	2.0250	7.9	0.1875	4.9	0.63	1107.7	50.3	1123.9	53.8	1155.5	122.7	1155.5	122.7	95.9
Trial-1_0695	51	17912	3.1	13.0839	5.1	1.9002	6.5	0.1803	4.0	0.62	1068.7	39.8	1081.2	43.3	1106.4	101.7	1106.4	101.7	96.6
Trial-1_0696	62	53666	1.5	8.3513	3.2	5.7785	4.1	0.3500	2.7	0.64	1934.6	44.4	1943.2	35.7	1952.4	56.3	1952.4	56.3	99.1
Trial-1_0697	20	38604	0.8	12.6037	8.1	2.1087	10.4	0.1928	6.5	0.62	1136.3	67.4	1151.7	71.7	1180.7	161.0	1180.7	161.0	96.2
Trial-1_0698	116	98203	2.4	8.7287	2.4	5.2948	3.1	0.3352	2.0	0.64	1863.5	32.1	1868.0	26.3	1873.0	42.6	1873.0	42.6	99.5
Trial-1_0699	107	353879	2.5	15.5417	5.1	0.8878	6.4	0.1001	3.9	0.61	614.8	22.8	645.2	30.5	752.9	107.1	614.8	22.8	81.7
Trial-1_0700	350	183542	1.6	12.4694	2.0	2.2259	2.5	0.2013	1.6	0.62	1182.3	16.8	1189.2	17.5	1201.8	38.5	1201.8	38.5	98.4
Trial-1_0701	518	217381	1.6	16.8370	2.7	0.6248	3.3	0.0763	2.0	0.60	474.0	9.2	492.9	13.1	581.5	58.0	474.0	9.2	81.5
Trial-1_0702	44	57906	2.3	12.8094	5.5	2.0285	7.0	0.1884	4.4	0.62	1113.0	44.7	1125.1	47.8	1148.6	109.3	1148.6	109.3	96.9
Trial-1_0703	137	13050	2.1	20.1501	8.3	0.1969	10.3	0.0288	6.1	0.59	182.9	11.0	182.5	17.2	177.6	193.5	182.9	11.0	NA
Trial-1_0704	288	94651	38.7	12.4055	2.1	2.2032	2.7	0.1982	1.7	0.62	1165.8	18.2	1182.1	19.1	1212.0	42.1	1212.0	42.1	96.2
Trial-1_0705	103	56740	4.4	12.5599	3.8	1.8882	4.9	0.1720	3.1	0.62	1023.1	28.9	1076.9	32.6	1187.6	75.9	1187.6	75.9	86.2
Trial-1_0706	78	56541	2.8	12.4370	4.0	2.2209	5.1	0.2003	3.2	0.62	1177.1	33.9	1187.7	35.4	1207.0	77.9	1207.0	77.9	97.5
Trial-1_0707	48	33838	1.0	8.0077	3.5	6.2422	4.6	0.3625	3.0	0.65	1994.1	50.7	2010.4	40.1	2027.1	61.9	2027.1	61.9	98.4
Trial-1_0708	42	26058	2.9	13.3242	5.1	1.8626	6.5	0.1800	4.1	0.63	1067.0	40.0	1067.9	42.9	1069.9	101.8	1069.9	101.8	99.7
Trial-1_0709	27	16440	0.8	8.8776	4.7	4.9859	6.2	0.3210	4.0	0.64	1794.7	62.2	1816.9	52.1	1842.5	85.1	1842.5	85.1	97.4
Trial-1_0710	27	21751	1.4	13.3289	6.7	1.8111	8.6	0.1751	5.3	0.62	1040.1	51.4	1049.5	56.2	1069.2	135.0	1069.2	135.0	97.3
Trial-1_0711	21	16471	1.5	13.3550	7.3	1.8358	9.4	0.1778	5.9	0.63	1055.0	57.3	1058.4	61.9	1065.3	147.8	1065.3	147.8	99.0
Trial-1_0712	20	6726	2.9	12.9191	8.0	1.9970	10.2	0.1871	6.4	0.62	1105.7	64.7	1114.5	69.2	1131.6	159.1	1131.6	159.1	97.7
Trial-1_0713	83	39739	16.9	9.9072	3.3	3.7962	4.3	0.2728	2.7	0.63	1554.9	37.4	1591.9	34.4	1641.4	61.5	1641.4	61.5	94.7
Trial-1_0714	71	52077	2.4	12.9292	4.5	1.9229	5.8	0.1803	3.6	0.62	1068.7	35.4	1089.1	38.7	1130.1	90.3	1130.1	90.3	94.6
Trial-1_0715	65	28908	2.1	13.5831	4.3	1.7183	5.5	0.1693	3.4	0.63	1008.1	32.0	1015.4	35.2	1031.1	86.4	1031.1	86.4	97.8
Trial-1_0716	24	3680	1.9	20.4946	16.8	0.3006	17.1	0.0447	3.4	0.20	281.8	9.4	266.9	40.2	137.9	396.4	281.8	9.4	NA
Trial-1_0717	114	8741	0.9	19.1998	7.7	0.3090	9.6	0.0430	5.7	0.60	271.5	15.2	273.4	23.0	289.1	175.8	271.5	15.2	NA
Trial-1_0718	27	21639	1.9	9.7415	5.7	3.9187	7.4	0.2769	4.7	0.63	1575.5	65.1	1617.5	59.7	1672.6	105.7	1672.6	105.7	94.2
Trial-1_0719	143	9609	1.6	18.6815	7.0	0.3021	8.8	0.0409	5.2	0.60	258.6	13.3	268.0	20.7	351.3	159.4	258.6	13.3	NA
Trial-1_0720	269	36263	1.3	17.6806	3.7	0.5368	4.6	0.0688	2.8	0.60	429.1	11.6	436.3	16.4	474.4	81.5	429.1	11.6	90.5
Trial-1_0721	63	275095	2.9	10.9301	3.9	3.2177	5.1	0.2551	3.2	0.63	1464.6	42.0	1461.4	39.4	1456.8	75.0	1456.8	75.0	100.5
Trial-1_0722	20	108888	1.4	12.8993	8.4	2.0355	10.8	0.1904	6.7	0.62	1123.7	69.1	1127.4	73.4	1134.7	168.1	1134.7	168.1	99.0
Trial-1_0723	70	69547	2.7	8.6885	3.1	5.1157	4.0	0.3224	2.6	0.64	1801.3	40.6	1838.7	34.1	1881.4	55.4	1881.4	55.4	95.7
Trial-1_0724	61	52664	1.1	12.9555	4.3	1.9067	5.5	0.1792	3.5	0.63	1062.4	34.0	1083.4	37.0	1126.0	86.2	1126.0	86.2	94.3
Trial-1_0725	52	44655	1.3	9.8669	4.2	3.6850	5.5	0.2637	3.5	0.63	1508.8	46.6	1568.1	43.7	1649.0	78.5	1649.0	78.5	91.5
Trial-1_0726	134	16220	2.3	17.2506	5.4	0.6046	6.7	0.0756	4.0	0.60	470.1	18.2	480.2	25.6	528.6	117.4	470.1	18.2	88.9
Trial-1_0727	15	8576	1.5	13.2528	3.0	1.8248	7.8	0.1754	7.2	0.92	1041.8	69.1	1054.4	51.2	1080.7	60.8	1080.7	60.8	96.4
Trial-1_0728	304	320224	3.3	12.7835	1.5	1.8935	1.9	0.1756	1.2	0.62	1042.6	11.2	1078.8	12.4	1152.6	29.1	1152.6	29.1	90.5
Trial-1_0729	57	91724	2.0	9.7842	2.6	3.7867	3.4	0.2687	2.2	0.64	1534.3	29.4	1589.9	27.2	1664.5	48.4	1664.5	48.4	92.2
Trial-1_0730	56	141650	1.7	13.4213	3.6	1.8178	4.6	0.1769	2.8	0.62	1050.2	27.4	1051.9	30.0	1055.3	72.4	1055.3	72.4	99.5
Trial-1_0731	166	161811	4.9	18.4728	3.6	0.4516	4.4	0.0605	2.7	0.60	378.7	9.8	378.4	14.1	376.6	80.1	378.7	9.8	NA
Trial-1_0732	57	137447	1.5	12.7574	3.2	2.1574	4.1	0.1996	2.5	0.62	1173.2	27.3	1167.4	28.3	1156.6	63.3	1156.6	63.3	101.4
Trial-1_0733	18	45660	1.5	13.7134	6.2	1.7069	7.9	0.1698	4.9	0.62	1010.8	45.4	1011.1	50.3	1011.7	125.2	1011.7	125.2	99.9
Trial-1_0734	203	191408	1.1	19.3929	3.9	0.2827	4.9	0.0398	2.9	0.60	251.4	7.2	252.8	11.0	266.2	90.4	251.4	7.2	NA
Trial-1_0735	93	206349	1.4	18.2682	4.6	0.4859	5.8	0.0644	3.5	0.60	402.2	13.5	402.1	19.1	401.6	103.3	402.2	13.5	100.2
Trial-1_0736	73	59925	2.9	13.6581	3.1	1.6758	4.0	0.1660	2.4	0.62	990.1	22.5	999.4	25.1	1019.9	62.9	1019.9	62.9	97.1
Trial-1_0737	43	187624	1.1	13.1727	3.8	1.9081	4.9	0.1823	3.0	0.62	1079.5	30.1	1083.9	32.5	1092.8	76.6	1092.8	76.6	98.8
Trial-1_0738	98	26025	1.1	18.1538	4.4	0.4929	5.6	0.0649	3.3	0.60	405.3	13.1	406.9	18.7	415.7	99.4	405.3	13.1	97.5
Trial-1_0739	124	269137	1.8	12.9627	2.3	1.9685	2.9	0.1851	1.8	0.62	1094.6	18.1	1104.8	19.5	1124.9	45.2	1124.9	45.2	97.3
Trial-1_0740	167	1072017	2.4	5.6625	1.1	11.4083	1.5	0.4685	1.0	0.66	2477.1	20.5	2557.1	14.1	2621.2	18.9	2621.2	18.9	94.5
Trial-1_0741	68	144263	1.1	13.4481	3.1	1.8188	3.9	0.1774	2.4	0.62	1052.7	23.7	1052.2	25.8	1051.3	62.2	1051.3	62.2	100.1
Trial-1_0742	108	141737	2.6	13.8448	2.5	1.6506	3.2	0.1657	2.0	0.62	988.6	18.0	989.8	20.1	992.4	50.7	992.4	50.7	99.6
Trial-1_0743	60	36963	0.9	17.9642	5.1	0.6326	6.4	0.0824	3.8	0.60	510.5	18.8	497.7	25.0	439.1	113.0	510.5	18.8	116.3
Trial-1_0744	48	43795	1.6	13.0588	3.6	2.0124	4.6	0.1906	2.9	0.62	1124.6	29.5	1119.7	31.1	1110.2	71.8	1110.2	71.8	101.3
Trial-1_0745	237	743034	1.8	11.0118	1.4	3.0719	1.8	0.2453	1.1	0.63	1414.4	14.4	1425.7	13.8	1442.6	26.5	1442.6	26.5	98.0
Trial-1_0746	62	83110	2.1	13.1248	3.2	2.0093	4.0	0.1913	2.5	0.62	1128.2	26.0	1118.6	27.4	1100.1	63.4	1100.1	63.4	102.6
Trial-1_0747	54	1091649	1.6	9.7742	2.6	4.1190	3.4	0.2920	2.2	0.64	1651.5	31.7	1658.1	27.9	1666.4	48.8	1666.4	48.8	99.1
Trial-1_0748	23	35812	0.3	5.3226	2.8	13.6728	3.7	0.5278	2.5	0.66	2732.2	55.3	2727.3	35.5	2723.7	46.2	2723.7	46.2	100.3
Trial-1_0749	34	156640	1.4	9.7621	3.4	3.8834	4.4	0.2750	2.8	0.64	1565.9	38.8	1610.2	35.5	1668.7	62.8	1668.7	62.8	93.8
Trial-1_0750	70	757084	2.5	5.3096	1.7	12.7743	2.2	0.4919	1.5	0.66	2579.0	31.5	2663.1	21.0	2727.7	27.5	2727.7	27.5	94.5
Trial-1_0751	111	94528	1.7	13.6369	2.6	1.5701	3.3	0.1553	2.1	0.62	930.5	17.8	958.5	20.7	1023.1	53.1	1023.1	53.1	91.0
Trial-1_0752	130	370452	1.0	6.0140	1.3	10.1898	1.7	0.4445	1.1	0.66	2370.6	22.5	2452.2	16.0	2520.5	21.8	2520.5	21.8	94.1
Trial-1_0753	23	110018	2.6	13.3697	5.2	1.9137	6.7	0.1856	4.2	0.62	1097.3	42.0	1085.9	44.7	1063.0	105.6	1063.0	105.6	103.2
Trial-1_0754	24	48457	1.6	13.4334	5.4	1.7611	6.8	0.1716	4.2	0.62	1020.8	39.9	1031.2	44.3	1053.5	108.2	1053.5	108.2	96.9
Trial-1_0755	191	134009	1.1	20.5310	5.1	0.1783	6.4	0.0266	3.8	0.59	169.0	6.3	166.6	9.8	133.7	121.1	169.0	6.3	NA
Trial-1_0756	359	801256	25.1	11.9739	1.3	2.3094	1.6	0.2006	1.0	0.63	1178.3	10.9	1215.2	11.5	1281.3	24.6	1281.3	24.6	92.0
Trial-1_0757	77	79098	1.1	13.5042	2.9	1.8266	3.7	0.1789	2.3	0.62	1061.0	22.4	1055.1	24.2	1042.8	58.4	1042.8	58.4	101.7
Trial-1_0758	288	263575	2.7	18.1414	2.5	0.5151	3.2	0.0678	1.9	0.60	422.7	7.8	421.9	10.9	417.2	56.6	422.7	7.8	101.3
Trial-1_0759	73	42170	3.0	19.4339	6.1	0.3421	7.5	0.0482	4.5	0.60	303.6	13.3	298.8	19.5	261.4	139.2	303.6	13.3	NA
Trial-1_0760	164	55056	1.5	20.1029	4.8	0.2806	5.9	0.0409	3.5	0.59	258.4	8.9	251.1	13.2	183.0	110.9	258.4	8.9	NA
Trial-1_0761	65	205906	3.6	5.3229	1.7	13.3019	2.3	0.5135	1.5	0.66	2671.6	32.8	2701.3	21.3	2723.6	27.8	2723.6	27.8	98.1
Trial-1_0762	139	77656	0.4	18.7244	4.4	0.3009	5.6	0.0409	3.4	0.62	258.2	8.7	267.1	13.1	346.1	99.3	258.2	8.7	NA
Trial-1_0763	61	68240	10.7	13.5062	3.4	1.7352	4.3	0.1700	2.7	0.62	1012.0	25.0	1021.7	27.7	1042.5	68.2	1042.5	68.2	97.1
Trial-1_0764	142	163687	0.6	12.9732	2.1	1.9716	2.7	0.1855	1.7	0.62	1097.0	16.7	1105.9	18.0	1123.3	41.6	1123.3	41.6	97.7
Trial-1_0765	219	174848	1.5	17.4035	2.6	0.6661	3.3	0.0841	2.0	0.60	520.4	9.8	518.3	13.2	509.2	57.2	520.4	9.8	102.2
Trial-1_0766	80	86756	1.1	16.4218	3.9	0.8534	4.9	0.1016	3.0	0.61	624.1	17.6	626.5	22.8	635.5	83.5	624.1	17.6	98.2
Trial-1_0767	132	188482	1.2	13.2844	2.2	1.8785	2.8	0.1810	1.8	0.62	1072.4	17.3	1073.5	18.7	1075.9	44.5	1075.9	44.5	99.7
Trial-1_0768	444	961518	4.7	16.5587	1.8	0.7009	2.3	0.0842	1.4	0.61	521.0	7.0	539.3	9.6	617.6	39.5	521.0	7.0	84.4
Trial-1_0769	68	156304	1.4	11.0566	2.6	3.0494	3.3	0.2445	2.1	0.63	1410.2	26.4	1420.1	25.3	1434.9	49.0	1434.9	49.0	98.3
Trial-1_0770	99	211684	1.7	9.2027	1.9	4.7206	2.4	0.3151	1.5	0.64	1765.6	23.9	1770.9	20.2	1777.1	33.8	1777.1	33.8	99.4
Trial-1_0771	64	45180	0.4	17.8055	4.8	0.6430	6.0	0.0830	3.6	0.60	514.2	18.0	504.1	24.0	458.8	106.8	514.2	18.0	112.1
Trial-1_0772	154	581446	1.2	9.7156	1.7	4.0493	2.2	0.2853	1.4	0.63	1618.1	19.9	1644.2	17.9	1677.5	31.6	1677.5	31.6	96.5
Trial-1_0773	225	326701	1.8	13.2345	1.8	1.7247	2.2	0.1655	1.4	0.62	987.5	12.7	1017.8	14.4	1083.4	35.2	1083.4	35.2	91.1
Trial-1_0774	247	359027	1.6	13.3174	1.7	1.7256	2.2	0.1667	1.3	0.62	993.8	12.3	1018.1	13.9	1070.9	34.1	1070.9	34.1	92.8
Trial-1_0775	100	127866	1.2	10.9114	2.1	3.2442	2.6	0.2567	1.7	0.63	1473.1	22.1	1467.8	20.6	1460.1	39.0	1460.1	39.0	100.9
Trial-1_0776	517	45778	1.1	15.8609	1.3	1.0421	1.6	0.1199	1.0	0.62	729.8	7.1	724.9	8.5	709.9	27.5	729.8	7.1	102.8
Trial-1_0777	162	150854	2.7	13.2514	2.0	1.7978	2.6	0.1728	1.6	0.62	1027.4	15.2	1044.7	16.8	1080.9	40.6	1080.9	40.6	95.1
Trial-1_0778	120	122749	2.7	19.8467	5.6	0.2724	7.0	0.0392	4.1	0.59	248.0	10.1	244.6	15.2	212.8	130.2	248.0	10.1	NA
Trial-1_0779	298	736888	2.5	13.7716	1.7	1.4053	2.1	0.1404	1.3	0.62	846.7	10.4	891.2	12.7	1003.2	34.0	1003.2	34.0	84.4
Trial-1_0780	635	1852251	3.2	16.7711	1.6	0.6634	2.0	0.0807	1.2	0.61	500.3	5.7	516.7	7.9	590.0	33.7	500.3	5.7	84.8
Trial-1_0781	29	89656	1.3	12.8619	4.5	2.0666	5.8	0.1928	3.6	0.62	1136.4	37.4	1137.8	39.4	1140.5	89.6	1140.5	89.6	99.6
Trial-1_0782	570	242725	4.6	16.5242	1.6	0.7248	2.0	0.0869	1.2	0.61	537.0	6.1	553.5	8.4	622.1	33.7	537.0	6.1	86.3
Trial-1_0783	174	333030	2.4	13.5323	2.0	1.6860	2.6	0.1655	1.6	0.62	987.1	14.6	1003.2	16.4	1038.6	40.9	1038.6	40.9	95.0
Trial-1_0784	28	156866	2.8	13.4281	4.7	1.8530	6.0	0.1805	3.7	0.62	1069.5	36.9	1064.5	39.8	1054.3	95.4	1054.3	95.4	101.4
Trial-1_0785	177	448758	1.4	10.8771	1.6	3.0322	2.1	0.2392	1.3	0.63	1382.6	16.5	1415.8	16.1	1466.1	31.0	1466.1	31.0	94.3
Trial-1_0786	25	29327	1.3	13.2817	5.0	1.8270	6.4	0.1760	4.0	0.62	1045.0	38.3	1055.2	42.0	1076.3	100.6	1076.3	100.6	97.1
Trial-1_0787	68	102101	2.0	13.3142	3.0	1.8698	3.9	0.1806	2.4	0.62	1070.0	23.6	1070.5	25.5	1071.4	60.7	1071.4	60.7	99.9
Trial-1_0788	93	124146	1.9	13.3321	2.6	1.8563	3.3	0.1795	2.1	0.62	1064.2	20.3	1065.7	22.0	1068.7	52.5	1068.7	52.5	99.6
Trial-1_0789	165	234627	1.5	11.0240	1.7	2.8522	2.2	0.2280	1.4	0.63	1324.3	16.8	1369.4	16.8	1440.5	33.0	1440.5	33.0	91.9
Trial-1_0790	92	103351	1.1	11.0961	2.2	2.9450	2.9	0.2370	1.8	0.63	1371.1	22.5	1393.6	21.9	1428.1	42.8	1428.1	42.8	96.0
Trial-1_0791	20	40439	1.5	9.3248	3.8	4.5211	5.0	0.3058	3.2	0.65	1719.8	48.4	1734.9	41.3	1753.0	69.4	1753.0	69.4	98.1
Trial-1_0792	131	252276	0.4	12.3601	2.1	2.2411	2.7	0.2009	1.7	0.63	1180.1	18.2	1194.0	18.9	1219.2	41.4	1219.2	41.4	96.8
Trial-1_0793	118	271114	3.0	12.7658	2.3	2.0390	2.9	0.1888	1.8	0.62	1114.8	18.6	1128.6	19.9	1155.3	45.3	1155.3	45.3	96.5
Trial-1_0794	21	25860	0.6	10.9240	4.5	3.0694	5.8	0.2432	3.7	0.63	1403.2	46.3	1425.1	44.5	1457.9	85.7	1457.9	85.7	96.3
Trial-1_0795	53	177397	0.6	10.8965	3.0	3.0539	3.8	0.2413	2.4	0.63	1393.7	30.2	1421.2	29.2	1462.7	56.2	1462.7	56.2	95.3
Trial-1_0796	44	52578	1.8	9.7315	2.9	4.1250	3.8	0.2911	2.4	0.64	1647.2	35.3	1659.3	31.2	1674.5	54.5	1674.5	54.5	98.4
Trial-1_0797	76	867784	1.3	13.1299	2.9	1.9601	3.7	0.1867	2.3	0.62	1103.3	23.3	1101.9	24.9	1099.3	58.0	1099.3	58.0	100.4
Trial-1_0798	85	54867	5.0	13.6707	3.0	1.5384	3.8	0.1525	2.4	0.62	915.1	20.1	945.9	23.5	1018.1	60.9	1018.1	60.9	89.9
Trial-1_0799	276	170573	2.6	18.9138	2.9	0.3959	3.6	0.0543	2.2	0.60	340.9	7.2	338.7	10.4	323.3	65.7	340.9	7.2	NA
Trial-1_0800	253	239930	1.2	18.2606	2.8	0.4990	3.5	0.0661	2.1	0.60	412.5	8.3	411.0	11.7	402.5	62.1	412.5	8.3	102.5
Trial-1_0801	49	57381	1.5	11.2153	3.0	3.0901	3.9	0.2513	2.4	0.63	1445.4	31.5	1430.2	29.6	1407.7	57.2	1407.7	57.2	102.7
Trial-1_0802	59	36062	1.6	13.3862	3.3	1.7739	4.3	0.1722	2.6	0.62	1024.3	25.0	1035.9	27.7	1060.6	67.3	1060.6	67.3	96.6
Trial-1_0803	125	50932	0.8	19.8859	5.3	0.2081	6.8	0.0300	4.3	0.63	190.6	8.0	191.9	12.0	208.3	123.5	190.6	8.0	NA
Trial-1_0804	113	349713	1.9	12.3068	2.2	2.2422	2.8	0.2001	1.8	0.63	1176.0	18.8	1194.4	19.7	1227.6	42.9	1227.6	42.9	95.8
Trial-1_0805	200	583664	1.3	16.6146	2.5	0.8335	3.1	0.1004	1.9	0.61	617.0	11.2	615.5	14.4	610.3	53.7	617.0	11.2	101.1
Trial-1_0806	98	159591	2.5	12.8394	2.4	2.0207	3.1	0.1882	1.9	0.62	1111.4	19.6	1122.5	20.9	1143.9	47.7	1143.9	47.7	97.2
Trial-1_0807	19	19344	1.4	13.4714	5.6	1.7654	7.1	0.1725	4.4	0.62	1025.8	42.1	1032.8	46.3	1047.8	112.9	1047.8	112.9	97.9
Trial-1_0808	482	160335	1.4	19.6335	2.6	0.2879	3.2	0.0410	1.9	0.60	259.0	4.9	256.9	7.3	237.8	59.9	259.0	4.9	NA
Trial-1_0809	98	140819	3.1	13.5483	2.6	1.7580	3.3	0.1727	2.1	0.62	1027.2	19.6	1030.1	21.5	1036.3	52.8	1036.3	52.8	99.1
Trial-1_0810	123	140187	2.8	13.3054	2.3	1.8195	2.9	0.1756	1.8	0.62	1042.8	17.3	1052.5	19.0	1072.7	45.7	1072.7	45.7	97.2
Trial-1_0811	60	69216	2.3	19.0013	5.6	0.4887	7.0	0.0674	4.2	0.60	420.2	17.0	404.0	23.3	312.8	127.2	420.2	17.0	134.3
Trial-1_0812	255	467682	2.5	10.9153	1.3	3.0692	1.7	0.2430	1.1	0.63	1402.1	13.5	1425.0	12.9	1459.4	24.9	1459.4	24.9	96.1
Trial-1_0813	59	72087	1.7	10.9824	2.8	3.1593	3.5	0.2516	2.2	0.63	1446.9	29.0	1447.3	27.4	1447.7	52.4	1447.7	52.4	99.9
Trial-1_0814	102	86422	2.5	10.9550	2.1	3.1756	2.7	0.2523	1.7	0.63	1450.4	22.3	1451.3	21.0	1452.5	40.0	1452.5	40.0	99.9
Trial-1_0815	27	842624	1.1	9.3814	3.2	4.3440	4.2	0.2956	2.7	0.65	1669.3	39.5	1701.8	34.3	1742.0	58.0	1742.0	58.0	95.8
Trial-1_0816	160	369153	2.3	13.5330	2.0	1.8033	2.6	0.1770	1.6	0.62	1050.5	15.5	1046.7	16.9	1038.5	40.9	1038.5	40.9	101.2
Trial-1_0817	350	314298	5.0	13.3028	1.4	1.7274	1.8	0.1667	1.1	0.62	993.7	10.2	1018.8	11.5	1073.1	28.1	1073.1	28.1	92.6
Trial-1_0818	43	165979	1.8	7.8575	2.5	6.4326	3.2	0.3666	2.1	0.65	2013.3	36.4	2036.7	28.5	2060.5	43.6	2060.5	43.6	97.7
Trial-1_0819	157	231182	2.4	12.5895	1.9	2.1233	2.5	0.1939	1.5	0.62	1142.3	16.1	1156.4	17.0	1182.9	38.0	1182.9	38.0	96.6
Trial-1_0820	67	266873	1.3	11.1399	2.7	3.1095	3.5	0.2512	2.2	0.63	1444.8	28.6	1435.0	26.9	1420.6	52.0	1420.6	52.0	101.7
Trial-1_0821	83	68883	2.3	13.1313	2.8	2.0457	3.6	0.1948	2.2	0.62	1147.5	23.6	1130.9	24.6	1099.1	56.6	1099.1	56.6	104.4

Trial-1_0822	132	206046	1.3	11.0267	2.0	3.0715	2.5	0.2456	1.6	0.63	1416.0	20.3	1425.6	19.4	1440.1	37.4	1440.1	37.4	98.3
Trial-1_0823	182	427992	1.2	10.9743	1.6	3.1100	2.1	0.2475	1.3	0.63	1425.7	17.2	1435.2	16.3	1449.1	31.4	1449.1	31.4	98.4
Trial-1_0824	355	1097110	2.5	10.9021	1.2	3.0688	1.5	0.2427	1.0	0.63	1400.5	12.2	1424.9	11.7	1461.7	22.5	1461.7	22.5	95.8
Trial-1_0825	79	17090	2.2	21.4168	11.8	0.0967	14.6	0.0150	8.6	0.59	96.1	8.2	93.7	13.1	33.5	283.4	96.1	8.2	NA
Trial-1_0826	177	92229	0.7	19.1825	3.9	0.3800	4.8	0.0529	2.9	0.60	332.1	9.4	327.0	13.5	291.1	88.7	332.1	9.4	NA
Trial-1_0827	256	281944	1.0	19.9140	3.9	0.2788	4.8	0.0403	2.9	0.59	254.5	7.2	249.7	10.7	205.0	90.5	254.5	7.2	NA
Trial-1_0828	20	19712	2.8	12.9523	5.8	1.9984	7.4	0.1877	4.6	0.62	1109.1	47.3	1115.0	50.4	1126.5	116.1	1126.5	116.1	98.5
Trial-1_0829	112	155571	2.8	13.6273	2.6	1.7221	3.4	0.1702	2.1	0.62	1013.2	19.5	1016.8	21.5	1024.5	53.3	1024.5	53.3	98.9
Trial-1_0830	98	192101	2.6	13.5082	2.7	1.7845	3.5	0.1748	2.2	0.62	1038.7	20.8	1039.8	22.7	1042.2	55.3	1042.2	55.3	99.7
Trial-1_0831	41	79950	2.9	12.0378	3.7	2.4910	4.8	0.2175	3.0	0.63	1268.5	34.3	1269.4	34.5	1270.9	72.3	1270.9	72.3	99.8
Trial-1_0832	19	30349	1.6	13.4961	6.1	1.8573	7.8	0.1818	4.9	0.62	1076.8	48.1	1066.0	51.6	1044.0	123.7	1044.0	123.7	103.1
Trial-1_0833	88	93758	1.8	13.3363	2.8	1.9076	3.6	0.1845	2.2	0.62	1091.6	22.4	1083.7	23.9	1068.1	56.5	1068.1	56.5	102.2
Trial-1_0834	104	32513	1.1	20.3898	5.9	0.2698	7.3	0.0399	4.3	0.59	252.2	10.7	242.6	15.7	149.9	137.5	252.2	10.7	NA
Trial-1_0835	179	196389	3.5	12.1911	1.8	2.3276	2.4	0.2058	1.5	0.63	1206.4	16.2	1220.8	16.7	1246.2	36.0	1246.2	36.0	96.8
Trial-1_0836	62	146772	3.2	9.3456	2.6	4.3130	3.3	0.2923	2.1	0.64	1653.2	31.0	1695.8	27.5	1749.0	46.9	1749.0	46.9	94.5
Trial-1_0837	534	376216	22.3	12.8589	1.2	1.8319	1.5	0.1708	0.9	0.62	1016.8	8.8	1057.0	9.9	1140.9	23.4	1140.9	23.4	89.1
Trial-1_0838	41	25219	0.9	12.6935	4.0	2.0930	5.1	0.1927	3.2	0.62	1135.9	32.9	1146.5	35.0	1166.6	79.1	1166.6	79.1	97.4
Trial-1_0839	67	21472	9.3	19.1869	5.9	0.4235	7.4	0.0589	4.4	0.60	369.1	15.7	358.6	22.2	290.6	135.0	369.1	15.7	NA
Trial-1_0840	122	27116	2.4	19.9229	5.5	0.2670	6.8	0.0386	4.1	0.59	244.0	9.7	240.3	14.6	203.9	127.7	244.0	9.7	NA
Trial-1_0841	84	83982	2.6	13.5116	3.0	1.8167	3.8	0.1780	2.3	0.62	1056.2	22.8	1051.5	24.7	1041.7	59.8	1041.7	59.8	101.4
Trial-1_0842	127	295292	2.1	11.0917	2.0	3.0756	2.5	0.2474	1.6	0.63	1425.1	20.5	1426.6	19.4	1428.9	37.5	1428.9	37.5	99.7
Trial-1_0843	329	261556	4.5	14.2055	1.6	1.4597	2.1	0.1504	1.3	0.62	903.2	10.7	913.9	12.4	939.9	33.1	939.9	33.1	96.1
Trial-1_0844	62	89006	6.8	9.1392	2.4	4.5450	3.1	0.3013	2.0	0.64	1697.5	30.1	1739.2	26.1	1789.8	43.9	1789.8	43.9	94.8
Trial-1_0845	100	79523	1.5	12.9242	2.6	1.9990	3.4	0.1874	2.1	0.62	1107.1	21.2	1115.2	22.7	1130.8	52.4	1130.8	52.4	97.9
Trial-1_0846	147	556723	0.3	8.3912	1.5	5.5464	2.0	0.3375	1.3	0.64	1874.9	20.7	1907.8	17.0	1943.8	27.0	1943.8	27.0	96.5
Trial-1_0847	53	109602	0.5	5.4967	2.0	12.7455	2.6	0.5081	1.8	0.66	2648.5	38.0	2661.0	24.9	2670.5	32.8	2670.5	32.8	99.2
Trial-1_0848	95	82125	1.6	13.4244	2.7	1.7641	3.5	0.1718	2.2	0.62	1021.8	20.5	1032.4	22.7	1054.8	55.3	1054.8	55.3	96.9
Trial-1_0849	136	410494	0.9	11.0274	1.9	3.1673	2.4	0.2533	1.5	0.63	1455.6	20.0	1449.2	18.7	1439.9	35.8	1439.9	35.8	101.1
Trial-1_0850	72	12042	2.4	20.1351	6.7	0.3001	8.3	0.0438	5.0	0.59	276.5	13.4	266.5	19.5	179.3	156.2	276.5	13.4	NA
Trial-1_0851	25	75307	2.0	5.4093	2.9	12.6941	3.8	0.4980	2.5	0.66	2605.3	54.2	2657.2	35.9	2697.0	47.1	2697.0	47.1	96.6
Trial-1_0852	30	56031	1.2	13.4753	4.8	1.7971	6.1	0.1756	3.8	0.62	1043.1	36.5	1044.4	39.9	1047.2	96.7	1047.2	96.7	99.6
Trial-1_0853	98	1122447	3.3	13.7388	2.7	1.6992	3.5	0.1693	2.2	0.62	1008.3	20.2	1008.2	22.4	1008.0	55.8	1008.0	55.8	100.0
Trial-1_0854	87	101695	1.9	13.0238	2.8	1.8956	3.6	0.1791	2.2	0.62	1061.8	21.7	1079.6	23.7	1115.5	55.8	1115.5	55.8	95.2
Trial-1_0855	174	330016	1.9	9.4663	1.5	4.2746	2.0	0.2935	1.3	0.64	1658.9	18.6	1688.5	16.4	1725.4	28.2	1725.4	28.2	96.1
Trial-1_0856	90	99294	1.7	10.9781	2.2	3.1168	2.9	0.2482	1.8	0.63	1429.0	23.3	1436.8	22.1	1448.5	42.3	1448.5	42.3	98.7
Trial-1_0857	124	97045	2.2	12.9364	2.3	2.0218	3.0	0.1897	1.8	0.62	1119.7	19.0	1122.9	20.2	1129.0	46.3	1129.0	46.3	99.2
Trial-1_0858	26	25079	1.3	13.4356	5.2	1.7789	6.6	0.1733	4.1	0.62	1030.5	39.1	1037.8	43.1	1053.1	104.8	1053.1	104.8	97.9
Trial-1_0859	192	46053	2.8	20.7466	5.3	0.1759	6.5	0.0265	3.9	0.59	168.4	6.4	164.6	10.0	109.1	124.8	168.4	6.4	NA
Trial-1_0860	73	65933	1.7	13.4783	3.1	1.7518	4.0	0.1712	2.5	0.62	1019.0	23.2	1027.8	25.7	1046.8	62.8	1046.8	62.8	97.3
Trial-1_0861	101	134846	2.2	13.1640	2.6	1.8860	3.3	0.1801	2.0	0.62	1067.3	20.0	1076.2	21.7	1094.2	51.3	1094.2	51.3	97.5
Trial-1_0862	105	125099	0.8	17.2918	3.8	0.7203	4.7	0.0903	2.9	0.61	557.5	15.3	550.9	20.1	523.3	82.6	557.5	15.3	106.5
Trial-1_0863	177	83976	15.6	19.2777	4.0	0.3573	5.0	0.0500	3.0	0.60	314.2	9.1	310.2	13.3	279.9	91.4	314.2	9.1	NA
Trial-1_0864	48	52839	1.6	11.0388	3.4	2.9376	4.3	0.2352	2.7	0.63	1361.6	33.3	1391.6	32.8	1438.0	64.2	1438.0	64.2	94.7
Trial-1_0865	75	8867	1.8	20.5897	7.0	0.2640	8.7	0.0394	5.1	0.59	249.2	12.6	237.9	18.4	127.1	164.3	249.2	12.6	NA
Trial-1_0866	41	149430	1.1	11.0988	3.5	3.0694	4.5	0.2471	2.8	0.63	1423.4	36.1	1425.1	34.4	1427.6	66.5	1427.6	66.5	99.7
Trial-1_0867	123	311997	7.8	13.8550	2.5	1.6633	3.2	0.1671	2.0	0.62	996.3	18.4	994.6	20.4	990.9	51.4	990.9	51.4	100.5
Trial-1_0868	277	368329	2.5	11.1351	1.4	2.6623	1.9	0.2150	1.2	0.63	1255.4	13.3	1318.0	13.7	1421.4	27.4	1421.4	27.4	88.3
Trial-1_0869	95	126932	1.4	13.6579	2.8	1.6935	3.6	0.1678	2.2	0.62	999.7	20.7	1006.1	23.1	1020.0	57.6	1020.0	57.6	98.0
Trial-1_0870	34	41726	2.3	13.3652	4.6	1.8282	5.9	0.1772	3.6	0.62	1051.7	35.2	1055.6	38.5	1063.7	92.6	1063.7	92.6	98.9
Trial-1_0871	15	14490	1.3	12.8493	6.6	2.0856	8.5	0.1944	5.3	0.62	1145.0	55.4	1144.1	58.3	1142.4	132.1	1142.4	132.1	100.2
Trial-1_0872	101	277006	4.1	13.1375	2.6	1.7363	3.3	0.1654	2.0	0.62	986.9	18.6	1022.1	21.1	1098.2	51.2	1098.2	51.2	89.9
Trial-1_0873	156	42896	0.4	18.8261	3.8	0.4905	4.8	0.0670	2.9	0.60	417.9	11.6	405.2	15.9	333.8	86.6	417.9	11.6	125.2
Trial-1_0874	82	206968	0.8	5.3890	1.6	12.6004	2.2	0.4925	1.4	0.66	2581.4	30.3	2650.2	20.2	2703.2	26.6	2703.2	26.6	95.5
Trial-1_0875	235	142980	0.7	20.6569	4.7	0.1797	5.8	0.0269	3.5	0.59	171.2	5.8	167.8	9.0	119.4	110.7	171.2	5.8	NA
Trial-1_0876	69	105047	0.9	8.7659	2.2	5.3659	2.9	0.3411	1.9	0.64	1892.2	30.8	1879.4	25.0	1865.3	40.4	1865.3	40.4	101.4
Trial-1_0877	168	405075	1.8	11.0279	1.7	3.0835	2.2	0.2466	1.4	0.63	1421.0	17.7	1428.6	16.9	1439.9	32.5	1439.9	32.5	98.7
Trial-1_0878	191	187845	2.7	13.3357	2.0	1.8581	2.5	0.1797	1.5	0.62	1065.4	15.2	1066.3	16.5	1068.2	39.4	1068.2	39.4	99.7
Trial-1_0879	119	26269	1.1	18.8105	4.2	0.4884	5.2	0.0666	3.1	0.60	415.9	12.6	403.8	17.4	335.7	94.8	415.9	12.6	123.9
Trial-1_0880	247	88046	2.1	19.0474	3.1	0.4366	3.9	0.0603	2.3	0.60	377.5	8.5	367.8	11.9	307.3	70.4	377.5	8.5	NA
Trial-1_0881	269	312725	9.7	13.4400	1.7	1.6833	2.2	0.1641	1.3	0.62	979.4	12.2	1002.2	13.8	1052.5	34.4	1052.5	34.4	93.1
Trial-1_0882	19	15587	2.0	9.3889	4.6	4.4356	5.9	0.3020	3.8	0.64	1701.4	56.7	1719.0	49.3	1740.5	84.0	1740.5	84.0	97.8
Trial-1_0883	29	51727	1.5	12.3896	4.5	2.4065	5.8	0.2162	3.6	0.63	1262.0	41.8	1244.5	41.8	1214.4	89.5	1214.4	89.5	103.9
Trial-1_0884	166	1552131	0.9	5.8953	1.2	10.7127	1.6	0.4580	1.1	0.66	2430.9	21.4	2498.5	14.9	2553.9	20.2	2553.9	20.2	95.2
Trial-1_0885	131	95777	1.4	12.8214	2.2	2.0891	2.9	0.1943	1.8	0.62	1144.4	18.7	1145.2	19.7	1146.7	44.6	1146.7	44.6	99.8
Trial-1_0886	136	59495	3.1	16.6689	3.1	0.8506	3.9	0.1028	2.4	0.61	631.0	14.3	625.0	18.4	603.3	67.6	631.0	14.3	104.6
Trial-1_0887	43	59068	2.4	12.9147	4.0	1.9877	5.1	0.1862	3.2	0.62	1100.6	32.3	1111.3	34.7	1132.3	80.2	1132.3	80.2	97.2
Trial-1_0888	299	1608230	3.1	9.4683	1.1	4.2967	1.5	0.2951	1.0	0.64	1666.7	14.0	1692.7	12.2	1725.0	20.9	1725.0	20.9	96.6
Trial-1_0889	621	434578	1.3	16.7959	1.8	0.5451	2.2	0.0664	1.3	0.61	414.5	5.4	441.8	7.9	586.9	38.3	414.5	5.4	70.6
Trial-1_0890	56	52832	1.8	13.1574	3.5	1.7770	4.4	0.1696	2.8	0.62	1009.8	25.8	1037.1	28.8	1095.2	69.6	1095.2	69.6	92.2
Trial-1_0891	45	158959	0.6	8.7100	2.8	5.3312	3.6	0.3368	2.3	0.64	1871.2	37.9	1873.9	31.1	1876.9	50.2	1876.9	50.2	99.7
Trial-1_0892	214	40487	1.3	20.6543	4.9	0.1815	6.1	0.0272	3.6	0.59	172.9	6.1	169.3	9.4	119.6	115.1	172.9	6.1	NA
Trial-1_0893	9	214566	1.0	13.5218	8.9	1.7413	11.4	0.1708	7.0	0.62	1016.3	66.3	1023.9	73.4	1040.2	180.3	1040.2	180.3	97.7
Trial-1_0894	101	118319	1.2	10.8026	2.1	3.3028	2.8	0.2588	1.7	0.63	1483.5	23.2	1481.7	21.6	1479.1	40.6	1479.1	40.6	100.3
Trial-1_0895	160	163647	3.1	12.9437	2.1	2.0136	2.6	0.1890	1.6	0.62	1116.1	16.8	1120.1	17.8	1127.9	41.0	1127.9	41.0	99.0
Trial-1_0896	134	196624	0.5	12.7897	2.2	2.0928	2.8	0.1941	1.8	0.62	1143.7	18.4	1146.4	19.4	1151.6	43.7	1151.6	43.7	99.3
Trial-1_0897	54	53599	1.8	13.3770	3.6	1.8684	4.6	0.1813	2.9	0.62	1073.9	28.4	1070.0	30.6	1061.9	73.1	1061.9	73.1	101.1
Trial-1_0898	139	115237	1.7	9.5856	1.7	4.3130	2.2	0.2998	1.4	0.64	1690.6	20.6	1695.9	17.9	1702.4	30.8	1702.4	30.8	99.3
Trial-1_0899	110	251642	1.5	8.6484	1.8	5.3934	2.3	0.3383	1.5	0.64	1878.5	24.2	1883.8	19.8	1889.7	31.9	1889.7	31.9	99.4
Trial-1_0900	10	17343	1.5	10.0191	3.5	4.0239	6.4	0.2924	5.3	0.84	1653.5	77.6	1639.0	51.7	1620.5	64.6	1620.5	64.6	102.0
Trial-1_0901	221	205441	3.1	13.0461	1.8	1.8909	2.3	0.1789	1.4	0.62	1061.1	13.7	1077.9	15.0	1112.1	35.3	1112.1	35.3	95.4
Trial-1_0902	65	83304	2.5	13.6624	3.4	1.7654	4.3	0.1749	2.6	0.62	1039.2	25.4	1032.8	27.7	1019.3	68.0	1019.3	68.0	102.0
Trial-1_0903	177	156397	2.5	13.8082	2.1	1.6018	2.7	0.1604	1.7	0.62	959.1	15.0	970.9	17.0	997.8	43.4	997.8	43.4	96.1
Trial-1_0904	39	38195	1.4	15.3627	5.3	1.0809	6.7	0.1204	4.1	0.61	733.1	28.5	744.1	35.5	777.3	112.0	733.1	28.5	94.3
Trial-1_0905	55	96584	1.7	11.0883	3.1	2.9408	3.9	0.2365	2.5	0.63	1368.5	30.5	1392.5	29.8	1429.4	58.4	1429.4	58.4	95.7
Trial-1_0906	64	205054	1.5	13.1032	3.2	1.9373	4.1	0.1841	2.5	0.62	1089.4	25.5	1094.1	27.4	1103.4	64.1	1103.4	64.1	98.7
Trial-1_0907	49	71935	1.1	8.6154	2.7	5.3227	3.5	0.3326	2.2	0.64	1850.9	36.0	1872.5	29.8	1896.6	48.0	1896.6	48.0	97.6
Trial-1_0908	162	387307	2.0	13.1499	2.1	1.7787	2.7	0.1696	1.7	0.62	1010.1	15.5	1037.7	17.4	1096.3	42.0	1096.3	42.0	92.1
Trial-1_0909	168	216184	2.1	9.1491	1.5	4.6146	2.0	0.3062	1.3	0.64	1722.0	19.0	1751.9	16.4	1787.8	27.4	1787.8	27.4	96.3
Trial-1_0910	13	71100	1.3	13.2360	7.5	1.9044	9.6	0.1828	5.9	0.62	1082.3	59.1	1082.6	64.0	1083.2	151.5	1083.2	151.5	99.9
Trial-1_0911	90	172542	1.5	13.5920	2.9	1.7947	3.6	0.1769	2.3	0.62	1050.1	21.8	1043.5	23.7	1029.7	57.6	1029.7	57.6	102.0
Trial-1_0912	48	278613	2.7	9.2188	2.6	4.7092	3.4	0.3149	2.2	0.64	1764.6	34.2	1768.9	28.9	1773.9	48.2	1773.9	48.2	99.5
Trial-1_0913	44	56985	2.3	13.4670	3.9	1.8050	5.0	0.1763	3.1	0.62	1046.7	29.9	1047.3	32.6	1048.4	78.9	1048.4	78.9	99.8
Trial-1_0914	75	77908	1.6	13.2985	3.4	1.5588	4.3	0.1503	2.7	0.62	902.9	22.5	954.0	26.7	1073.8	68.1	1073.8	68.1	84.1
Trial-1_0915	173	219599	1.7	9.5612	1.5	4.1518	2.0	0.2879	1.3	0.64	1631.0	18.5	1664.6	16.4	1707.1	28.5	1707.1	28.5	95.5
Trial-1_0916	223	170627	1.4	18.2203	3.0	0.5329	3.7	0.0704	2.2	0.60	438.7	9.5	433.7	13.1	407.5	66.3	438.7	9.5	107.7
Trial-1_0917	64	77717	2.1	10.9535	2.8	3.1131	3.6	0.2473	2.3	0.63	1424.6	29.0	1435.9	27.7	1452.7	53.1	1452.7	53.1	98.1
Trial-1_0918	21	184130	0.6	9.7347	4.6	3.8620	5.9	0.2727	3.8	0.63	1554.3	51.9	1605.8	47.9	1673.9	84.9	1673.9	84.9	92.9
Trial-1_0919	66	347418	3.2	13.3830	3.2	1.8240	4.1	0.1770	2.6	0.62	1050.8	24.8	1054.1	27.0	1061.0	65.0	1061.0	65.0	99.0
Trial-1_0920	81	94217	1.5	13.8821	3.1	1.6486	4.0	0.1660	2.5	0.62	989.9	22.6	989.0	25.2	986.9	63.8	986.9	63.8	100.3
Trial-1_0921	32	327772	2.8	13.3960	4.6	1.8396	5.9	0.1787	3.6	0.62	1060.1	35.6	1059.7	38.6	1059.1	92.6	1059.1	92.6	100.1
Trial-1_0922	54	89989	2.2	6.0841	2.1	10.1989	2.8	0.4500	1.9	0.66	2395.4	37.3	2453.0	26.3	2501.0	36.1	2501.0	36.1	95.8
Trial-1_0923	160	206398	3.4	13.4775	2.1	1.8179	2.7	0.1777	1.7	0.62	1054.4	16.3	1051.9	17.6	1046.9	42.6	1046.9	42.6	100.7
Trial-1_0924	20	14445	0.5	11.9201	5.3	2.5016	6.9	0.2163	4.3	0.63	1262.2	49.3	1272.5	49.8	1290.0	104.1	1290.0	104.1	97.8
Trial-1_0925	32	53650	0.7	10.9254	3.8	3.0731	4.9	0.2435	3.1	0.63	1404.9	38.9	1426.0	37.3	1457.6	71.8	1457.6	71.8	96.4
Trial-1_0926	61	156728	1.1	5.3739	1.8	13.1474	2.5	0.5124	1.6	0.66	2667.0	35.6	2690.3	23.2	2707.9	30.4	2707.9	30.4	98.5
Trial-1_0927	167	220761	1.7	13.0672	2.0	1.9429	2.6	0.1841	1.6	0.62	1089.5	16.1	1096.0	17.3	1108.9	40.5	1108.9	40.5	98.3
Trial-1_0928	67	44670	3.3	10.9892	2.7	3.0763	3.5	0.2452	2.2	0.63	1413.6	28.0	1426.8	26.9	1446.5	51.9	1446.5	51.9	97.7
Trial-1_0929	514	155846	4.4	16.7116	1.8	0.7058	2.3	0.0855	1.4	0.61	529.1	7.1	542.2	9.6	597.7	39.6	529.1	7.1	88.5
Trial-1_0930	76	307079	1.9	13.3208	3.1	1.8290	3.9	0.1767	2.4	0.62	1049.0	23.5	1055.9	25.7	1070.4	61.7	1070.4	61.7	98.0
Trial-1_0931	70	49657	1.3	13.0655	3.1	1.9290	4.0	0.1828	2.5	0.62	1082.2	24.9	1091.2	26.8	1109.2	62.8	1109.2	62.8	97.6
Trial-1_0932	59	145874	2.2	17.6926	5.3	0.6882	6.6	0.0883	4.0	0.60	545.6	20.8	531.7	27.4	472.9	116.7	545.6	20.8	115.4
Trial-1_0933	62	121074	0.5	5.1264	1.9	14.2814	2.5	0.5310	1.6	0.66	2745.6	36.8	2768.6	23.6	2785.3	30.4	2785.3	30.4	98.6
Trial-1_0934	176	448622	1.2	13.6072	2.1	1.7515	2.7	0.1728	1.7	0.62	1027.8	16.0	1027.7	17.6	1027.5	43.3	1027.5	43.3	100.0
Trial-1_0935	56	15710	1.0	18.9521	6.4	0.4957	8.0	0.0681	4.8	0.60	424.9	19.7	408.8	26.9	318.7	145.6	424.9	19.7	133.3
Trial-1_0936	109	122725	0.7	8.7522	1.8	5.2319	2.4	0.3321	1.5	0.64	1848.6	24.9	1857.8	20.6	1868.2	33.4	1868.2	33.4	99.0
Trial-1_0937	100	126666	1.3	8.6605	1.9	5.3293	2.5	0.3347	1.6	0.64	1861.3	26.4	1873.6	21.7	1887.2	35.1	1887.2	35.1	98.6
Trial-1_0938	15	25899	1.2	7.6778	4.5	6.8035	5.9	0.3788	3.8	0.65	2070.9	67.5	2086.2	52.1	2101.3	78.6	2101.3	78.6	98.6
Trial-1_0939	144	203471	1.5	12.6527	2.2	2.0997	2.9	0.1927	1.8	0.62	1135.9	18.5	1148.7	19.6	1173.0	44.2	1173.0	44.2	96.8
Trial-1_0940	122	227957	2.1	13.4152	2.5	1.7662	3.2	0.1718	2.0	0.62	1022.3	18.6	1033.1	20.6	1056.2	50.2	1056.2	50.2	96.8
Trial-1_0941	73	101529	1.0	11.1010	2.7	3.0806	3.5	0.2480	2.2	0.63	1428.3	28.3	1427.9	26.9	1427.2	52.1	1427.2	52.1	100.1
Trial-1_0942	42	10781	2.2	20.8839	9.6	0.2675	11.9	0.0405	7.0	0.59	256.1	17.7	240.7	25.6	93.6	228.3	256.1	17.7	NA
Trial-1_0943	21	46672	1.0	7.7896	3.9	6.6771	5.1	0.3772	3.3	0.65	2063.3	58.3	2069.6	45.0	2075.8	68.4	2075.8	68.4	99.4
Trial-1_0944	350	520529	6.7	13.4342	1.5	1.7188	2.0	0.1675	1.2	0.62	998.2	11.2	1015.6	12.6	1053.4	30.9	1053.4	30.9	94.8
Trial-1_0945	153	117408	1.6	16.1207	2.9	0.9895	3.6	0.1157	2.2	0.61	705.7	14.8	698.5	18.3	675.2	61.7	705.7	14.8	104.5
Trial-1_0946	128	221200	2.1	13.6943	2.6	1.6888	3.2	0.1677	2.0	0.62	999.6	18.6	1004.3	20.7	1014.6	51.7	1014.6	51.7	98.5
Trial-1_0947	58	1059172	0.8	11.1412	3.0	3.0573	3.8	0.2470	2.4	0.63	1423.2	30.6	1422.1	29.1	1420.3	56.5	1420.3	56.5	100.2
Trial-1_0948	31	21407	1.7	19.8795	11.1	0.2809	13.8	0.0405	8.2	0.59	255.9	20.5	251.4	30.7	209.0	258.3	255.9	20.5	NA
Trial-1_0949	12	12408	2.0	13.1835	8.0	1.9161	10.2	0.1832	6.3	0.62	1084.5	63.0	1086.7	68.2	1091.2	161.1	1091.2	161.1	99.4
Trial-1_0950	62	30841	0.8	11.0951	3.0	2.9379	3.9	0.2364	2.4	0.63	1368.0	30.1	1391.7	29.4	1428.3	57.7	1428.3	57.7	95.8
Trial-1_0951	372	82711	3.9	17.9030	2.5	0.5232	3.1	0.0679	1.8	0.60	423.7	7.6	427.3	10.7	446.6	54.6	423.7	7.6	94.9
Trial-1_0952	16	19208	1.6	13.2890	6.9	1.8312	8.8	0.1765	5.4	0.62	1047.8	52.5	1056.7	57.6	1075.2	138.2	1075.2	138.2	97.5
Trial-1_0953	34	34132	2.0	12.1212	4.3	2.3773	5.5	0.2090	3.5	0.63	1223.4	38.5	1235.8	39.5	1257.4	84.3	1257.4	84.3	97.3
Trial-1_0954	78	47542	1.5	13.6191	3.2	1.6750	4.0	0.1654	2.5	0.62	987.0	22.8	999.1	25.6	1025.7	64.1	1025.7	64.1	96.2
Trial-1_0955	36	152760	2.0	13.2765	4.5	1.8376	5.7	0.1769	3.6	0.62	1050.2	34.4	1059.0	37.7	1077.1	90.4	1077.1	90.4	97.5
Trial-1_0956	371	782402	7.1	13.5267	1.5	1.6382	1.9	0.1607	1.2	0.62	960.8	10.7	985.0	12.3	1039.5	30.9	1039.5	30.9	92.4
Trial-1_0957	72	277724	3.7	9.1651	2.3	4.6864	3.0	0.3115	1.9	0.64	1748.2	29.4	1764.8	25.1	1784.6	41.9	1784.6	41.9	98.0
Trial-1_0958	30	112207	0.9	13.5356	5.0	1.7627	6.4	0.1730	4.0	0.62	1028.9	37.8	1031.8	41.6	1038.1	101.7	1038.1	101.7	99.1
Trial-1_0959	30	49777	3.0	18.1711	8.3	0.5546	10.3	0.0731	6.2	0.60	454.8	27.2	448.0	37.4	413.5	184.8	454.8	27.2	110.0

Trial-1_0960	65	56392	1.3	16.7536	4.8	0.8250	6.0	0.1002	3.6	0.61	615.9	21.3	610.8	27.5	592.3	103.4	615.9	21.3	104.0
Trial-1_0961	87	40758	2.1	18.4498	4.9	0.5257	6.1	0.0703	3.6	0.60	438.2	15.4	429.0	21.3	379.4	109.4	438.2	15.4	115.5
Trial-1_0962	50	138906	1.7	13.2981	4.0	1.8754	5.0	0.1809	3.1	0.62	1071.8	30.8	1072.4	33.4	1073.8	79.4	1073.8	79.4	99.8
Trial-1_0963	127	380021	1.1	8.9959	1.8	4.8578	2.3	0.3169	1.5	0.64	1774.8	22.9	1795.0	19.4	1818.5	32.1	1818.5	32.1	97.6
Trial-1_0964	53	130332	1.6	12.8421	3.5	2.0515	4.5	0.1911	2.8	0.62	1127.2	29.2	1132.8	30.9	1143.5	70.5	1143.5	70.5	98.6
Trial-1_0965	61	519311	1.2	11.1075	2.9	3.1212	3.8	0.2514	2.4	0.63	1445.9	30.7	1437.9	28.9	1426.1	55.7	1426.1	55.7	101.4
Trial-1_0966	126	53318	1.2	16.6642	3.5	0.8171	4.3	0.0988	2.6	0.61	607.1	15.2	606.4	19.8	603.9	74.8	607.1	15.2	100.5
Trial-1_0967	109	46674	2.5	20.5127	6.7	0.1923	8.3	0.0286	4.9	0.59	181.8	8.8	178.6	13.6	135.8	157.8	181.8	8.8	NA
Trial-1_0968	87	257971	0.7	16.4322	4.1	0.8715	5.1	0.1039	3.1	0.61	637.0	18.8	636.4	24.2	634.2	87.5	637.0	18.8	100.5
Trial-1_0969	37	117999	1.4	13.3460	4.6	1.8392	5.9	0.1780	3.6	0.62	1056.1	35.3	1059.6	38.5	1066.6	92.6	1066.6	92.6	99.0
Trial-1_0970	319	61570	2.4	20.1114	4.1	0.1879	5.1	0.0274	3.0	0.59	174.3	5.2	174.9	8.3	182.0	96.7	174.3	5.2	NA
Trial-1_0971	245	62631	1.2	19.5451	3.9	0.2919	4.8	0.0414	2.9	0.60	261.4	7.3	260.0	11.0	248.2	89.1	261.4	7.3	NA
Trial-1_0972	109	19397	1.2	18.3195	4.6	0.5066	5.7	0.0673	3.4	0.60	420.0	13.9	416.2	19.5	395.3	102.8	420.0	13.9	106.2
Trial-1_0973	185	27373	1.2	20.7471	5.6	0.1786	6.9	0.0269	4.1	0.59	171.0	6.9	166.9	10.6	109.1	131.5	171.0	6.9	NA
Trial-1_0974	55	38888	1.0	12.9556	3.6	2.0329	4.5	0.1910	2.8	0.62	1126.9	29.2	1126.6	30.9	1126.0	70.8	1126.0	70.8	100.1
Trial-1_0975	146	1033794	1.2	10.8945	1.9	3.1427	2.4	0.2483	1.5	0.63	1429.8	19.5	1443.2	18.6	1463.0	35.6	1463.0	35.6	97.7
Trial-1_0976	49	105313	3.7	9.2018	2.8	4.4200	3.7	0.2950	2.3	0.64	1666.4	34.3	1716.1	30.3	1777.3	51.2	1777.3	51.2	93.8
Trial-1_0977	35	53516	1.6	12.9846	4.5	1.9698	5.7	0.1855	3.6	0.62	1097.0	36.0	1105.2	38.7	1121.6	89.7	1121.6	89.7	97.8
Trial-1_0978	122	1240823	4.4	5.2649	1.3	13.5066	1.8	0.5157	1.2	0.66	2681.1	26.0	2715.7	16.9	2741.6	22.0	2741.6	22.0	97.8
Trial-1_0979	241	525883	2.3	13.0488	1.8	1.9055	2.2	0.1803	1.4	0.62	1068.8	13.7	1083.0	14.9	1111.7	35.0	1111.7	35.0	96.1
Trial-1_0980	58	84869	8.5	9.3212	2.6	4.5819	3.4	0.3098	2.2	0.64	1739.5	32.8	1746.0	28.1	1753.7	47.4	1753.7	47.4	99.2
Trial-1_0981	50	61552	3.9	9.2809	2.8	4.5943	3.6	0.3093	2.3	0.64	1737.0	35.2	1748.2	30.2	1761.7	50.9	1761.7	50.9	98.6
Trial-1_0982	76	196260	1.3	5.2901	1.7	13.5375	2.2	0.5194	1.5	0.66	2696.6	32.7	2717.9	21.2	2733.7	27.5	2733.7	27.5	98.6
Trial-1_0983	286	332022	2.1	13.1202	1.7	1.8174	2.1	0.1729	1.3	0.62	1028.3	12.6	1051.7	14.0	1100.8	33.6	1100.8	33.6	93.4
Trial-1_0984	87	162803	2.4	13.1018	2.9	1.9596	3.7	0.1862	2.3	0.62	1100.8	23.5	1101.7	25.1	1103.6	58.6	1103.6	58.6	99.7
Trial-1_0985	73	53597	1.3	11.0319	2.7	2.9924	3.5	0.2394	2.2	0.63	1383.7	27.4	1405.7	26.6	1439.2	51.9	1439.2	51.9	96.1
Trial-1_0986	39	151028	3.9	5.9995	2.5	10.9050	3.3	0.4745	2.2	0.66	2503.3	45.1	2515.1	30.8	2524.6	41.9	2524.6	41.9	99.2
Trial-1_0987	106	168246	0.9	13.5980	2.8	1.7436	3.5	0.1720	2.2	0.62	1022.9	20.7	1024.8	22.9	1028.9	56.3	1028.9	56.3	99.4
Trial-1_0988	40	48916	2.6	12.6672	4.0	2.1417	5.1	0.1968	3.2	0.62	1157.9	33.9	1162.4	35.5	1170.8	79.5	1170.8	79.5	98.9
Trial-1_0989	124	143940	2.4	9.2196	1.8	4.6010	2.3	0.3077	1.5	0.64	1729.2	22.6	1749.5	19.4	1773.8	32.7	1773.8	32.7	97.5
Trial-1_0990	210	106906	3.5	13.1560	1.9	1.8895	2.5	0.1803	1.5	0.62	1068.5	15.0	1077.4	16.4	1095.4	38.7	1095.4	38.7	97.6
Trial-1_0991	122	41097	1.5	19.8558	8.2	0.1165	10.3	0.0168	6.3	0.61	107.2	6.7	111.9	10.9	211.8	189.3	107.2	6.7	NA
Trial-1_0992	61	482699	0.9	5.1639	1.9	14.2947	2.5	0.5354	1.7	0.66	2764.0	37.9	2769.5	24.1	2773.4	31.2	2773.4	31.2	99.7
Trial-1_0993	95	248408	1.5	12.5811	2.7	2.2077	3.4	0.2014	2.1	0.62	1183.1	22.9	1183.5	23.8	1184.2	52.6	1184.2	52.6	99.9
Trial-1_0994	92	141915	2.1	13.3925	2.9	1.7830	3.6	0.1732	2.3	0.62	1029.7	21.5	1039.3	23.7	1059.6	57.6	1059.6	57.6	97.2
Trial-1_0995	146	79840	4.3	16.7500	3.2	0.7879	4.1	0.0957	2.5	0.60	589.2	13.9	590.0	18.2	592.8	70.3	589.2	13.9	99.4
Trial-1_0996	20	26145	1.4	13.2799	6.0	1.9006	7.6	0.1831	4.7	0.62	1083.6	47.2	1081.3	50.8	1076.6	120.3	1076.6	120.3	100.7
Trial-1_0997	94	81906	1.8	12.9388	2.7	1.9531	3.5	0.1833	2.2	0.62	1084.9	21.7	1099.5	23.4	1128.6	54.5	1128.6	54.5	96.1
Trial-1_0998	41	64406	3.1	13.9377	4.6	1.5835	5.8	0.1601	3.6	0.62	957.2	32.0	963.8	36.3	978.8	93.5	978.8	93.5	97.8
Trial-1_0999	24	16875	1.1	13.4181	5.7	1.8029	7.3	0.1755	4.5	0.62	1042.1	43.4	1046.5	47.7	1055.8	115.7	1055.8	115.7	98.7
Trial-1_1000	42	41682	2.1	13.3359	4.3	1.8442	5.5	0.1784	3.4	0.62	1058.1	33.2	1061.4	36.3	1068.1	87.0	1068.1	87.0	99.1
Trial-1_1001	41	56947	1.0	11.1210	3.6	3.0233	4.7	0.2439	2.9	0.63	1406.7	37.2	1413.5	35.8	1423.8	69.6	1423.8	69.6	98.8
Trial-1_1002	151	215801	3.1	13.2625	2.3	1.8697	2.9	0.1798	1.8	0.62	1066.1	17.8	1070.4	19.4	1079.2	46.2	1079.2	46.2	98.8
Trial-1_1003	143	27602	2.9	19.6525	5.2	0.2887	6.4	0.0411	3.8	0.59	259.9	9.7	257.5	14.6	235.6	119.1	259.9	9.7	NA
Trial-1_1004	174	398978	2.8	10.9047	1.9	3.0511	2.4	0.2413	1.5	0.63	1393.5	18.6	1420.5	18.2	1461.2	35.2	1461.2	35.2	95.4
Trial-1_1005	120	512606	1.1	18.2076	4.1	0.5551	5.2	0.0733	3.1	0.60	456.1	13.7	448.4	18.8	409.0	92.6	456.1	13.7	111.5
Trial-1_1006	97	281347	1.7	16.2762	3.8	0.8778	4.8	0.1036	2.9	0.61	635.6	17.7	639.8	22.9	654.6	82.3	635.6	17.7	97.1
Trial-1_1007	28	60715	1.1	13.3707	5.2	1.7679	6.6	0.1714	4.1	0.62	1020.0	38.8	1033.8	43.1	1062.9	105.0	1062.9	105.0	96.0
Trial-1_1008	52	336768	0.4	5.3426	2.1	13.2800	2.8	0.5146	1.8	0.66	2676.1	40.1	2699.8	26.1	2717.5	34.1	2717.5	34.1	98.5
Trial-1_1009	135	303124	2.0	12.6195	2.2	2.1142	2.9	0.1935	1.8	0.62	1140.3	18.6	1153.4	19.6	1178.2	44.1	1178.2	44.1	96.8
Trial-1_1010	98	561018	0.6	5.1051	1.5	14.1080	2.0	0.5224	1.3	0.66	2709.2	29.7	2757.0	19.2	2792.2	24.8	2792.2	24.8	97.0
Trial-1_1011	234	1335837	1.4	10.8345	1.5	3.1091	2.0	0.2443	1.2	0.63	1409.1	15.7	1434.9	15.1	1473.5	28.9	1473.5	28.9	95.6
Trial-1_1012	201	700584	1.7	8.8032	1.4	5.0307	1.8	0.3212	1.2	0.64	1795.6	18.3	1824.5	15.4	1857.7	25.2	1857.7	25.2	96.7
Trial-1_1013	34	71675	0.7	9.1980	3.2	4.4791	4.2	0.2988	2.7	0.64	1685.4	40.1	1727.1	34.9	1778.0	58.8	1778.0	58.8	94.8
Trial-1_1014	207	722725	0.9	9.4266	1.4	4.3829	1.9	0.2997	1.2	0.64	1689.6	17.6	1709.1	15.3	1733.1	26.1	1733.1	26.1	97.5
Trial-1_1015	528	1358496	0.8	19.5591	2.7	0.2866	3.3	0.0407	2.0	0.59	256.9	4.9	255.9	7.5	246.5	61.1	256.9	4.9	NA
Trial-1_1016	50	6195	0.7	21.5562	10.0	0.1855	12.3	0.0290	7.2	0.59	184.3	13.1	172.8	19.6	17.9	240.0	184.3	13.1	NA
Trial-1_1017	31	117377	3.2	13.2622	5.0	1.8605	6.3	0.1790	3.9	0.62	1061.2	38.4	1067.2	41.9	1079.3	100.1	1079.3	100.1	98.3
Trial-1_1018	217	192696	3.1	13.1824	1.9	1.8980	2.4	0.1815	1.5	0.62	1075.0	14.6	1080.4	15.8	1091.3	37.4	1091.3	37.4	98.5
Trial-1_1019	95	13166	1.0	20.0520	6.5	0.2675	8.0	0.0389	4.8	0.59	246.0	11.5	240.7	17.2	188.9	150.5	246.0	11.5	NA
Trial-1_1020	630	925665	3.2	13.0770	1.1	1.7573	1.5	0.1667	0.9	0.62	993.7	8.3	1029.8	9.4	1107.4	22.8	1107.4	22.8	89.7
Trial-1_1021	128	18165	0.9	19.5138	5.4	0.3002	6.7	0.0425	4.0	0.60	268.2	10.4	266.5	15.6	251.9	123.2	268.2	10.4	NA
Trial-1_1022	108	315627	1.3	12.6515	2.5	2.0820	3.1	0.1910	2.0	0.62	1127.0	20.3	1142.9	21.6	1173.2	48.7	1173.2	48.7	96.1
Trial-1_1023	54	85854	1.6	13.0088	3.6	1.9340	4.6	0.1825	2.9	0.62	1080.5	28.4	1092.9	30.8	1117.9	71.8	1117.9	71.8	96.7
Trial-1_1024	41	201776	0.9	5.8326	2.4	11.4880	3.2	0.4860	2.1	0.66	2553.2	44.8	2563.6	30.2	2571.8	40.7	2571.8	40.7	99.3
Trial-1_1025	126	22760	1.0	19.7785	5.6	0.2730	6.9	0.0392	4.1	0.59	247.6	10.0	245.1	15.1	220.8	128.8	247.6	10.0	NA
Trial-1_1026	119	106109	2.7	11.9505	2.2	2.5446	2.9	0.2205	1.8	0.63	1284.8	20.9	1284.9	20.9	1285.1	43.6	1285.1	43.6	100.0
Trial-1_1027	732	821947	1.6	17.4513	1.7	0.5605	2.1	0.0709	1.3	0.60	441.8	5.3	451.9	7.6	503.2	36.5	441.8	5.3	87.8
Trial-1_1028	34	305880	0.9	8.7922	3.3	5.2333	4.3	0.3337	2.7	0.64	1856.4	44.2	1858.1	36.4	1859.9	59.1	1859.9	59.1	99.8
Trial-1_1029	71	194879	1.5	11.0098	2.6	3.1182	3.4	0.2490	2.2	0.63	1433.3	27.6	1437.2	26.2	1443.0	50.4	1443.0	50.4	99.3

Trial 2: CP-40 Nu Plasma HR TC						Isotope ratios							Apparent ages (Ma)						
Analysis	U (ppm)	206Pb 204Pb	U/Th	206Pb* 207Pb*	± (%)	207Pb* 235U*	± (%)	206Pb* 238U	± (%)	error corr.	206Pb* 238U*	± (Ma)	207Pb* 235U	± (Ma)	206Pb* 207Pb*	± (Ma)	Best age (Ma)	± (Ma)	Conc (%)
Trial-2_0001	23	328283	1.5	9.1946	4.8	4.5494	6.3	0.3034	4.0	0.64	1708.1	60.2	1740.1	52.2	1778.7	87.8	1778.7	87.8	96.0
Trial-2_0002	66	66897	2.6	12.9704	3.9	1.8437	5.0	0.1734	3.1	0.62	1031.0	29.4	1061.2	32.7	1123.7	77.5	1123.7	77.5	91.8
Trial-2_0003	53	65015	1.6	13.1081	4.4	1.9174	5.6	0.1823	3.5	0.62	1079.5	34.4	1087.2	37.1	1102.7	87.1	1102.7	87.1	97.9
Trial-2_0004	68	48203	1.9	13.4366	4.0	1.8000	5.1	0.1754	3.1	0.62	1041.9	30.1	1045.5	33.0	1053.0	79.9	1053.0	79.9	98.9
Trial-2_0005	59	74757	2.6	10.9470	3.5	2.9998	4.5	0.2382	2.9	0.63	1377.2	35.5	1407.6	34.5	1453.9	66.7	1453.9	66.7	94.7
Trial-2_0006	29	5125	1.5	20.0619	10.2	0.4884	12.8	0.0711	7.6	0.60	442.6	32.6	403.8	42.6	187.8	239.0	442.6	32.6	235.7
Trial-2_0007	86	19210	2.9	18.8018	5.8	0.5152	7.3	0.0702	4.4	0.60	437.7	18.6	421.9	25.2	336.7	132.4	437.7	18.6	130.0
Trial-2_0008	31	24897	1.8	9.3888	4.4	4.5763	5.7	0.3116	3.6	0.64	1748.7	55.8	1745.0	47.6	1740.5	80.5	1740.5	80.5	100.5
Trial-2_0009	38	3574	0.4	22.0269	10.8	0.2959	13.4	0.0473	7.9	0.59	297.8	22.8	263.2	31.0	-34.2	262.8	297.8	22.8	NA
Trial-2_0010	110	61338	2.7	5.1615	2.0	11.9350	2.7	0.4468	1.8	0.66	2380.9	35.9	2599.3	25.6	2774.2	33.6	2774.2	33.6	85.8
Trial-2_0011	510	114733	4.2	16.7584	2.2	0.7132	2.8	0.0867	1.7	0.61	535.9	8.6	546.6	11.7	591.7	47.7	535.9	8.6	90.6
Trial-2_0012	31	11908	0.6	19.7100	6.2	0.1881	6.7	0.0269	2.6	0.39	171.1	4.4	175.0	10.8	228.8	142.6	171.1	4.4	NA
Trial-2_0013	27	9281	1.2	13.4090	6.2	1.8179	7.9	0.1768	4.9	0.62	1049.4	47.7	1051.9	52.1	1057.1	125.6	1057.1	125.6	99.3
Trial-2_0014	192	94574	6.8	12.1459	2.2	2.2650	2.8	0.1995	1.7	0.63	1172.8	18.6	1201.5	19.6	1253.4	42.3	1253.4	42.3	93.6
Trial-2_0015	16	13593	2.6	13.1975	8.3	1.9238	10.6	0.1841	6.6	0.62	1089.6	66.0	1089.4	71.0	1089.0	166.9	1089.0	166.9	100.0
Trial-2_0016	54	144125	0.6	5.2591	2.4	13.4908	3.1	0.5146	2.1	0.66	2676.2	45.8	2714.6	29.8	2743.4	38.7	2743.4	38.7	97.5
Trial-2_0017	73	125824	2.6	13.7451	4.0	1.7143	5.1	0.1709	3.2	0.62	1017.0	29.8	1013.9	32.8	1007.1	81.5	1007.1	81.5	101.0
Trial-2_0018	456	899675	17.9	12.9949	1.6	1.6822	2.0	0.1585	1.2	0.62	948.7	11.0	1001.8	12.7	1120.0	31.3	1120.0	31.3	84.7
Trial-2_0019	44	168380	1.8	7.7514	3.2	6.6249	4.2	0.3724	2.7	0.65	2040.9	47.2	2062.7	36.7	2084.5	55.7	2084.5	55.7	97.9
Trial-2_0020	60	14837	2.5	20.5287	9.1	0.2944	11.3	0.0438	6.7	0.59	276.6	18.1	262.1	26.0	134.0	213.6	276.6	18.1	NA
Trial-2_0021	35	46443	0.5	5.6179	3.0	12.1416	4.0	0.4947	2.7	0.66	2591.0	57.0	2615.4	37.9	2634.3	50.3	2634.3	50.3	98.4
Trial-2_0022	44	43966	0.5	11.1692	4.2	2.9518	5.4	0.2391	3.4	0.63	1382.1	42.1	1395.3	40.7	1415.5	79.7	1415.5	79.7	97.6
Trial-2_0023	318	1236985	2.7	13.6885	1.9	1.5813	2.4	0.1570	1.5	0.62	940.0	13.2	962.9	15.2	1015.4	38.9	1015.4	38.9	92.6
Trial-2_0024	24	41412	2.8	13.3647	6.7	1.8409	8.5	0.1784	5.3	0.62	1058.4	51.6	1060.2	56.0	1063.8	134.0	1063.8	134.0	99.5
Trial-2_0025	22	11375	2.1	13.5184	6.9	1.8653	8.8	0.1829	5.5	0.62	1082.7	54.6	1068.9	58.4	1040.7	139.7	1040.7	139.7	104.0
Trial-2_0026	36	79439	0.7	8.7017	3.8	5.1864	5.0	0.3273	3.2	0.64	1825.4	51.0	1850.4	42.5	1878.6	68.9	1878.6	68.9	97.2
Trial-2_0027	122	41482	2.2	13.6535	3.1	1.7104	4.0	0.1694	2.5	0.62	1008.6	23.0	1012.4	25.5	1020.6	63.2	1020.6	63.2	98.8
Trial-2_0028	44	83464	1.2	11.1970	4.2	2.9890	5.4	0.2427	3.4	0.63	1400.9	42.6	1404.8	40.9	1410.8	79.8	1410.8	79.8	99.3
Trial-2_0029	122	71920	2.1	13.2482	3.0	1.8320	3.8	0.1760	2.4	0.62	1045.3	23.1	1057.0	25.3	1081.4	60.4	1081.4	60.4	96.7
Trial-2_0030	38	17153	0.8	16.8206	7.5	0.8124	9.4	0.0991	5.7	0.61	609.2	33.1	603.8	42.8	583.6	162.4	609.2	33.1	104.4
Trial-2_0031	509	27620	1.8	20.0953	3.4	0.2057	4.2	0.0300	2.5	0.60	190.4	4.7	190.0	7.3	183.9	79.0	190.4	4.7	NA
Trial-2_0032	96	29673	0.9	16.6875	4.7	0.7961	5.9	0.0963	3.6	0.61	593.0	20.3	594.6	26.6	600.8	101.5	593.0	20.3	98.7
Trial-2_0033	66	125341	1.5	9.5749	3.0	4.2935	4.0	0.2982	2.5	0.64	1682.2	37.4	1692.1	32.6	1704.5	56.1	1704.5	56.1	98.7
Trial-2_0034	105	103764	4.8	9.2939	2.3	4.4992	3.0	0.3033	1.9	0.64	1707.5	29.1	1730.8	25.2	1759.1	42.6	1759.1	42.6	97.1
Trial-2_0035	209	123479	1.1	10.9029	1.9	3.0721	2.4	0.2429	1.5	0.63	1401.9	19.5	1425.8	18.7	1461.5	36.0	1461.5	36.0	95.9
Trial-2_0036	12	5394	1.6	12.5326	8.3	2.0998	9.1	0.1909	3.7	0.41	1126.0	38.2	1148.7	62.9	1191.9	165.0	1191.9	165.0	94.5
Trial-2_0037	50	24159	1.9	13.1554	4.6	1.9447	5.9	0.1856	3.6	0.62	1097.2	36.7	1096.6	39.3	1095.5	91.9	1095.5	91.9	100.2
Trial-2_0038	37	46961	1.1	12.9064	5.3	2.0219	6.8	0.1893	4.2	0.62	1117.4	43.3	1122.9	46.1	1133.6	105.6	1133.6	105.6	98.6
Trial-2_0039	56	85092	3.3	9.3035	3.2	4.5868	4.2	0.3095	2.7	0.64	1738.2	40.6	1746.9	34.7	1757.2	58.5	1757.2	58.5	98.9
Trial-2_0040	201	959182	1.3	5.2160	1.3	12.9358	1.7	0.4894	1.1	0.66	2567.9	24.1	2675.0	16.2	2757.0	21.1	2757.0	21.1	93.1
Trial-2_0041	91	93455	1.4	10.9721	2.9	2.9940	3.8	0.2383	2.4	0.63	1377.6	29.7	1406.1	28.9	1449.5	56.0	1449.5	56.0	95.0
Trial-2_0042	104	452036	2.0	9.0132	2.3	4.8522	3.0	0.3172	1.9	0.64	1776.0	29.9	1794.0	25.3	1815.0	41.8	1815.0	41.8	97.8
Trial-2_0043	108	423397	1.6	13.1556	3.1	1.8836	4.0	0.1797	2.5	0.62	1065.5	24.6	1075.3	26.7	1095.4	63.0	1095.4	63.0	97.3
Trial-2_0044	99	45056	2.1	13.7121	3.5	1.6573	4.5	0.1648	2.8	0.62	983.5	25.1	992.3	28.2	1011.9	70.9	1011.9	70.9	97.2
Trial-2_0045	113	72473	1.6	13.5705	3.3	1.6251	4.2	0.1599	2.6	0.62	956.5	23.0	980.0	26.2	1032.9	66.1	1032.9	66.1	92.6
Trial-2_0046	103	102339	1.6	9.6029	2.4	4.1922	3.2	0.2920	2.0	0.64	1651.4	29.5	1672.5	26.0	1699.1	45.0	1699.1	45.0	97.2
Trial-2_0047	88	187011	1.1	13.1855	3.6	1.9359	4.6	0.1851	2.8	0.62	1094.9	28.5	1093.6	30.5	1090.8	71.5	1090.8	71.5	100.4
Trial-2_0048	19	24667	1.2	13.2289	7.3	1.9313	9.3	0.1853	5.8	0.62	1095.9	58.0	1092.0	62.2	1084.3	146.1	1084.3	146.1	101.1
Trial-2_0049	11	12596	1.0	11.0873	8.7	2.9963	11.1	0.2409	7.0	0.63	1391.6	87.6	1406.7	84.9	1429.6	165.5	1429.6	165.5	97.3
Trial-2_0050	337	164728	2.7	13.4056	1.9	1.6302	2.4	0.1585	1.5	0.62	948.4	12.9	981.9	14.9	1057.6	37.3	1057.6	37.3	89.7
Trial-2_0051	231	79394	2.5	13.7017	2.3	1.6340	2.9	0.1624	1.8	0.62	970.0	16.1	983.4	18.2	1013.5	45.9	1013.5	45.9	95.7
Trial-2_0052	168	110258	1.0	13.2970	2.5	1.8134	3.2	0.1749	2.0	0.62	1038.9	19.4	1050.3	21.2	1074.0	51.0	1074.0	51.0	96.7
Trial-2_0053	62	37563	1.9	12.8247	4.1	1.9916	5.2	0.1852	3.3	0.62	1095.6	32.8	1112.7	35.2	1146.2	80.9	1146.2	80.9	95.6
Trial-2_0054	80	94712	2.2	13.0926	3.5	1.9808	4.5	0.1881	2.8	0.62	1111.0	28.6	1109.0	30.3	1105.0	70.3	1105.0	70.3	100.5
Trial-2_0055	45	36149	1.4	11.1128	4.2	3.0438	5.4	0.2453	3.4	0.63	1414.3	43.7	1418.7	41.6	1425.2	80.6	1425.2	80.6	99.2
Trial-2_0056	91	41748	0.9	16.6184	4.6	0.8156	5.8	0.0983	3.5	0.61	604.5	20.2	605.6	26.3	609.8	99.1	604.5	20.2	99.1
Trial-2_0057	48	30100	1.1	13.6130	4.8	1.7802	6.1	0.1758	3.8	0.62	1043.8	36.6	1038.2	39.8	1026.6	97.1	1026.6	97.1	101.7
Trial-2_0058	136	223370	1.3	10.9640	2.4	3.0324	3.1	0.2411	1.9	0.63	1392.6	24.2	1415.8	23.3	1450.9	45.1	1450.9	45.1	96.0
Trial-2_0059	51	49320	1.3	11.4134	3.9	2.8375	5.1	0.2349	3.2	0.63	1360.0	39.3	1365.5	38.2	1374.1	76.0	1374.1	76.0	99.0
Trial-2_0060	1346	28126	4.0	11.8930	1.1	2.1375	1.4	0.1844	0.9	0.62	1090.8	9.1	1161.0	10.0	1294.5	22.0	1294.5	22.0	84.3

Trial-2_0061	196	278443	1.6	8.2846	1.7	5.5291	2.2	0.3322	1.4	0.65	1849.1	22.6	1905.1	18.7	1966.7	29.7	1966.7	29.7	94.0
Trial-2_0062	49	35218	1.2	13.4430	4.7	1.8368	6.0	0.1791	3.7	0.62	1062.0	36.5	1058.7	39.5	1052.0	95.0	1052.0	95.0	100.9
Trial-2_0063	313	689516	6.2	9.9564	1.6	3.4028	2.0	0.2457	1.3	0.64	1416.4	16.4	1505.0	15.9	1632.2	29.1	1632.2	29.1	86.8
Trial-2_0064	32	53401	0.6	11.0818	4.7	3.0927	6.1	0.2486	3.9	0.63	1431.1	49.4	1430.9	46.8	1430.5	90.4	1430.5	90.4	100.0
Trial-2_0065	45	21639	1.2	11.1445	4.1	3.0526	5.3	0.2467	3.3	0.63	1421.6	42.6	1420.9	40.5	1419.8	78.4	1419.8	78.4	100.1
Trial-2_0066	20	37188	1.9	11.7574	6.3	2.6900	8.1	0.2294	5.1	0.63	1331.3	61.4	1325.7	60.1	1316.7	122.3	1316.7	122.3	101.1
Trial-2_0067	18	204803	1.4	13.4658	7.7	1.8193	9.8	0.1777	6.1	0.62	1054.3	59.1	1052.4	64.1	1048.6	154.5	1048.6	154.5	100.5
Trial-2_0068	26	36764	0.8	8.7281	4.5	5.2217	5.9	0.3305	3.8	0.64	1841.0	60.4	1856.2	50.0	1873.2	80.9	1873.2	80.9	98.3
Trial-2_0069	91	247804	0.9	11.0320	2.9	3.0297	3.7	0.2424	2.3	0.63	1399.2	29.5	1415.1	28.4	1439.1	54.9	1439.1	54.9	97.2
Trial-2_0070	401	29573	2.5	15.3104	2.0	1.0825	2.5	0.1202	1.6	0.62	731.7	10.8	744.8	13.3	784.5	41.3	731.7	10.8	93.3
Trial-2_0071	4	3190	-13.9	16.5817	8.2	0.7342	8.3	0.0883	1.8	0.21	545.4	9.3	559.0	35.9	614.6	176.4	545.4	9.3	88.7
Trial-2_0072	125	118430	0.9	13.2851	3.0	1.8198	3.8	0.1753	2.4	0.62	1041.5	23.0	1052.6	25.2	1075.8	60.5	1075.8	60.5	96.8
Trial-2_0073	40	58428	0.8	8.7201	3.5	5.1548	4.6	0.3260	3.0	0.64	1819.0	47.2	1845.2	39.4	1874.8	63.8	1874.8	63.8	97.0
Trial-2_0074	138	181435	12.0	18.7178	4.8	0.4939	6.0	0.0670	3.6	0.60	418.3	14.5	407.6	20.0	346.9	108.1	418.3	14.5	120.6
Trial-2_0075	201	73527	1.2	16.8227	3.2	0.7753	4.0	0.0946	2.4	0.61	582.6	13.6	582.8	17.9	583.4	69.5	582.6	13.6	99.9
Trial-2_0076	31	294958	0.9	5.9842	3.3	10.1314	4.5	0.4397	2.9	0.66	2349.4	57.8	2446.8	41.2	2528.9	56.2	2528.9	56.2	92.9
Trial-2_0077	49	23718	0.6	16.9255	6.5	0.7911	8.2	0.0971	5.0	0.61	597.4	28.3	591.8	36.6	570.2	141.2	597.4	28.3	104.8
Trial-2_0078	206	30667	1.9	20.1423	5.3	0.2656	6.5	0.0388	3.9	0.60	245.4	9.4	239.2	14.0	178.5	122.7	245.4	9.4	NA
Trial-2_0079	34	27320	1.2	17.9367	8.4	0.6451	10.5	0.0839	6.3	0.60	519.5	31.7	505.5	41.8	442.5	186.1	519.5	31.7	117.4
Trial-2_0080	34	30618	0.7	10.9443	4.7	3.2046	6.0	0.2544	3.8	0.63	1461.0	49.7	1458.3	46.6	1454.4	88.8	1454.4	88.8	100.5
Trial-2_0081	985	290490	4.7	17.9261	1.8	0.4719	2.3	0.0613	1.4	0.60	383.8	5.1	392.5	7.4	443.8	40.2	383.8	5.1	NA
Trial-2_0082	58	28234	1.8	13.8739	4.6	1.5789	5.9	0.1589	3.6	0.62	950.5	32.2	962.0	36.5	988.1	93.9	988.1	93.9	96.2
Trial-2_0083	66	34368	3.2	13.9884	4.1	1.5919	5.2	0.1615	3.3	0.62	965.2	29.1	967.1	32.7	971.4	84.1	971.4	84.1	99.4
Trial-2_0084	123	357266	0.7	13.1812	2.9	1.8553	3.8	0.1774	2.3	0.62	1052.6	22.7	1065.3	24.8	1091.5	58.9	1091.5	58.9	96.4
Trial-2_0085	9	4784	1.0	12.8667	1.6	2.0084	8.5	0.1874	8.3	0.98	1107.4	84.7	1118.4	57.6	1139.7	32.7	1139.7	32.7	97.2
Trial-2_0086	302	203773	3.7	13.1346	1.9	1.8005	2.5	0.1715	1.5	0.62	1020.5	14.6	1045.6	16.2	1098.6	38.9	1098.6	38.9	92.9
Trial-2_0087	35	36117	1.5	5.4617	3.0	12.4871	4.0	0.4946	2.6	0.66	2590.7	56.4	2641.8	37.5	2681.1	49.4	2681.1	49.4	96.6
Trial-2_0088	191	13522	0.6	20.6836	5.6	0.2684	7.0	0.0403	4.2	0.59	254.4	10.4	241.4	15.0	116.3	132.8	254.4	10.4	NA
Trial-2_0089	331	478961	2.4	12.7364	1.8	1.9753	2.3	0.1825	1.4	0.62	1080.4	14.1	1107.1	15.3	1159.9	35.2	1159.9	35.2	93.1
Trial-2_0090	90	150679	1.4	11.0307	3.0	2.9685	3.8	0.2375	2.4	0.63	1373.6	30.0	1399.6	29.2	1439.4	56.9	1439.4	56.9	95.4
Trial-2_0091	80	27458	1.3	20.5828	8.5	0.2620	10.6	0.0391	6.3	0.59	247.3	15.3	236.3	22.4	127.8	201.3	247.3	15.3	NA
Trial-2_0092	35	25625	1.1	13.7395	5.5	1.7312	7.0	0.1725	4.3	0.62	1025.9	41.2	1020.2	45.1	1007.9	111.4	1007.9	111.4	101.8
Trial-2_0093	120	37598	2.0	17.3076	4.6	0.6373	5.8	0.0800	3.5	0.61	496.1	16.6	500.6	22.8	521.3	100.6	496.1	16.6	95.2
Trial-2_0094	120	44658	2.8	13.6550	3.1	1.7465	3.9	0.1730	2.4	0.62	1028.4	23.2	1025.9	25.4	1020.4	62.6	1020.4	62.6	100.8
Trial-2_0095	47	88491	0.5	13.0499	4.6	1.9043	5.9	0.1802	3.7	0.62	1068.3	36.2	1082.6	39.3	1111.5	92.2	1111.5	92.2	96.1
Trial-2_0096	77	224501	2.5	5.4179	2.0	12.7609	2.7	0.5014	1.8	0.66	2619.9	38.0	2662.2	25.1	2694.4	32.9	2694.4	32.9	97.2
Trial-2_0097	66	96605	1.9	13.3174	4.2	1.8136	5.3	0.1752	3.3	0.62	1040.6	31.8	1050.4	34.8	1070.9	83.7	1070.9	83.7	97.2
Trial-2_0098	382	360640	2.8	13.1466	1.7	1.7980	2.2	0.1714	1.4	0.62	1020.0	12.7	1044.7	14.2	1096.8	34.0	1096.8	34.0	93.0
Trial-2_0099	169	177092	2.2	12.6736	2.4	2.0635	3.1	0.1897	1.9	0.62	1119.6	20.0	1136.8	21.3	1169.7	48.2	1169.7	48.2	95.7
Trial-2_0100	44	13251	1.5	20.4322	9.9	0.3735	12.3	0.0553	7.3	0.60	347.2	24.7	322.2	33.9	145.0	231.8	347.2	24.7	NA
Trial-2_0101	19	56976	1.5	13.4763	7.5	1.7501	9.6	0.1711	5.9	0.62	1017.9	55.9	1027.2	61.8	1047.0	151.1	1047.0	151.1	97.2
Trial-2_0102	697	53749	3.0	15.6161	2.4	0.3908	3.0	0.0443	1.9	0.61	279.2	5.1	335.0	8.7	742.8	51.2	279.2	5.1	NA
Trial-2_0103	36	180944	1.5	5.2748	2.9	13.6753	3.9	0.5232	2.6	0.66	2712.6	56.9	2727.5	36.6	2738.5	47.7	2738.5	47.7	99.1
Trial-2_0104	375	165732	1.0	18.1230	2.9	0.4988	3.6	0.0656	2.2	0.60	409.4	8.7	410.9	12.3	419.4	65.0	409.4	8.7	97.6
Trial-2_0105	53	46218	1.4	11.1578	3.6	3.0420	4.7	0.2462	3.0	0.63	1418.7	37.7	1418.2	35.8	1417.5	69.5	1417.5	69.5	100.1
Trial-2_0106	68	247858	2.2	13.2865	4.0	1.7793	5.1	0.1715	3.2	0.62	1020.1	29.8	1037.9	33.1	1075.6	80.2	1075.6	80.2	94.8
Trial-2_0107	169	55260	0.9	17.9517	4.0	0.5589	5.0	0.0728	3.0	0.60	452.8	13.3	450.8	18.4	440.6	89.7	452.8	13.3	102.8
Trial-2_0108	23	5541	0.9	20.1667	11.9	0.4452	14.8	0.0651	8.9	0.60	406.7	34.9	373.9	46.4	175.6	278.4	406.7	34.9	231.5
Trial-2_0109	138	8728	1.1	22.4305	10.1	0.0935	12.5	0.0152	7.4	0.59	97.3	7.1	90.7	10.9	-78.4	247.7	97.3	7.1	NA
Trial-2_0110	206	275111	1.5	10.8893	2.0	3.0179	2.5	0.2383	1.6	0.63	1378.1	19.9	1412.1	19.3	1463.9	37.4	1463.9	37.4	94.1
Trial-2_0111	512	213749	4.5	16.4580	2.1	0.7286	2.6	0.0870	1.6	0.61	537.6	8.3	555.7	11.3	630.8	45.2	537.6	8.3	85.2
Trial-2_0112	88	41909	1.3	12.5033	3.2	2.2593	4.1	0.2049	2.6	0.62	1201.5	28.4	1199.7	29.2	1196.5	63.8	1196.5	63.8	100.4
Trial-2_0113	24	10738	1.8	13.2961	6.8	1.8534	8.6	0.1787	5.3	0.62	1060.0	52.2	1064.6	56.9	1074.1	136.1	1074.1	136.1	98.7
Trial-2_0114	157	142971	1.4	9.4390	2.0	4.2581	2.6	0.2915	1.6	0.64	1649.0	23.8	1685.3	21.1	1730.7	36.1	1730.7	36.1	95.3
Trial-2_0115	60	64618	1.6	13.0423	4.0	1.9853	5.1	0.1878	3.2	0.62	1109.4	32.6	1110.5	34.7	1112.7	80.4	1112.7	80.4	99.7
Trial-2_0116	28	39209	1.1	13.3090	6.0	1.8883	7.7	0.1823	4.8	0.62	1079.4	47.8	1077.0	51.3	1072.2	121.5	1072.2	121.5	100.7
Trial-2_0117	47	50093	1.3	11.0745	4.1	3.0472	5.3	0.2448	3.3	0.63	1411.4	42.3	1419.5	40.5	1431.8	78.4	1431.8	78.4	98.6
Trial-2_0118	212	37429	1.1	18.6328	3.9	0.4854	4.9	0.0656	2.9	0.60	409.6	11.7	401.8	16.2	357.2	88.1	409.6	11.7	114.7
Trial-2_0119	42	25359	1.2	13.0505	2.1	1.9595	4.5	0.1855	4.0	0.89	1096.8	40.5	1101.7	30.4	1111.4	41.2	1111.4	41.2	98.7
Trial-2_0120	434	320980	2.8	12.7644	1.6	1.9140	2.0	0.1772	1.3	0.62	1051.6	12.2	1086.0	13.4	1155.6	31.1	1155.6	31.1	91.0
Trial-2_0121	271	247956	0.8	18.2335	3.4	0.5253	4.3	0.0695	2.6	0.60	432.9	10.9	428.7	15.1	405.8	77.2	432.9	10.9	106.7
Trial-2_0122	116	141361	1.5	11.0965	2.6	3.0353	3.3	0.2443	2.1	0.63	1408.9	26.8	1416.5	25.6	1428.0	49.6	1428.0	49.6	98.7
Trial-2_0123	125	142214	1.0	8.7778	2.1	5.0836	2.7	0.3236	1.7	0.64	1807.5	27.4	1833.4	22.9	1862.9	37.4	1862.9	37.4	97.0
Trial-2_0124	167	84280	1.4	19.0752	4.6	0.4249	5.7	0.0588	3.4	0.60	368.3	12.3	359.6	17.4	304.0	104.9	368.3	12.3	NA
Trial-2_0125	366	110272	0.6	18.4482	3.0	0.4990	3.7	0.0668	2.2	0.60	416.6	9.0	411.0	12.6	379.6	66.7	416.6	9.0	109.8
Trial-2_0126	248	1333554	1.6	9.3012	1.6	4.2900	2.1	0.2894	1.3	0.64	1638.5	19.0	1691.4	16.9	1757.7	28.8	1757.7	28.8	93.2
Trial-2_0127	117	98129	1.5	10.6460	2.5	3.3972	3.2	0.2623	2.1	0.63	1501.6	27.5	1503.7	25.4	1506.7	47.3	1506.7	47.3	99.7
Trial-2_0128	87	154406	1.6	5.0431	1.9	14.2445	2.5	0.5210	1.7	0.67	2703.4	36.6	2766.1	23.6	2812.2	30.3	2812.2	30.3	96.1
Trial-2_0129	22	11549	1.4	13.3154	3.0	1.8403	6.2	0.1777	5.5	0.88	1054.5	53.4	1060.0	41.1	1071.2	59.7	1071.2	59.7	98.4
Trial-2_0130	124	266848	1.3	5.0347	1.6	14.1528	2.1	0.5168	1.4	0.67	2685.6	30.6	2760.0	19.8	2814.9	25.5	2814.9	25.5	95.4
Trial-2_0131	124	96969	2.3	11.0703	2.4	3.0301	3.1	0.2433	2.0	0.63	1403.7	24.9	1415.2	23.9	1432.5	46.2	1432.5	46.2	98.0
Trial-2_0132	26	60717	1.1	5.2450	3.6	13.2333	4.8	0.5034	3.2	0.66	2628.4	68.9	2696.4	45.4	2747.8	59.2	2747.8	59.2	95.7
Trial-2_0133	181	736286	1.3	5.1229	1.3	13.2099	1.8	0.4908	1.2	0.67	2574.2	25.1	2694.8	16.8	2786.5	21.7	2786.5	21.7	92.4
Trial-2_0134	339	370484	11.1	9.2552	1.4	4.2194	1.8	0.2832	1.1	0.64	1607.6	16.1	1677.8	14.5	1766.7	24.7	1766.7	24.7	91.0
Trial-2_0135	152	873075	1.2	13.4774	2.7	1.8190	3.5	0.1778	2.2	0.62	1055.0	20.9	1052.3	22.7	1046.9	54.7	1046.9	54.7	100.8
Trial-2_0136	125	10494	0.8	20.7823	6.6	0.2550	8.2	0.0384	4.9	0.59	243.1	11.7	230.6	16.9	105.0	156.2	243.1	11.7	NA
Trial-2_0137	20	17201	0.7	10.9196	1.3	3.0882	5.3	0.2446	5.1	0.97	1410.4	64.7	1429.8	40.5	1458.6	25.1	1458.6	25.1	96.7
Trial-2_0138	91	34491	2.4	19.1244	5.9	0.4354	7.4	0.0604	4.4	0.60	378.0	16.3	367.0	22.8	298.1	134.9	378.0	16.3	NA
Trial-2_0139	117	107163	1.3	13.1322	3.0	1.9610	3.8	0.1868	2.4	0.62	1103.9	24.3	1102.2	25.9	1099.0	60.2	1099.0	60.2	100.4
Trial-2_0140	100	79432	1.0	11.1960	2.8	2.9933	3.6	0.2431	2.3	0.63	1402.6	28.5	1405.9	27.3	1411.0	53.1	1411.0	53.1	99.4
Trial-2_0141	357	246359	3.6	11.0417	1.5	2.9537	1.9	0.2365	1.2	0.63	1368.7	15.0	1395.8	14.5	1437.5	28.3	1437.5	28.3	95.2
Trial-2_0142	289	241139	6.8	13.1319	1.8	1.8981	2.4	0.1808	1.5	0.63	1071.2	14.8	1080.4	15.7	1099.0	36.5	1099.0	36.5	97.5
Trial-2_0143	386	206436	2.7	12.4001	1.6	2.0940	2.1	0.1883	1.3	0.63	1112.3	13.2	1146.8	14.2	1212.8	31.7	1212.8	31.7	91.7
Trial-2_0144	142	210870	1.5	13.8143	2.8	1.6628	3.6	0.1666	2.2	0.62	993.3	20.4	994.4	22.6	996.9	56.7	996.9	56.7	99.6
Trial-2_0145	84	88095	2.0	13.6700	3.7	1.6723	4.7	0.1658	2.9	0.62	988.9	26.9	998.1	30.0	1018.2	75.1	1018.2	75.1	97.1
Trial-2_0146	137	14509	0.9	21.7311	7.4	0.1724	9.2	0.0272	5.4	0.59	172.8	9.2	161.5	13.7	-1.5	178.0	172.8	9.2	NA
Trial-2_0147	63	24487	1.0	13.5603	4.3	1.7066	5.5	0.1678	3.4	0.62	1000.2	31.6	1011.0	35.2	1034.5	87.2	1034.5	87.2	96.7
Trial-2_0148	99	130819	1.5	13.7271	3.3	1.7183	4.2	0.1711	2.6	0.62	1018.0	24.7	1015.4	27.1	1009.7	67.2	1009.7	67.2	100.8
Trial-2_0149	380	63096	1.6	20.3025	4.0	0.2735	5.0	0.0403	3.0	0.60	254.5	7.5	245.5	10.9	160.0	94.3	254.5	7.5	NA
Trial-2_0150	104	16468	1.6	14.0197	3.5	1.5473	4.4	0.1573	2.7	0.62	942.0	24.1	949.4	27.4	966.8	71.1	966.8	71.1	97.4
Trial-2_0151	257	89316	2.4	12.5543	2.0	2.0746	2.5	0.1889	1.6	0.63	1115.4	16.2	1140.5	17.3	1188.4	39.0	1188.4	39.0	93.9
Trial-2_0152	126	166495	2.0	13.6371	3.1	1.6883	3.9	0.1670	2.4	0.62	995.5	22.5	1004.1	25.1	1023.1	62.4	1023.1	62.4	97.3
Trial-2_0153	111	211685	1.9	12.8600	2.9	2.0681	3.7	0.1929	2.3	0.62	1137.0	24.4	1138.3	25.6	1140.8	58.1	1140.8	58.1	99.7
Trial-2_0154	105	48378	0.8	8.8872	2.3	4.9615	3.0	0.3198	1.9	0.64	1788.7	29.8	1812.8	25.1	1840.5	41.1	1840.5	41.1	97.2
Trial-2_0155	155	65677	4.4	9.1346	2.2	4.1264	2.9	0.2734	1.8	0.64	1557.9	25.3	1659.5	23.4	1790.7	40.1	1790.7	40.1	87.0
Trial-2_0156	62	95833	1.1	11.1935	3.4	3.0542	4.4	0.2479	2.8	0.63	1427.9	35.9	1421.3	33.9	1411.4	65.6	1411.4	65.6	101.2
Trial-2_0157	57	33343	1.6	12.9942	4.3	2.0097	5.5	0.1894	3.4	0.62	1118.1	35.0	1118.8	37.1	1120.1	85.4	1120.1	85.4	99.8
Trial-2_0158	74	76741	1.1	9.1467	2.9	4.7246	3.7	0.3134	2.4	0.64	1757.5	36.9	1771.6	31.4	1788.3	52.3	1788.3	52.3	98.3
Trial-2_0159	63	208568	1.7	13.5122	4.2	1.8317	5.4	0.1795	3.3	0.62	1064.3	32.6	1056.9	35.2	1041.6	84.6	1041.6	84.6	102.2
Trial-2_0160	35	229098	1.0	5.4290	3.0	12.9708	4.0	0.5107	2.6	0.66	2659.7	57.5	2677.5	37.5	2691.0	49.1	2691.0	49.1	98.8
Trial-2_0161	172	204045	2.8	4.9499	1.4	14.3074	1.9	0.5136	1.2	0.67	2672.1	27.0	2770.3	17.6	2842.6	22.5	2842.6	22.5	94.0
Trial-2_0162	222	214698	2.7	18.5152	3.7	0.5082	4.6	0.0682	2.8	0.60	425.5	11.4	417.2	15.7	371.4	82.5	425.5	11.4	114.6
Trial-2_0163	75	79178	1.0	9.8892	2.9	4.0943	3.8	0.2937	2.4	0.64	1659.8	35.2	1653.2	30.8	1644.8	53.9	1644.8	53.9	100.9
Trial-2_0164	39	8292	0.9	24.1540	12.5	0.2398	15.4	0.0420	9.1	0.59	265.3	23.6	218.3	30.3	-262.8	317.3	265.3	23.6	NA
Trial-2_0165	42	148920	1.1	11.2251	4.2	3.0300	5.4	0.2467	3.4	0.63	1421.3	43.7	1415.2	41.5	1406.0	80.7	1406.0	80.7	101.1
Trial-2_0166	101	24925	2.0	18.0342	5.0	0.6358	6.3	0.0832	3.8	0.60	515.0	18.8	499.7	24.8	430.4	111.7	515.0	18.8	119.7
Trial-2_0167	36	16390	1.4	12.1933	5.0	2.4088	6.4	0.2130	4.0	0.63	1244.9	45.2	1245.2	45.8	1245.8	97.5	1245.8	97.5	99.9
Trial-2_0168	60	53029	1.6	11.7864	3.7	2.7633	4.8	0.2362	3.0	0.63	1367.0	36.9	1345.7	35.5	1311.9	71.8	1311.9	71.8	104.2
Trial-2_0169	308	203436	5.4	13.2771	1.9	1.7576	2.5	0.1692	1.5	0.62	1008.0	14.4	1030.0	16.1	1077.0	39.0	1077.0	39.0	93.6
Trial-2_0170	44	164994	0.8	5.3708	2.7	12.9465	3.6	0.5043	2.4	0.66	2632.3	52.1	2675.8	34.2	2708.8	44.8	2708.8	44.8	97.2
Trial-2_0171	133	107465	2.2	9.3186	2.1	4.4760	2.7	0.3025	1.8	0.64	1703.7	26.2	1726.5	22.7	1754.3	38.4	1754.3	38.4	97.1
Trial-2_0172	298	187595	1.9	13.4247	2.0	1.7187	2.6	0.1673	1.6	0.62	997.5	14.7	1015.6	16.4	1054.8	40.4	1054.8	40.4	94.6
Trial-2_0173	139	82118	1.8	11.0994	2.3	3.0563	3.0	0.2460	1.9	0.63	1418.0	24.4	1421.8	23.2	1427.5	44.8	1427.5	44.8	99.3
Trial-2_0174	212	300811	1.2	5.1378	1.2	13.3083	1.7	0.4959	1.1	0.66	2596.2	23.5	2701.8	15.6	2781.7	20.3	2781.7	20.3	93.3
Trial-2_0175	23	17666	0.9	15.5474	2.1	1.0576	9.6	0.1193	9.3	0.98	726.3	64.0	732.6	49.9	752.1	43.9	726.3	64.0	96.6
Trial-2_0176	228	173752	1.1	12.6056	2.1	2.0695	2.7	0.1892	1.7	0.63	1117.0	17.3	1138.8	18.5	1180.4	41.6	1180.4	41.6	94.6
Trial-2_0177	81	175209	0.9	9.8180	2.8	4.0584	3.6	0.2890	2.3	0.64	1636.4	33.5	1646.0	29.6	1658.2	51.8	1658.2	51.8	98.7
Trial-2_0178	87	33398	1.0	17.7609	5.4	0.6458	6.8	0.0832	4.1	0.60	515.1	20.5	505.9	27.2	464.4	120.7	515.1	20.5	110.9
Trial-2_0179	141	528675	1.3	7.5819	1.8	6.7614	2.4	0.3718	1.6	0.65	2037.9	27.5	2080.7	21.4	2123.3	32.2	2123.3	32.2	96.0
Trial-2_0180	248	51773	1.1	20.7658	5.8	0.1764	7.2	0.0266	4.3	0.59	169.0	7.1	165.0	11.0	107.0	137.6	169.0	7.1	NA
Trial-2_0181	49	150987	1.3	11.2312	4.0	2.9961	5.2	0.2441	3.3	0.63	1407.7	41.3	1406.6	39.4	1405.0	76.8	1405.0	76.8	100.2
Trial-2_0182	246	724110	1.6	13.5865	2.2	1.6297	2.9	0.1606	1.8	0.62	960.0	15.9	981.7	18.0	1030.6	45.5	1030.6	45.5	93.2
Trial-2_0183	79	54638	0.9	12.7962	3.8	1.8616	4.8	0.1728	3.0	0.62	1027.4	28.5	1067.6	31.9	1150.6	75.2	1150.6	75.2	89.3
Trial-2_0184	139	190973	3.1	7.7599	1.9	6.3864	2.4	0.3594	1.6	0.65	1979.5	27.0	2030.4	21.5	2082.6	32.8	2082.6	32.8	95.0
Trial-2_0185	307	43416	1.7	19.8842	4.1	0.2879	5.1	0.0415	3.1	0.60	262.2	7.9	256.9	11.7	208.5	95.8	262.2	7.9	NA
Trial-2_0186	12	75808	1.1	12.1528	8.5	2.3510	10.9	0.2072	6.8	0.62	1214.0	75.5	1227.9	78.0	1252.3	167.5	1252.3	167.5	96.9
Trial-2_0187	107	10216	1.0	20.0980	7.2	0.2623	8.9	0.0382	5.3	0.59	241.9	12.6	236.5	18.8	183.6	167.6	241.9	12.6	NA
Trial-2_0188	58	252280	0.5	11.1806	3.6	3.1333	4.6	0.2541	2.9	0.63	1459.5	38.2	1440.9	35.7	1413.6	68.7	1413.6	68.7	103.2
Trial-2_0189	77	89005	1.6	9.2074	2.8	4.7048	3.6	0.3142	2.3	0.64	1761.2	35.5	1768.1	30.1	1776.2	50.3	1776.2	50.3	99.2
Trial-2_0190	83	20224	0.8	17.4074	5.2	0.7636	6.5	0.0964	3.9	0.61	593.3	22.3	576.1	28.5	508.7	113.3	593.3	22.3	116.6
Trial-2_0191	71	866857	0.8	5.4120	2.1	12.8590	2.9	0.5047	1.9	0.66	2634.1	41.2	2669.4	27.0	2696.2	35.5	2696.2	35.5	97.7
Trial-2_0192	178	28663	0.8	17.2260	3.6	0.7681	4.5	0.0960	2.7	0.61	590.7	15.4	578.7	19.8	531.7	78.3	590.7	15.4	111.1
Trial-2_0193	31	15783	0.6	11.1029	4.8	3.0301	6.2	0.2440	3.9	0.63	1407.4	49.2	1415.2	47.0	1426.9	91.1	1426.9	91.1	98.6
Trial-2_0194	74	236282	1.0	12.3162	3.5	2.2965	4.5	0.2051	2.8	0.63	1202.8	31.0	1211.2	31.9	1226.1	69.0	1226.1	69.0	98.1
Trial-2_0195	21	11941	2.1	12.4806	6.4	2.3567	8.2	0.2133	5.1	0.63	1246.5	58.3	1229.6	58.6	1200.0	126.3	1200.0	126.3	103.9
Trial-2_0196	247	100493	1.5	18.4209	3.6	0.4421	4.5	0.0591	2.7	0.60	369.9	9.8	371.7	14.1	382.9	81.2	369.9	9.8	NA
Trial-2_0197	38	12329	1.6	13.4479	5.2	1.8209	6.7	0.1776	4.2	0.62	1053.8	40.4	1053.0	43.9	1051.3	105.7	1051.3	105.7	100.2
Trial-2_0198	95	266071	0.8	13.5643	3.4	1.7396	4.4	0.1711	2.7	0.62	1018.4	25.7	1023.3	28.3	1033.9	69.7	1033.9	69.7	98.5

Trial-2_0199	31	11746	1.2	12.7505	2.9	1.9850	5.5	0.1836	4.7	0.85	1086.4	46.7	1110.4	37.2	1157.7	57.9	1157.7	57.9	93.8
Trial-2_0200	11	18149	1.1	13.2862	1.5	1.8118	8.3	0.1746	8.2	0.98	1037.3	78.4	1049.7	54.5	1075.6	30.7	1075.6	30.7	96.4
Trial-2_0201	295	75598	0.8	19.5659	4.2	0.2927	5.3	0.0415	3.1	0.60	262.4	8.1	260.7	12.2	245.8	97.9	262.4	8.1	NA
Trial-2_0202	67	121485	0.6	5.2129	2.1	14.0993	2.8	0.5331	1.9	0.67	2754.3	41.5	2756.4	26.4	2757.9	34.2	2757.9	34.2	99.9
Trial-2_0203	102	400583	0.5	5.0744	1.8	13.8392	2.4	0.5093	1.6	0.66	2653.7	34.1	2738.8	22.3	2802.1	28.8	2802.1	28.8	94.7
Trial-2_0204	92	125880	1.3	10.8582	2.8	3.1121	3.7	0.2451	2.3	0.63	1413.1	29.4	1435.7	28.2	1469.4	53.9	1469.4	53.9	96.2
Trial-2_0205	37	39325	2.0	13.3978	5.3	1.9313	6.7	0.1877	4.2	0.62	1108.7	42.6	1092.0	45.0	1058.8	106.1	1058.8	106.1	104.7
Trial-2_0206	47	19251	1.2	13.0563	4.6	2.0007	5.9	0.1895	3.7	0.62	1118.4	37.8	1115.8	40.1	1110.6	92.6	1110.6	92.6	100.7
Trial-2_0207	35	31884	1.2	13.5620	5.8	1.7653	7.4	0.1736	4.6	0.62	1032.1	43.5	1032.8	47.8	1034.2	116.9	1034.2	116.9	99.8
Trial-2_0208	81	120988	1.2	12.9185	3.6	2.0044	4.6	0.1878	2.9	0.62	1109.4	29.1	1117.0	31.1	1131.7	71.4	1131.7	71.4	98.0
Trial-2_0209	19	85928	0.8	8.3108	5.2	5.7922	6.8	0.3491	4.4	0.64	1930.4	73.2	1945.2	59.0	1961.0	93.1	1961.0	93.1	98.4
Trial-2_0210	41	51698	0.6	9.9228	3.9	4.0693	5.1	0.2929	3.3	0.64	1655.8	47.7	1648.2	41.8	1638.5	73.3	1638.5	73.3	101.1
Trial-2_0211	40	110384	0.8	9.9024	4.0	3.9102	5.2	0.2808	3.3	0.64	1595.5	47.0	1615.8	42.3	1642.3	74.8	1642.3	74.8	97.2
Trial-2_0212	34	62563	1.8	11.9613	4.9	2.2655	6.3	0.1965	3.9	0.63	1156.7	41.7	1201.6	44.2	1283.3	94.9	1283.3	94.9	90.1
Trial-2_0213	203	71316	1.3	13.3920	2.4	1.7910	3.0	0.1740	1.9	0.62	1033.9	17.9	1042.2	19.6	1059.7	47.5	1059.7	47.5	97.6
Trial-2_0214	244	148532	1.8	8.3174	1.5	5.3319	2.0	0.3216	1.3	0.65	1797.7	19.9	1874.0	16.8	1959.6	26.8	1959.6	26.8	91.7
Trial-2_0215	62	369348	1.1	5.3838	2.3	13.0223	3.0	0.5085	2.0	0.66	2650.1	44.0	2681.3	28.8	2704.8	37.6	2704.8	37.6	98.0
Trial-2_0216	151	282613	0.9	8.5449	1.9	5.2769	2.5	0.3270	1.6	0.64	1824.0	25.6	1865.1	21.3	1911.3	34.3	1911.3	34.3	95.4
Trial-2_0217	74	67242	1.6	13.0152	3.9	1.9811	5.0	0.1870	3.1	0.62	1105.1	31.5	1109.1	33.7	1116.9	77.9	1116.9	77.9	98.9
Trial-2_0218	67	75531	1.0	5.2606	2.2	13.5438	2.9	0.5167	2.0	0.66	2685.4	43.0	2718.3	27.8	2742.9	36.2	2742.9	36.2	97.9
Trial-2_0219	883	488095	3.4	13.4203	1.1	1.6660	1.5	0.1622	0.9	0.63	968.7	8.2	995.7	9.2	1055.4	22.8	1055.4	22.8	91.8
Trial-2_0220	40	100633	1.4	13.6661	5.4	1.7031	6.9	0.1688	4.3	0.62	1005.5	40.1	1009.7	44.5	1018.8	110.3	1018.8	110.3	98.7
Trial-2_0221	53	238289	1.2	5.2529	2.5	13.6043	3.3	0.5183	2.2	0.66	2691.9	48.9	2722.6	31.7	2745.3	41.1	2745.3	41.1	98.1
Trial-2_0222	93	66066	1.8	13.4984	3.6	1.7806	4.5	0.1743	2.8	0.62	1035.9	27.0	1038.4	29.6	1043.7	71.9	1043.7	71.9	99.3
Trial-2_0223	614	68932	0.8	17.4885	2.2	0.5497	2.8	0.0697	1.7	0.60	434.5	7.1	444.8	10.1	498.5	49.2	434.5	7.1	87.2
Trial-2_0224	90	329354	1.0	5.4841	2.0	12.1398	2.6	0.4829	1.7	0.66	2539.7	36.5	2615.3	24.6	2674.3	32.5	2674.3	32.5	95.0
Trial-2_0225	214	166509	1.4	11.3961	2.0	2.6638	2.6	0.2202	1.6	0.63	1282.8	19.2	1318.5	19.3	1377.0	39.0	1377.0	39.0	93.2
Trial-2_0226	40	23615	1.8	13.2109	5.1	2.0057	6.5	0.1922	4.1	0.62	1133.1	42.4	1117.4	44.4	1087.0	102.6	1087.0	102.6	104.2
Trial-2_0227	24	18404	0.4	10.7925	5.4	3.0767	7.0	0.2408	4.5	0.63	1391.0	56.0	1426.9	54.1	1480.9	103.4	1480.9	103.4	93.9
Trial-2_0228	130	252946	1.5	13.6152	2.9	1.7414	3.7	0.1720	2.3	0.62	1022.9	21.7	1024.0	23.8	1026.3	58.5	1026.3	58.5	99.7
Trial-2_0229	116	116005	2.7	9.2456	2.3	4.5236	3.0	0.3033	1.9	0.64	1707.8	28.6	1735.3	24.7	1768.6	41.7	1768.6	41.7	96.6
Trial-2_0230	397	209751	3.8	12.6866	1.7	1.9195	2.1	0.1766	1.3	0.62	1048.5	12.8	1087.9	14.1	1167.7	32.7	1167.7	32.7	89.8
Trial-2_0231	75	79451	2.8	13.2369	3.9	1.8057	5.0	0.1733	3.1	0.62	1030.5	29.5	1047.5	32.6	1083.1	78.3	1083.1	78.3	95.2
Trial-2_0232	161	75672	2.7	12.6903	2.5	2.1351	3.2	0.1965	2.0	0.63	1156.6	21.1	1160.2	22.0	1167.1	49.2	1167.1	49.2	99.1
Trial-2_0233	179	25536	1.2	19.9767	5.4	0.2802	6.7	0.0406	4.0	0.60	256.5	10.1	250.8	15.0	197.7	125.8	256.5	10.1	NA
Trial-2_0234	49	71889	2.0	9.3136	3.5	4.7495	4.5	0.3208	2.9	0.64	1793.8	45.4	1776.0	37.9	1755.2	63.5	1755.2	63.5	102.2
Trial-2_0235	51	150146	0.7	5.4845	2.5	12.8781	3.4	0.5123	2.3	0.66	2666.3	49.3	2670.8	32.1	2674.2	42.1	2674.2	42.1	99.7
Trial-2_0236	85	74478	1.1	21.6118	8.5	0.2427	10.5	0.0380	6.3	0.59	240.7	14.8	220.6	20.9	11.8	204.0	240.7	14.8	NA
Trial-2_0237	718	1778812	1.4	9.9082	1.1	4.0285	1.4	0.2895	0.9	0.64	1639.0	12.9	1640.0	11.4	1641.2	20.1	1641.2	20.1	99.9
Trial-2_0238	200	115346	1.6	10.9306	2.0	3.0716	2.6	0.2435	1.6	0.63	1404.9	20.5	1425.6	19.7	1456.7	37.8	1456.7	37.8	96.4
Trial-2_0239	33	24267	3.4	12.1978	5.2	2.3584	6.6	0.2086	4.2	0.63	1221.6	46.5	1230.1	47.5	1245.1	101.4	1245.1	101.4	98.1
Trial-2_0240	201	25475	1.7	20.1999	5.5	0.2636	6.9	0.0386	4.1	0.60	244.3	9.9	237.6	14.6	171.8	129.6	244.3	9.9	NA
Trial-2_0241	381	166205	4.6	12.4947	1.7	2.0711	2.1	0.1877	1.3	0.63	1108.8	13.6	1139.3	14.6	1197.8	32.7	1197.8	32.7	92.6
Trial-2_0242	35	16029	3.0	13.3208	5.8	1.8321	7.4	0.1770	4.6	0.62	1050.6	44.4	1057.1	48.6	1070.4	116.5	1070.4	116.5	98.1
Trial-2_0243	74	100078	2.2	6.1741	2.3	10.3328	3.0	0.4627	2.0	0.66	2451.4	40.3	2465.0	27.8	2476.3	38.0	2476.3	38.0	99.0
Trial-2_0244	73	33476	1.7	12.9353	3.7	2.1482	4.7	0.2015	3.0	0.62	1183.6	32.0	1164.5	32.8	1129.1	73.6	1129.1	73.6	104.8
Trial-2_0245	16	18291	1.2	12.2232	7.7	2.3526	9.8	0.2086	6.1	0.62	1221.2	68.2	1228.4	70.1	1241.0	150.4	1241.0	150.4	98.4
Trial-2_0246	44	34686	1.8	13.4735	5.0	1.8834	6.4	0.1840	4.0	0.62	1089.0	39.6	1075.2	42.2	1047.5	100.3	1047.5	100.3	104.0
Trial-2_0247	606	1167554	5.2	9.2488	1.1	3.9002	1.4	0.2616	0.9	0.64	1498.1	12.2	1613.7	11.5	1768.0	20.0	1768.0	20.0	84.7
Trial-2_0248	115	10492	1.2	20.7601	7.4	0.2729	9.2	0.0411	5.5	0.60	259.6	13.9	245.0	20.0	107.6	174.1	259.6	13.9	NA
Trial-2_0249	146	37112	1.1	17.8285	4.2	0.6561	5.3	0.0848	3.2	0.61	525.0	16.1	512.3	21.2	455.9	93.1	525.0	16.1	115.1
Trial-2_0250	143	351110	1.0	11.1797	2.4	2.9563	3.1	0.2397	2.0	0.63	1385.2	24.5	1396.5	23.6	1413.7	46.1	1413.7	46.1	98.0
Trial-2_0251	586	462811	0.5	18.1277	2.7	0.3910	3.4	0.0514	2.1	0.60	323.1	6.5	335.1	9.8	418.9	61.0	323.1	6.5	NA
Trial-2_0252	144	167665	2.4	5.2843	1.5	13.1784	2.0	0.5051	1.3	0.66	2635.5	29.1	2692.5	19.1	2735.6	24.9	2735.6	24.9	96.3
Trial-2_0253	47	27203	1.1	12.4997	4.8	2.1061	6.2	0.1909	3.8	0.62	1126.4	39.6	1150.8	42.4	1197.1	95.1	1197.1	95.1	94.1
Trial-2_0254	344	155847	47.9	16.5677	2.6	0.7994	3.3	0.0961	2.0	0.61	591.2	11.2	596.5	14.7	616.4	55.7	591.2	11.2	95.9
Trial-2_0255	148	144949	15.6	13.3356	2.7	1.8230	3.5	0.1763	2.2	0.62	1046.9	21.0	1053.8	22.9	1068.2	55.0	1068.2	55.0	98.0
Trial-2_0256	306	201890	0.8	16.0514	2.5	0.9364	3.2	0.1090	2.0	0.61	667.0	12.4	671.0	15.7	684.4	53.8	667.0	12.4	97.5
Trial-2_0257	53	114489	1.2	5.4813	2.5	12.9454	3.3	0.5146	2.2	0.66	2676.4	48.0	2675.7	31.1	2675.2	40.8	2675.2	40.8	100.0
Trial-2_0258	318	138865	1.9	13.4801	2.0	1.7195	2.5	0.1681	1.6	0.62	1001.7	14.5	1015.8	16.2	1046.5	39.8	1046.5	39.8	95.7
Trial-2_0259	89	68155	0.5	13.5406	3.7	1.7299	4.8	0.1699	2.9	0.62	1011.5	27.6	1019.7	30.6	1037.4	75.4	1037.4	75.4	97.5
Trial-2_0260	38	52598	2.0	13.1686	5.3	1.9954	6.7	0.1906	4.2	0.62	1124.5	43.3	1113.9	45.6	1093.5	105.4	1093.5	105.4	102.8
Trial-2_0261	58	66082	1.6	13.0129	4.2	2.0779	5.3	0.1961	3.3	0.62	1154.4	35.2	1141.5	36.6	1117.2	83.2	1117.2	83.2	103.3
Trial-2_0262	183	291716	1.5	10.9651	2.1	2.8603	2.7	0.2275	1.7	0.63	1321.2	20.7	1371.5	20.6	1450.7	40.3	1450.7	40.3	91.1
Trial-2_0263	231	110932	3.2	13.5175	2.3	1.7497	2.9	0.1715	1.8	0.62	1020.5	17.2	1027.0	18.9	1040.9	46.3	1040.9	46.3	98.0
Trial-2_0264	332	22034	0.8	20.1558	4.3	0.2853	5.4	0.0417	3.2	0.60	263.4	8.3	254.8	12.2	176.9	101.1	263.4	8.3	NA
Trial-2_0265	104	28947	2.0	20.0971	7.5	0.2779	9.4	0.0405	5.6	0.60	256.0	14.1	249.0	20.7	183.7	175.7	256.0	14.1	NA
Trial-2_0266	115	103000	2.3	13.8991	3.2	1.7082	4.1	0.1722	2.5	0.62	1024.2	23.9	1011.6	26.0	984.4	64.7	984.4	64.7	104.0
Trial-2_0267	184	259852	2.4	9.2098	1.8	4.4998	2.4	0.3006	1.5	0.64	1694.1	22.5	1730.9	19.5	1775.7	32.9	1775.7	32.9	95.4
Trial-2_0268	1120	373162	2.8	12.6574	1.1	2.1576	1.4	0.1981	0.9	0.62	1164.9	9.2	1167.5	9.6	1172.3	21.5	1172.3	21.5	99.4
Trial-2_0269	69	80984	2.6	12.7141	3.8	2.1393	4.9	0.1973	3.1	0.62	1160.6	32.4	1161.6	33.9	1163.4	75.8	1163.4	75.8	99.8
Trial-2_0270	287	616514	3.5	13.5190	2.0	1.7154	2.6	0.1682	1.6	0.62	1002.2	15.1	1014.3	16.8	1040.6	41.3	1040.6	41.3	96.3
Trial-2_0271	114	144013	1.9	8.6851	2.2	5.2286	2.9	0.3294	1.9	0.64	1835.2	29.7	1857.3	24.6	1882.0	39.7	1882.0	39.7	97.5
Trial-2_0272	570	111361	16.9	16.6379	2.0	0.7253	2.6	0.0875	1.6	0.61	540.9	8.1	553.8	10.9	607.3	43.8	540.9	8.1	89.1
Trial-2_0273	39	109634	0.8	11.0119	4.4	3.0724	5.7	0.2454	3.6	0.63	1414.6	46.1	1425.8	43.9	1442.6	84.4	1442.6	84.4	98.1
Trial-2_0274	152	424065	1.1	10.1012	2.1	3.6328	2.8	0.2661	1.8	0.64	1521.2	24.0	1556.8	22.2	1605.3	40.0	1605.3	40.0	94.8
Trial-2_0275	129	30563	1.0	17.6670	4.7	0.5709	5.9	0.0732	3.6	0.61	455.1	15.7	458.6	21.8	476.1	103.9	455.1	15.7	95.6
Trial-2_0276	61	90103	1.1	11.0753	3.5	3.1353	4.6	0.2518	2.9	0.63	1448.0	37.6	1441.4	35.3	1431.7	67.7	1431.7	67.7	101.1
Trial-2_0277	51	32706	1.5	9.3299	3.3	4.6770	4.3	0.3165	2.8	0.64	1772.5	43.0	1763.1	36.1	1752.0	60.6	1752.0	60.6	101.2
Trial-2_0278	101	522881	1.6	9.5703	2.5	4.3075	3.2	0.2990	2.1	0.64	1686.3	30.5	1694.8	26.5	1705.3	45.4	1705.3	45.4	98.9
Trial-2_0279	23	19654	2.6	16.0533	8.9	0.9744	11.2	0.1135	6.9	0.61	692.8	45.0	690.7	56.2	684.2	189.6	692.8	45.0	101.3
Trial-2_0280	211	693235	2.8	11.0081	1.9	2.9930	2.5	0.2390	1.6	0.63	1381.3	19.7	1405.8	19.0	1443.3	36.8	1443.3	36.8	95.7
Trial-2_0281	397	147014	2.4	13.3290	1.8	1.6897	2.3	0.1633	1.4	0.62	975.3	12.8	1004.6	14.5	1069.2	35.9	1069.2	35.9	91.2
Trial-2_0282	39	48701	1.5	8.9542	3.8	5.0277	5.0	0.3265	3.2	0.64	1821.4	50.7	1824.0	42.2	1826.9	69.3	1826.9	69.3	99.7
Trial-2_0283	294	39625	0.7	20.5004	4.4	0.2625	5.5	0.0390	3.3	0.60	246.8	7.9	236.7	11.6	137.3	103.4	246.8	7.9	NA
Trial-2_0284	37	26051	0.9	10.2524	4.1	3.9314	5.4	0.2923	3.4	0.64	1653.2	49.8	1620.2	43.4	1577.6	77.2	1577.6	77.2	104.8
Trial-2_0285	757	2463242	9.3	9.3578	1.0	4.5319	1.3	0.3076	0.8	0.64	1728.8	12.7	1736.8	10.9	1746.6	18.4	1746.6	18.4	99.0
Trial-2_0286	128	203531	0.9	11.4527	2.5	2.8437	3.2	0.2362	2.0	0.63	1366.9	25.0	1367.1	24.1	1367.5	48.0	1367.5	48.0	100.0
Trial-2_0287	106	361112	0.7	5.2820	1.8	13.1662	2.3	0.5044	1.6	0.66	2632.6	33.7	2691.6	22.1	2736.3	28.8	2736.3	28.8	96.2
Trial-2_0288	129	35821	2.4	13.8684	3.0	1.6919	3.8	0.1702	2.4	0.62	1013.1	22.2	1005.5	24.4	989.0	60.8	989.0	60.8	102.4
Trial-2_0289	85	32416	1.4	20.5112	7.9	0.2760	9.9	0.0411	5.9	0.60	259.4	15.0	247.5	21.7	136.0	186.7	259.4	15.0	NA
Trial-2_0290	134	193282	1.3	8.5961	2.1	5.2467	2.7	0.3271	1.7	0.64	1824.3	27.7	1860.2	23.1	1900.6	37.3	1900.6	37.3	96.0
Trial-2_0291	35	19665	1.0	11.0027	5.0	2.9393	6.5	0.2346	4.1	0.63	1358.3	49.6	1392.1	49.0	1444.2	95.9	1444.2	95.9	94.1
Trial-2_0292	17	24734	3879.8	5.9892	5.4	9.8380	7.0	0.4273	4.4	0.64	2293.7	85.8	2419.7	64.2	2527.5	89.9	2527.5	89.9	90.8
Trial-2_0293	48	26590	1.0	13.2013	5.0	1.8910	6.3	0.1811	3.9	0.62	1072.7	38.8	1077.9	42.1	1088.5	99.7	1088.5	99.7	98.6
Trial-2_0294	40	24860	1.6	10.0186	4.0	4.1498	5.2	0.3015	3.3	0.64	1698.9	49.4	1664.2	42.5	1620.6	74.6	1620.6	74.6	104.8
Trial-2_0295	299	142526	1.8	13.0363	2.0	1.8200	2.6	0.1721	1.6	0.62	1023.5	15.0	1052.7	16.7	1113.6	39.9	1113.6	39.9	91.9
Trial-2_0296	58	170584	2.2	12.6761	4.2	2.2052	5.3	0.2027	3.3	0.62	1190.0	36.0	1182.7	37.2	1169.3	82.4	1169.3	82.4	101.8
Trial-2_0297	74	65579	1.2	13.0210	4.0	1.9954	5.1	0.1884	3.2	0.62	1112.9	32.2	1114.0	34.3	1116.0	79.3	1116.0	79.3	99.7
Trial-2_0298	85	9467	1.6	19.1861	6.3	0.4723	7.8	0.0657	4.7	0.60	410.3	18.6	392.7	25.5	290.7	143.6	410.3	18.6	141.1
Trial-2_0299	130	52525	1.3	11.1818	2.6	2.9665	3.4	0.2406	2.1	0.63	1389.7	26.8	1399.1	25.8	1413.4	50.6	1413.4	50.6	98.3
Trial-2_0300	241	239257	1.5	16.2230	2.9	0.8917	3.6	0.1049	2.2	0.61	643.2	13.5	647.3	17.3	661.7	61.5	643.2	13.5	97.2
Trial-2_0301	59	96112	0.5	5.0428	2.4	14.5668	3.2	0.5328	2.1	0.67	2753.1	47.0	2787.4	29.9	2812.3	38.4	2812.3	38.4	97.9
Trial-2_0302	699	140252	4.4	19.3140	2.8	0.2950	3.5	0.0413	2.1	0.60	261.0	5.4	262.5	8.2	275.5	65.1	261.0	5.4	NA
Trial-2_0303	105	96074	2.1	13.4991	3.4	1.8117	4.3	0.1774	2.7	0.62	1052.6	26.2	1049.7	28.4	1043.6	68.8	1043.6	68.8	100.9
Trial-2_0304	72	52551	1.3	11.0249	3.3	3.2606	4.2	0.2607	2.7	0.63	1493.5	35.4	1471.7	32.6	1440.4	62.0	1440.4	62.0	103.7
Trial-2_0305	95	73534	1.3	16.7296	4.9	0.8105	6.2	0.0983	3.7	0.61	604.7	21.6	602.7	28.0	595.4	106.1	604.7	21.6	101.6
Trial-2_0306	34	28253	0.9	12.9540	5.9	2.0059	7.5	0.1885	4.7	0.62	1113.0	47.6	1117.5	50.9	1126.3	117.2	1126.3	117.2	98.8
Trial-2_0307	74	68041	0.5	8.9278	2.8	5.0658	3.7	0.3280	2.4	0.64	1828.7	37.5	1830.4	31.1	1832.3	51.0	1832.3	51.0	99.8
Trial-2_0308	98	196833	0.7	13.2405	3.5	1.8903	4.5	0.1815	2.8	0.62	1075.3	27.4	1077.7	29.6	1082.5	70.0	1082.5	70.0	99.3
Trial-2_0309	22	12843	1.9	9.8864	5.8	3.9974	7.5	0.2866	4.8	0.63	1624.6	68.5	1633.7	61.1	1645.3	107.8	1645.3	107.8	98.7
Trial-2_0310	45	37872	1.5	12.6386	4.8	2.2288	6.1	0.2043	3.8	0.62	1198.4	41.8	1190.1	42.9	1175.2	94.6	1175.2	94.6	102.0
Trial-2_0311	86	30619	1.6	12.7957	3.6	2.0626	4.6	0.1914	2.9	0.62	1129.0	29.8	1136.5	31.7	1150.7	72.0	1150.7	72.0	98.1
Trial-2_0312	251	278861	1.3	8.0624	1.5	5.7121	2.0	0.3340	1.3	0.65	1857.8	20.5	1933.2	17.0	2015.0	26.6	2015.0	26.6	92.2
Trial-2_0313	24	41805	2.2	12.7539	6.3	2.1009	8.1	0.1943	5.0	0.62	1144.8	53.0	1149.1	55.8	1157.2	126.0	1157.2	126.0	98.9
Trial-2_0314	21	192629	1.3	9.8731	5.7	3.8796	7.4	0.2778	4.7	0.63	1580.3	65.9	1609.4	60.0	1647.8	106.7	1647.8	106.7	95.9
Trial-2_0315	119	17717	1.6	20.6191	7.0	0.2839	8.7	0.0425	5.2	0.59	268.0	13.6	253.7	19.5	123.6	164.9	268.0	13.6	NA
Trial-2_0316	16	17258	1.5	12.7216	8.2	2.0921	10.5	0.1930	6.5	0.62	1137.7	68.2	1146.2	72.2	1162.2	163.0	1162.2	163.0	97.9
Trial-2_0317	94	130825	1.7	18.4909	5.6	0.5843	7.0	0.0784	4.2	0.60	486.4	19.7	467.2	26.1	374.4	125.1	486.4	19.7	129.9
Trial-2_0318	128	378993	1.1	8.5677	2.1	5.4308	2.8	0.3375	1.8	0.64	1874.5	28.9	1889.7	23.6	1906.5	37.8	1906.5	37.8	98.3
Trial-2_0319	118	86097	2.7	12.3303	2.9	2.2952	3.7	0.2053	2.3	0.63	1203.5	25.6	1210.8	26.3	1223.9	57.0	1223.9	57.0	98.3
Trial-2_0320	31	147860	2.5	9.9509	4.7	3.9862	6.1	0.2877	3.9	0.64	1630.0	55.7	1631.4	49.4	1633.2	87.1	1633.2	87.1	99.8
Trial-2_0321	589	596709	2.0	16.3692	2.0	0.7767	2.5	0.0922	1.5	0.61	568.6	8.4	583.6	11.2	642.4	43.1	568.6	8.4	88.5
Trial-2_0322	764	126691	0.6	18.9023	2.5	0.3884	3.1	0.0532	1.8	0.60	334.4	6.0	333.2	8.7	324.6	56.0	334.4	6.0	NA
Trial-2_0323	166	448656	1.4	10.9179	2.2	3.1607	2.8	0.2503	1.8	0.63	1439.9	23.1	1447.6	21.8	1458.9	41.6	1458.9	41.6	98.7
Trial-2_0324	85	27162	1.6	12.6147	3.6	2.0959	4.6	0.1918	2.8	0.62	1130.9	29.5	1147.5	31.4	1179.0	70.6	1179.0	70.6	95.9
Trial-2_0325	41	48082	2.2	13.2900	5.3	1.8878	6.8	0.1820	4.2	0.62	1077.7	41.6	1076.8	44.9	1075.1	106.4	1075.1	106.4	100.2
Trial-2_0326	53	29904	1.5	12.8705	4.7	2.0353	5.9	0.1900	3.7	0.62	1121.3	38.0	1127.4	40.5	1139.1	92.6	1139.1	92.6	98.4
Trial-2_0327	158	106900	0.9	13.9270	2.9	1.6464	3.7	0.1663	2.3	0.62	991.7	21.0	988.2	23.3	980.3	59.0	980.3	59.0	101.2
Trial-2_0328	187	287511	1.2	10.9777	2.2	2.8722	2.8	0.2287	1.8	0.63	1327.6	21.2	1374.7	21.0	1448.6	41.2	1448.6	41.2	91.6
Trial-2_0329	962	238335	4.9	15.3769	1.6	0.8123	2.0	0.0906	1.2	0.61	559.0	6.5	603.8	9.0	775.4	33.0	559.0	6.5	72.1
Trial-2_0330	86	190562	1.2	11.2142	3.0	3.1031	3.9	0.2524	2.5	0.63	1450.8	32.2	1433.5	30.1	1407.9	58.1	1407.9	58.1	103.0
Trial-2_0331	19	19000	0.9	13.5472	7.7	1.8114	9.9	0.1780	6.1	0.62	1055.9	59.7	1049.6	64.7	1036.4	156.6	1036.4	156.6	101.9
Trial-2_0332	286	341906	0.8	18.2308	3.3	0.5527	4.1	0.0731	2.5	0.60	454.7	10.9	446.8	14.9	406.2	73.6	454.7	10.9	111.9
Trial-2_0333	390	1062254	5.0	8.6283	1.5	5.2966	2.0	0.3314	1.3	0.64	1845.4	20.2	1868.3	16.7	1893.9	27.0	1893.9	27.0	97.4
Trial-2_0334	166	104328	1.5	13.8289	2.8	1.6676	3.6	0.1673	2.2	0.62	996.9	20.4	996.3	22.6	994.8	56.7	994.8	56.7	100.2
Trial-2_0335	169	335497	0.9	8.3015	1.9	5.1122	2.5	0.3078	1.6	0.65	1729.9	24.3	1838.1	21.1	1963.0	33.8	1963.0	33.8	88.1
Trial-2_0336	8	4924	1.5	13.1273	1.9	1.9673	9.4	0.1873	9.2	0.98	1106.8	93.9	1104.4	63.5	1099.8	38.0	1099.8	38.0	100.6

Trial-2_0337	305	193660	3.6	12.7363	1.9	1.9767	2.4	0.1826	1.5	0.63	1081.1	15.2	1107.6	16.5	1159.9	37.8	1159.9	37.8	93.2
Trial-2_0338	231	183672	7.2	13.4538	2.3	1.7531	2.9	0.1711	1.8	0.62	1018.0	17.2	1028.3	18.9	1050.4	46.2	1050.4	46.2	96.9
Trial-2_0339	90	90525	0.6	13.5187	3.7	1.7304	4.7	0.1697	2.9	0.62	1010.2	27.4	1019.9	30.3	1040.7	74.5	1040.7	74.5	97.1
Trial-2_0340	61	75857	1.7	12.5582	4.2	2.1795	5.3	0.1985	3.3	0.62	1167.3	35.4	1174.5	37.0	1187.8	82.0	1187.8	82.0	98.3
Trial-2_0341	53	96136	1.3	16.3283	6.2	0.9058	7.8	0.1073	4.7	0.61	656.9	29.6	654.8	37.5	647.8	132.5	656.9	29.6	101.4
Trial-2_0342	584	885149	1.4	12.3469	1.4	1.9382	1.8	0.1736	1.1	0.63	1031.7	10.9	1094.4	12.2	1221.2	28.0	1221.2	28.0	84.5
Trial-2_0343	178	14188	1.2	20.3618	5.9	0.2642	7.4	0.0390	4.4	0.60	246.7	10.6	238.1	15.6	153.2	138.4	246.7	10.6	NA
Trial-2_0344	309	328150	3.8	9.2533	1.5	4.2529	1.9	0.2854	1.2	0.64	1618.6	17.5	1684.3	15.6	1767.1	26.7	1767.1	26.7	91.6
Trial-2_0345	295	236680	1.6	10.8293	1.8	2.7470	2.3	0.2158	1.5	0.63	1259.4	16.7	1341.3	17.1	1474.4	33.8	1474.4	33.8	85.4
Trial-2_0346	102	118699	2.8	13.8974	3.6	1.6742	4.5	0.1687	2.8	0.62	1005.2	26.3	998.8	28.9	984.7	72.6	984.7	72.6	102.1
Trial-2_0347	98	525329	1.1	9.7577	2.7	3.8747	3.5	0.2742	2.3	0.64	1562.1	31.4	1608.4	28.6	1669.6	50.5	1669.6	50.5	93.6
Trial-2_0348	77	173433	1.4	6.0530	2.3	10.7825	3.0	0.4734	2.0	0.66	2498.3	41.0	2504.6	27.9	2509.6	37.9	2509.6	37.9	99.5
Trial-2_0349	234	28742	1.4	19.0263	4.0	0.4343	5.0	0.0599	3.0	0.60	375.2	11.1	366.2	15.5	309.8	91.8	375.2	11.1	NA
Trial-2_0350	162	395856	1.5	9.8080	2.1	3.9763	2.7	0.2828	1.7	0.64	1605.7	24.5	1629.4	21.9	1660.0	38.5	1660.0	38.5	96.7
Trial-2_0351	60	41126	1.2	11.8100	3.8	2.6695	5.0	0.2287	3.1	0.63	1327.5	37.4	1320.1	36.6	1308.1	74.7	1308.1	74.7	101.5
Trial-2_0352	54	26707	2.2	13.3771	4.7	1.8510	6.0	0.1796	3.7	0.62	1064.7	36.7	1063.8	39.6	1061.9	94.5	1061.9	94.5	100.3
Trial-2_0353	1799	33238	1.3	21.0322	3.0	0.1011	3.7	0.0154	2.2	0.59	98.7	2.1	97.8	3.4	76.7	70.7	98.7	2.1	NA
Trial-2_0354	146	10686	2.6	21.9210	7.7	0.1769	9.6	0.0281	5.7	0.59	178.8	10.0	165.4	14.6	-22.5	187.1	178.8	10.0	NA
Trial-2_0355	138	51391	1.7	19.5578	6.6	0.2762	8.2	0.0392	4.9	0.60	247.7	11.9	247.6	18.1	246.7	152.4	247.7	11.9	NA
Trial-2_0356	26	13417	1.7	12.8787	6.6	2.0697	8.4	0.1933	5.3	0.62	1139.3	55.0	1138.8	57.9	1137.9	131.4	1137.9	131.4	100.1
Trial-2_0357	32	17204	1.5	13.1805	5.9	1.9225	7.6	0.1838	4.7	0.62	1087.6	47.1	1088.9	50.6	1091.6	118.8	1091.6	118.8	99.6
Trial-2_0358	1431	19961	1.6	16.4003	1.2	0.8456	1.5	0.1006	0.9	0.63	617.8	5.5	622.3	6.9	638.4	25.0	617.8	5.5	96.8
Trial-2_0359	198	142883	2.6	12.6827	2.3	2.0760	2.9	0.1910	1.8	0.63	1126.5	19.1	1140.9	20.2	1168.3	45.6	1168.3	45.6	96.4
Trial-2_0360	149	20603	1.1	18.7266	4.7	0.5071	5.9	0.0689	3.6	0.60	429.4	14.8	416.5	20.2	345.8	106.8	429.4	14.8	124.2
Trial-2_0361	114	39682	4.7	13.8710	3.3	1.6972	4.2	0.1707	2.6	0.62	1016.2	24.4	1007.5	26.7	988.6	66.7	988.6	66.7	102.8
Trial-2_0362	9	8130	1.3	12.5619	1.2	2.1576	8.9	0.1966	8.8	0.99	1156.9	93.0	1167.5	61.6	1187.3	23.2	1187.3	23.2	97.4
Trial-2_0363	151	81413	1.1	17.2290	3.9	0.7217	4.9	0.0902	3.0	0.61	556.6	16.0	551.7	21.0	531.4	85.8	556.6	16.0	104.8
Trial-2_0364	88	36548	1.4	12.9838	3.5	2.0435	4.5	0.1924	2.8	0.62	1134.5	29.5	1130.1	31.0	1121.7	70.8	1121.7	70.8	101.1
Trial-2_0365	39	16688	1.3	13.3792	5.4	1.8429	6.9	0.1788	4.3	0.62	1060.5	42.2	1060.9	45.6	1061.6	109.3	1061.6	109.3	99.9
Trial-2_0366	27	19585	1.9	13.4989	7.0	1.7770	8.9	0.1740	5.5	0.62	1034.0	52.8	1037.1	58.1	1043.6	141.8	1043.6	141.8	99.1
Trial-2_0367	223	117543	3.1	12.3758	2.3	2.0864	3.0	0.1873	1.8	0.63	1106.6	18.8	1144.3	20.3	1216.7	45.3	1216.7	45.3	91.0
Trial-2_0368	128	26109	1.8	13.1191	3.1	1.8810	4.0	0.1790	2.5	0.62	1061.4	24.4	1074.4	26.5	1101.0	62.6	1101.0	62.6	96.4
Trial-2_0369	32	25718	1.1	7.8255	3.8	6.7768	5.0	0.3846	3.3	0.65	2097.9	58.6	2082.7	44.6	2067.7	67.5	2067.7	67.5	101.5
Trial-2_0370	169	180669	2.3	13.4307	2.7	1.7834	3.4	0.1737	2.1	0.62	1032.6	20.3	1039.4	22.2	1053.9	53.9	1053.9	53.9	98.0
Trial-2_0371	51	108006	1.5	5.7020	2.8	11.1692	3.8	0.4619	2.5	0.66	2447.9	51.0	2537.3	35.3	2609.6	47.4	2609.6	47.4	93.8
Trial-2_0372	7	2024	1.3	13.5563	3.5	1.7790	3.7	0.1749	1.3	0.35	1039.1	12.6	1037.8	24.1	1035.1	70.2	1035.1	70.2	100.4
Trial-2_0373	20	16019	1.3	13.2557	8.0	1.8594	10.1	0.1788	6.3	0.62	1060.2	61.4	1066.8	67.1	1080.2	160.1	1080.2	160.1	98.1
Trial-2_0374	73	882342	0.7	5.1938	2.2	13.5239	2.9	0.5094	1.9	0.67	2654.2	41.9	2717.0	27.4	2763.9	35.5	2763.9	35.5	96.0
Trial-2_0375	642	549250	3.4	9.2163	0.9	4.6653	1.2	0.3118	0.8	0.65	1749.8	12.3	1761.0	10.3	1774.4	17.1	1774.4	17.1	98.6
Trial-2_0376	446	114824	1.6	16.3197	2.2	0.8360	2.8	0.0990	1.7	0.61	608.3	10.0	616.9	13.1	648.9	48.2	608.3	10.0	93.7
Trial-2_0377	83	97467	1.0	13.6703	3.9	1.7635	5.0	0.1748	3.1	0.62	1038.7	29.9	1032.1	32.5	1018.1	79.6	1018.1	79.6	102.0
Trial-2_0378	215	28535	1.2	20.5158	5.5	0.2597	6.8	0.0386	4.0	0.59	244.4	9.7	234.4	14.2	135.5	128.2	244.4	9.7	NA
Trial-2_0379	64	24382	1.7	13.4916	4.3	1.8944	5.5	0.1854	3.4	0.62	1096.2	34.4	1079.1	36.5	1044.7	86.9	1044.7	86.9	104.9
Trial-2_0380	132	51558	1.8	9.0133	2.2	4.8129	2.9	0.3146	1.8	0.64	1763.4	28.5	1787.1	24.1	1815.0	40.0	1815.0	40.0	97.2
Trial-2_0381	64	212898	0.7	11.1247	3.5	3.0467	4.5	0.2458	2.9	0.63	1416.9	36.4	1419.4	34.6	1423.2	67.0	1423.2	67.0	99.6
Trial-2_0382	69	345368	2.7	12.2828	3.7	2.3498	4.7	0.2093	3.0	0.63	1225.2	33.1	1227.5	33.7	1231.5	72.5	1231.5	72.5	99.5
Trial-2_0383	244	119281	4.3	17.5731	4.4	0.3642	5.5	0.0464	3.3	0.60	292.5	9.5	315.4	14.9	487.8	97.1	292.5	9.5	NA
Trial-2_0384	102	172196	1.4	13.6080	3.1	1.6989	4.0	0.1677	2.5	0.62	999.3	22.8	1008.1	25.4	1027.4	63.1	1027.4	63.1	97.3
Trial-2_0385	35	69094	0.9	10.9398	4.4	3.0609	5.7	0.2429	3.6	0.63	1401.5	45.2	1423.0	43.5	1455.1	83.9	1455.1	83.9	96.3
Trial-2_0386	167	221917	19.4	12.4651	2.2	2.1490	2.8	0.1943	1.8	0.63	1144.5	18.4	1164.7	19.5	1202.5	43.3	1202.5	43.3	95.2
Trial-2_0387	167	793332	1.4	13.1329	2.3	1.9149	3.0	0.1824	1.8	0.62	1080.0	18.3	1086.3	19.7	1098.9	46.3	1098.9	46.3	98.3
Trial-2_0388	60	144324	1.1	13.1627	4.2	1.7548	5.3	0.1675	3.3	0.62	998.4	30.2	1028.9	34.2	1094.4	83.3	1094.4	83.3	91.2
Trial-2_0389	67	146574	0.9	5.4303	2.1	12.5761	2.8	0.4953	1.8	0.66	2593.6	39.3	2648.4	26.1	2690.6	34.3	2690.6	34.3	96.4
Trial-2_0390	78	106293	2.6	9.4230	2.5	4.3951	3.3	0.3004	2.1	0.64	1693.1	31.2	1711.4	27.0	1733.9	46.1	1733.9	46.1	97.7
Trial-2_0391	228	309814	2.4	13.7242	2.1	1.6267	2.7	0.1619	1.7	0.62	967.4	15.1	980.6	17.0	1010.2	43.1	1010.2	43.1	95.8
Trial-2_0392	31	63830	2.3	13.1665	5.5	1.8465	7.0	0.1763	4.3	0.62	1046.9	41.7	1062.2	45.9	1093.8	109.4	1093.8	109.4	95.7
Trial-2_0393	121	89186	1.8	12.8346	2.7	2.0577	3.5	0.1915	2.2	0.62	1129.7	22.5	1134.8	23.8	1144.7	54.0	1144.7	54.0	98.7
Trial-2_0394	38	22378	1.3	13.1600	5.2	1.8587	6.6	0.1774	4.1	0.62	1052.8	39.7	1066.5	43.5	1094.8	103.6	1094.8	103.6	96.2
Trial-2_0395	250	400138	2.0	8.7371	1.5	4.8033	1.9	0.3044	1.2	0.64	1713.0	18.6	1785.5	16.2	1871.3	26.6	1871.3	26.6	91.5
Trial-2_0396	506	532384	0.5	12.6115	1.4	1.9188	1.8	0.1755	1.1	0.62	1042.4	10.9	1087.7	12.2	1179.5	28.2	1179.5	28.2	88.4
Trial-2_0397	147	30428	0.8	21.2579	6.0	0.2003	7.4	0.0309	4.4	0.59	196.1	8.5	185.4	12.6	51.3	142.9	196.1	8.5	NA
Trial-2_0398	22	37001	2.7	9.9282	5.0	3.8970	6.4	0.2806	4.1	0.64	1594.4	58.0	1613.1	52.2	1637.5	92.4	1637.5	92.4	97.4
Trial-2_0399	49	11558	1.5	13.2959	4.8	1.7761	6.1	0.1713	3.7	0.62	1019.1	35.3	1036.7	39.4	1074.2	95.7	1074.2	95.7	94.9
Trial-2_0400	60	53218	0.8	13.3679	3.8	1.9095	4.8	0.1851	3.0	0.62	1094.9	30.2	1084.4	32.1	1063.3	75.9	1063.3	75.9	103.0
Trial-2_0401	107	156959	2.2	13.1811	2.9	1.9657	3.7	0.1879	2.3	0.62	1110.1	23.6	1103.8	25.0	1091.5	58.2	1091.5	58.2	101.7
Trial-2_0402	698	1140473	14.3	12.1510	1.1	1.9377	1.4	0.1708	0.9	0.63	1016.3	8.5	1094.2	9.6	1252.6	21.9	1252.6	21.9	81.1
Trial-2_0403	176	77983	0.9	20.1400	5.3	0.2679	6.6	0.0391	3.9	0.60	247.4	9.6	241.0	14.2	178.7	123.7	247.4	9.6	NA
Trial-2_0404	358	245522	1.6	13.3289	1.7	1.6790	2.2	0.1623	1.4	0.62	969.6	12.3	1000.6	14.0	1069.2	34.7	1069.2	34.7	90.7
Trial-2_0405	268	147630	1.0	10.7090	1.7	2.9700	2.2	0.2307	1.4	0.63	1338.1	16.8	1400.0	16.7	1495.6	32.1	1495.6	32.1	89.5
Trial-2_0406	359	445885	2.4	10.8820	1.3	2.9947	1.7	0.2364	1.1	0.64	1367.7	13.3	1406.3	12.9	1465.2	24.9	1465.2	24.9	93.3
Trial-2_0407	105	51188	2.3	13.3302	3.2	1.8445	4.1	0.1783	2.5	0.62	1057.8	24.7	1061.5	26.9	1069.0	64.3	1069.0	64.3	99.0
Trial-2_0408	36	63942	0.6	5.5266	2.8	12.7369	3.8	0.5105	2.5	0.66	2658.9	54.5	2660.4	35.5	2661.5	46.8	2661.5	46.8	99.9
Trial-2_0409	47	86687	1.4	3.0191	2.0	33.7644	2.7	0.7393	1.8	0.68	3568.0	50.2	3603.2	26.4	3622.8	30.0	3622.8	30.0	98.5
Trial-2_0410	105	142509	0.7	11.0048	2.5	3.0248	3.2	0.2414	2.0	0.63	1394.1	25.6	1413.9	24.6	1443.8	47.6	1443.8	47.6	96.6
Trial-2_0411	118	755888	1.5	11.7279	2.5	2.6045	3.2	0.2215	2.0	0.63	1290.0	23.4	1301.9	23.3	1321.6	47.7	1321.6	47.7	97.6
Trial-2_0412	417	119284	1.0	17.4074	2.7	0.4872	3.5	0.0615	2.1	0.61	384.8	7.9	403.0	11.5	508.7	60.3	384.8	7.9	NA
Trial-2_0413	108	227860	2.8	9.3958	2.2	4.4327	2.9	0.3021	1.9	0.64	1701.5	28.0	1718.5	24.2	1739.1	41.1	1739.1	41.1	97.8
Trial-2_0414	112	192579	0.8	11.0993	2.4	3.1143	3.2	0.2507	2.0	0.63	1442.1	25.8	1436.2	24.3	1427.5	46.7	1427.5	46.7	101.0
Trial-2_0415	270	17057	0.9	20.5446	4.4	0.2550	5.5	0.0380	3.3	0.60	240.4	7.7	230.7	11.3	132.2	103.2	240.4	7.7	NA
Trial-2_0416	98	72982	2.0	12.9643	3.1	1.9979	4.0	0.1879	2.5	0.62	1109.8	25.3	1114.8	27.0	1124.7	62.2	1124.7	62.2	98.7
Trial-2_0417	58	52624	1.8	13.3754	4.0	1.9406	5.2	0.1883	3.2	0.62	1111.9	32.9	1095.2	34.7	1062.2	81.5	1062.2	81.5	104.7
Trial-2_0418	21	22148	0.3	6.2215	4.0	10.2576	5.3	0.4628	3.5	0.66	2452.1	71.2	2458.3	49.1	2463.4	67.4	2463.4	67.4	99.5
Trial-2_0419	32	21479	1.5	12.3566	5.3	2.3322	6.8	0.2090	4.3	0.63	1223.5	47.6	1222.1	48.6	1219.7	104.8	1219.7	104.8	100.3
Trial-2_0420	214	40249	1.0	18.9005	3.8	0.4365	4.8	0.0598	2.9	0.60	374.6	10.5	367.8	14.8	324.9	87.2	374.6	10.5	NA
Trial-2_0421	285	212497	2.0	12.3926	1.8	2.1788	2.3	0.1958	1.4	0.63	1152.9	14.9	1174.3	15.7	1214.0	34.5	1214.0	34.5	95.0
Trial-2_0422	98	112504	1.7	13.3143	3.0	1.9188	3.9	0.1853	2.4	0.62	1095.8	24.5	1087.6	26.0	1071.4	61.3	1071.4	61.3	102.3
Trial-2_0423	64	136188	1.4	11.0878	3.4	3.0852	4.4	0.2481	2.8	0.63	1428.7	35.3	1429.0	33.5	1429.5	64.7	1429.5	64.7	99.9
Trial-2_0424	153	111679	3.4	13.4900	2.8	1.5630	3.5	0.1529	2.2	0.62	917.3	18.7	955.7	21.9	1045.0	56.1	1045.0	56.1	87.8
Trial-2_0425	128	37255	0.7	20.9437	7.5	0.1967	9.3	0.0299	5.5	0.59	189.8	10.3	182.3	15.5	86.8	177.6	189.8	10.3	NA
Trial-2_0426	80	55878	0.8	18.5812	6.0	0.5097	7.5	0.0687	4.5	0.60	428.3	18.8	418.3	25.8	363.4	135.5	428.3	18.8	117.9
Trial-2_0427	295	379722	4.4	5.1877	1.1	12.7995	1.4	0.4816	1.0	0.66	2534.1	20.0	2665.0	13.6	2765.9	17.7	2765.9	17.7	91.6
Trial-2_0428	215	105521	2.1	12.7188	2.0	2.0951	2.6	0.1933	1.6	0.63	1139.0	16.8	1147.2	17.7	1162.7	39.9	1162.7	39.9	98.0
Trial-2_0429	31	24671	0.6	23.2794	13.2	0.2409	16.4	0.0407	9.7	0.59	257.0	24.5	219.1	32.4	-170.1	331.2	257.0	24.5	NA
Trial-2_0430	182	40533	0.9	20.8622	5.2	0.2773	6.5	0.0420	3.9	0.59	264.9	10.0	248.5	14.3	96.0	123.9	264.9	10.0	NA
Trial-2_0431	15	6162	1.2	13.5179	8.5	1.8061	10.8	0.1771	6.7	0.62	1050.9	64.6	1047.7	70.6	1040.8	171.6	1040.8	171.6	101.0
Trial-2_0432	259	438089	1.9	9.3506	1.5	4.1539	1.9	0.2817	1.2	0.64	1599.9	17.6	1665.0	15.9	1748.0	27.3	1748.0	27.3	91.5
Trial-2_0433	22	19035	1.5	4.7823	3.7	16.5935	4.9	0.5755	3.3	0.67	2930.5	77.7	2911.6	47.4	2898.6	59.8	2898.6	59.8	101.1
Trial-2_0434	34	9924	0.9	16.5502	7.1	0.8600	9.0	0.1032	5.5	0.61	633.3	32.9	630.1	42.2	618.7	154.4	633.3	32.9	102.4
Trial-2_0435	483	369736	2.7	11.6906	1.4	2.2685	1.8	0.1923	1.1	0.63	1134.0	11.8	1202.5	12.7	1327.8	27.1	1327.8	27.1	85.4
Trial-2_0436	117	72633	2.1	13.5953	3.0	1.8211	3.9	0.1796	2.4	0.62	1064.6	23.6	1053.1	25.4	1029.3	61.5	1029.3	61.5	103.4
Trial-2_0437	394	309024	2.5	11.2613	1.4	2.5502	1.8	0.2083	1.2	0.63	1219.7	12.9	1286.5	13.4	1399.8	27.3	1399.8	27.3	87.1
Trial-2_0438	453	228742	1.0	9.8634	1.3	4.0309	1.6	0.2884	1.0	0.63	1633.3	15.0	1640.5	13.3	1649.6	23.5	1649.6	23.5	99.0
Trial-2_0439	98	127273	2.1	13.8258	3.3	1.6890	4.2	0.1694	2.6	0.62	1008.6	24.4	1004.4	26.8	995.2	67.1	995.2	67.1	101.4
Trial-2_0440	48	26533	3.9	19.2482	8.0	0.4779	10.0	0.0667	6.0	0.60	416.4	24.3	396.7	32.9	283.4	183.7	416.4	24.3	146.9
Trial-2_0441	309	407973	7.1	12.4018	1.7	2.1379	2.2	0.1923	1.4	0.63	1133.8	14.5	1161.1	15.4	1212.5	34.2	1212.5	34.2	93.5
Trial-2_0442	15	29343	1.0	6.6029	4.9	8.9935	6.5	0.4307	4.3	0.66	2308.8	83.0	2337.3	59.7	2362.3	84.2	2362.3	84.2	97.7
Trial-2_0443	212	127919	3.0	13.3271	2.2	1.8030	2.8	0.1743	1.7	0.62	1035.6	16.4	1046.6	18.0	1069.5	43.4	1069.5	43.4	96.8
Trial-2_0444	44	20808	1.2	11.3467	3.8	2.9953	5.0	0.2465	3.1	0.63	1420.4	40.0	1406.4	37.8	1385.3	73.9	1385.3	73.9	102.5
Trial-2_0445	161	453123	2.4	13.6721	2.5	1.6906	3.2	0.1676	2.0	0.62	999.1	18.4	1005.0	20.4	1017.9	50.7	1017.9	50.7	98.2
Trial-2_0446	256	214397	2.9	9.3813	1.5	4.2441	1.9	0.2888	1.2	0.64	1635.4	17.7	1682.6	15.7	1742.0	27.0	1742.0	27.0	93.9
Trial-2_0447	391	54075	0.9	19.4911	3.3	0.3106	4.1	0.0439	2.5	0.60	277.0	6.7	274.7	9.9	254.6	75.7	277.0	6.7	NA
Trial-2_0448	81	109657	1.9	13.7019	3.5	1.7270	4.5	0.1716	2.8	0.62	1021.1	26.5	1018.6	29.1	1013.5	71.8	1013.5	71.8	100.8
Trial-2_0449	133	63639	1.8	13.2741	2.6	1.9121	3.3	0.1841	2.1	0.62	1089.2	20.9	1085.3	22.3	1077.5	52.6	1077.5	52.6	101.1
Trial-2_0450	78	42893	1.5	11.2413	3.0	2.9413	3.9	0.2398	2.5	0.63	1385.7	30.6	1392.6	29.5	1403.2	57.8	1403.2	57.8	98.7
Trial-2_0451	142	788363	1.2	8.2338	1.8	5.4862	2.4	0.3276	1.6	0.65	1826.8	24.8	1898.4	20.8	1977.6	32.9	1977.6	32.9	92.4
Trial-2_0452	90	87408	2.0	5.4313	2.3	11.1578	3.0	0.4395	2.0	0.65	2348.5	38.7	2536.4	28.2	2690.3	38.0	2690.3	38.0	87.3
Trial-2_0453	543	169581	1.0	20.3802	3.4	0.2092	4.3	0.0309	2.5	0.59	196.3	4.9	192.9	7.5	151.1	80.8	196.3	4.9	NA
Trial-2_0454	332	235060	2.0	12.9258	1.7	1.9092	2.2	0.1790	1.4	0.62	1061.4	13.6	1084.3	14.8	1130.6	34.7	1130.6	34.7	93.9
Trial-2_0455	76	37696	1.3	13.6304	3.6	1.7756	4.5	0.1755	2.8	0.62	1042.5	27.1	1036.6	29.5	1024.0	72.0	1024.0	72.0	101.8
Trial-2_0456	201	179335	1.2	9.3496	1.7	4.2856	2.3	0.2906	1.4	0.64	1644.5	20.9	1690.6	18.5	1748.2	31.7	1748.2	31.7	94.1
Trial-2_0457	175	119564	2.3	20.3871	5.5	0.2568	6.8	0.0380	4.1	0.59	240.2	9.6	232.1	14.2	150.3	128.9	240.2	9.6	NA
Trial-2_0458	59	170205	1.5	5.2665	2.2	13.5623	3.0	0.5180	2.0	0.66	2690.8	43.2	2719.6	27.9	2741.1	36.3	2741.1	36.3	98.2
Trial-2_0459	198	28370	1.5	21.0262	5.4	0.2648	6.7	0.0404	4.0	0.59	255.2	10.0	238.5	14.3	77.4	129.0	255.2	10.0	NA
Trial-2_0460	58	36073	1.3	10.9901	3.4	3.0967	4.4	0.2468	2.8	0.63	1422.1	35.7	1431.9	34.0	1446.4	65.4	1446.4	65.4	98.3
Trial-2_0461	70	144800	0.6	8.8664	2.7	4.9887	3.5	0.3208	2.2	0.64	1793.6	35.0	1817.4	29.4	1844.8	48.2	1844.8	48.2	97.2
Trial-2_0462	306	224625	1.8	19.0863	3.4	0.3857	4.2	0.0534	2.5	0.60	335.3	8.2	331.2	11.8	302.6	76.4	335.3	8.2	NA
Trial-2_0463	112	43983	2.7	13.9449	3.2	1.6362	4.1	0.1655	2.5	0.62	987.2	23.0	984.3	25.6	977.7	65.0	977.7	65.0	101.0
Trial-2_0464	71	19533	1.0	21.1700	8.3	0.2479	10.4	0.0381	6.2	0.59	240.8	14.6	224.8	20.9	61.2	199.2	240.8	14.6	NA
Trial-2_0465	184	104315	1.7	11.9951	2.6	2.0716	3.4	0.1802	2.1	0.63	1068.2	20.8	1139.5	23.1	1277.8	51.4	1277.8	51.4	83.6
Trial-2_0466	95	51238	1.3	13.8200	3.3	1.6800	4.2	0.1684	2.6	0.62	1003.2	24.0	1001.0	26.5	996.1	66.3	996.1	66.3	100.7
Trial-2_0467	120	53566	0.4	11.0080	2.4	2.9801	3.1	0.2379	2.0	0.63	1375.9	24.5	1402.5	23.8	1443.3	46.1	1443.3	46.1	95.3
Trial-2_0468	11	31981	0.9	20.2276	6.4	0.2672	6.7	0.0392	2.2	0.32	247.9	5.3	240.5	14.4	168.6	149.4	247.9	5.3	NA
Trial-2_0469	142	317183	1.9	5.3516	1.5	12.1694	2.0	0.4723	1.3	0.66	2493.8	27.1	2617.6	18.5	2714.7	24.3	2714.7	24.3	91.9
Trial-2_0470	26	40505	1.3	7.7205	4.0	6.7752	5.2	0.3794	3.4	0.65	2073.3	59.9	2082.5	46.1	2091.5	69.7	2091.5	69.7	99.1
Trial-2_0471	70	37227	2.0	13.2274	3.7	1.8966	4.7	0.1820	2.9	0.62	1077.6	28.9	1079.9	31.1	1084.5	73.5	1084.5	73.5	99.4
Trial-2_0472	223	846049	1.4	9.0586	1.9	4.7292	2.5	0.3107	1.6	0.64	1744.2	24.0	1772.4	20.6	1805.9	34.4	1805.9	34.4	96.6
Trial-2_0473	212	207011	0.9	11.2064	1.9	2.7939	2.4	0.2271	1.5	0.63	1319.2	18.1	1353.9	18.0	1409.2	35.7	1409.2	35.7	93.6
Trial-2_0474	154	134296	2.0	11.9585	2.2	2.5266	2.9	0.2191	1.8	0.63	1277.3	21.0	1279.7	21.0	1283.8	43.6	1283.8	43.6	99.5

Trial-2_0475	28	32148	0.9	13.2650	6.0	1.8414	7.6	0.1772	4.7	0.62	1051.4	45.9	1060.4	50.3	1078.8	120.2	1078.8	120.2	97.5
Trial-2_0476	258	585908	3.2	13.2823	2.0	1.7503	2.5	0.1686	1.6	0.62	1004.4	14.6	1027.3	16.3	1076.2	39.7	1076.2	39.7	93.3
Trial-2_0477	79	33045	1.3	13.0633	3.3	1.9726	4.3	0.1869	2.7	0.62	1104.5	26.9	1106.2	28.7	1109.5	66.4	1109.5	66.4	99.6
Trial-2_0478	210	31235	1.1	18.6840	3.7	0.4960	4.6	0.0672	2.8	0.60	419.3	11.3	409.0	15.5	351.0	83.4	419.3	11.3	119.5
Trial-2_0479	50	202551	1.2	8.8141	3.0	5.2223	3.9	0.3338	2.5	0.64	1857.0	40.2	1856.3	33.1	1855.5	53.8	1855.5	53.8	100.1
Trial-2_0480	18	10086	1.7	10.8949	6.3	3.1215	8.1	0.2467	5.1	0.63	1421.2	64.9	1438.0	62.3	1462.9	119.6	1462.9	119.6	97.1
Trial-2_0481	90	86887	0.6	9.0272	2.4	4.8401	3.2	0.3169	2.1	0.64	1774.5	31.8	1791.9	26.9	1812.2	44.5	1812.2	44.5	97.9
Trial-2_0482	221	150414	1.9	10.4876	1.8	2.9981	2.3	0.2280	1.5	0.63	1324.2	17.4	1407.1	17.4	1535.0	33.3	1535.0	33.3	86.3
Trial-2_0483	111	103738	1.2	18.5921	5.1	0.5105	6.4	0.0688	3.8	0.60	429.2	15.9	418.8	21.8	362.1	114.5	429.2	15.9	118.5
Trial-2_0484	357	856896	1.4	11.9471	1.6	2.2315	2.0	0.1934	1.3	0.63	1139.5	13.3	1191.0	14.2	1285.6	30.7	1285.6	30.7	88.6
Trial-2_0485	18	26603	-19.3	16.2132	4.0	0.9053	5.1	0.1065	3.2	0.62	652.1	19.9	654.6	24.8	663.0	86.1	652.1	19.9	98.4
Trial-2_0486	141	82234	2.5	13.8206	2.7	1.6599	3.5	0.1664	2.2	0.62	992.2	19.9	993.3	22.1	996.0	55.5	996.0	55.5	99.6
Trial-2_0487	148	151464	1.9	20.9442	5.7	0.2559	7.1	0.0389	4.2	0.59	245.8	10.2	231.3	14.8	86.7	136.4	245.8	10.2	NA
Trial-2_0488	53	50195	1.7	13.0755	4.2	1.9574	5.4	0.1856	3.4	0.62	1097.7	33.9	1101.0	36.4	1107.6	84.9	1107.6	84.9	99.1
Trial-2_0489	136	247461	1.6	9.4152	2.0	4.2805	2.6	0.2923	1.6	0.64	1653.0	23.9	1689.6	21.1	1735.4	36.0	1735.4	36.0	95.3
Trial-2_0490	47	89412	0.8	12.6397	4.3	2.1113	5.5	0.1935	3.4	0.62	1140.6	35.8	1152.5	37.9	1175.1	85.0	1175.1	85.0	97.1
Trial-2_0491	29	45360	2.1	13.5866	5.9	1.7347	7.5	0.1709	4.7	0.62	1017.3	43.8	1021.5	48.5	1030.6	119.8	1030.6	119.8	98.7
Trial-2_0492	127	123729	2.1	9.3880	2.0	4.4774	2.6	0.3049	1.7	0.64	1715.3	25.4	1726.8	21.9	1740.7	37.1	1740.7	37.1	98.5
Trial-2_0493	20	38133	1.0	12.9351	6.8	2.0158	8.7	0.1891	5.4	0.62	1116.5	55.3	1120.8	58.8	1129.2	135.2	1129.2	135.2	98.9
Trial-2_0494	80	41935	0.5	13.7197	3.6	1.7307	4.6	0.1722	2.8	0.62	1024.3	26.8	1020.0	29.3	1010.8	72.4	1010.8	72.4	101.3
Trial-2_0495	107	47095	2.9	18.2872	4.5	0.6389	5.6	0.0847	3.4	0.60	524.4	17.2	501.7	22.3	399.3	100.8	524.4	17.2	131.3
Trial-2_0496	110	11642	1.3	21.5184	6.8	0.2571	8.5	0.0401	5.0	0.59	253.6	12.5	232.3	17.6	22.2	164.4	253.6	12.5	NA
Trial-2_0497	66	28448	2.1	12.5012	3.6	2.1984	4.6	0.1993	2.9	0.62	1171.7	30.7	1180.5	32.1	1196.8	70.8	1196.8	70.8	97.9
Trial-2_0498	231	143319	3.5	13.3248	2.0	1.8326	2.6	0.1771	1.6	0.62	1051.1	15.7	1057.2	17.1	1069.8	40.8	1069.8	40.8	98.3
Trial-2_0499	107	267574	2.2	13.1970	2.9	1.8729	3.7	0.1793	2.3	0.62	1063.0	22.7	1071.6	24.7	1089.2	58.4	1089.2	58.4	97.6
Trial-2_0500	68	11048	2.6	20.4607	7.1	0.2962	8.9	0.0439	5.3	0.60	277.3	14.4	263.4	20.7	141.8	167.9	277.3	14.4	NA
Trial-2_0501	42	13185	1.6	19.0661	7.2	0.6142	9.0	0.0849	5.4	0.60	525.5	27.4	486.2	34.8	305.0	163.8	525.5	27.4	172.3
Trial-2_0502	59	142353	1.3	13.1627	3.9	1.8960	5.0	0.1810	3.1	0.62	1072.4	30.7	1079.7	33.2	1094.4	78.3	1094.4	78.3	98.0
Trial-2_0503	287	44858	0.9	19.9383	4.2	0.2770	5.3	0.0401	3.1	0.60	253.2	7.8	248.3	11.6	202.2	98.1	253.2	7.8	NA
Trial-2_0504	189	139513	1.5	8.6547	1.7	4.9720	2.2	0.3121	1.4	0.64	1751.0	21.3	1814.6	18.2	1888.4	29.7	1888.4	29.7	92.7
Trial-2_0505	42	67324	1.0	12.0390	4.2	2.5419	5.4	0.2219	3.4	0.63	1292.2	39.9	1284.1	39.5	1270.7	82.2	1270.7	82.2	101.7
Trial-2_0506	55	63720	1.4	13.2354	3.9	1.9315	4.9	0.1854	3.1	0.62	1096.5	31.1	1092.1	33.0	1083.3	77.4	1083.3	77.4	101.2
Trial-2_0507	43	131223	1.8	12.5120	4.4	2.2255	5.6	0.2020	3.5	0.62	1185.8	37.9	1189.1	39.3	1195.1	86.4	1195.1	86.4	99.2
Trial-2_0508	31	74508	2.4	13.3382	5.5	1.8269	7.0	0.1767	4.4	0.62	1049.1	42.3	1055.2	46.2	1067.8	110.9	1067.8	110.9	98.2
Trial-2_0509	411	404703	29.0	13.2659	1.6	1.7109	2.0	0.1646	1.3	0.62	982.3	11.6	1012.6	13.1	1078.7	32.1	1078.7	32.1	91.1
Trial-2_0510	927	101049	1.5	20.1099	2.5	0.2122	3.1	0.0309	1.9	0.60	196.5	3.6	195.4	5.5	182.2	58.3	196.5	3.6	NA
Trial-2_0511	155	19032	0.5	21.7911	6.8	0.1704	8.5	0.0269	5.0	0.59	171.3	8.5	159.7	12.5	-8.1	164.9	171.3	8.5	NA
Trial-2_0512	164	39052	0.5	19.9262	4.6	0.3653	5.8	0.0528	3.5	0.60	331.7	11.2	316.2	15.8	203.6	107.9	331.7	11.2	NA
Trial-2_0513	59	30193	3.9	13.2145	4.0	1.9121	5.1	0.1833	3.2	0.62	1084.7	31.8	1085.3	34.2	1086.5	80.4	1086.5	80.4	99.8
Trial-2_0514	293	274632	0.8	9.4754	1.4	4.0264	1.9	0.2767	1.2	0.64	1574.7	16.6	1639.5	15.1	1723.7	26.1	1723.7	26.1	91.4
Trial-2_0515	83	146841	1.6	8.7461	2.3	5.2529	3.0	0.3332	2.0	0.65	1853.9	31.6	1861.2	25.9	1869.4	41.9	1869.4	41.9	99.2
Trial-2_0516	25	59683	2.4	4.5128	3.2	17.7509	4.2	0.5810	2.8	0.67	2952.8	67.0	2976.3	40.8	2992.3	50.9	2992.3	50.9	98.7
Trial-2_0517	23	23198	1.4	15.8754	8.1	0.9808	10.2	0.1129	6.2	0.61	689.7	40.7	694.0	51.4	707.9	172.3	689.7	40.7	97.4
Trial-2_0518	122	201294	1.8	9.8174	2.1	3.9517	2.8	0.2814	1.8	0.64	1598.3	25.2	1624.3	22.6	1658.3	39.7	1658.3	39.7	96.4
Trial-2_0519	190	316642	1.6	10.9064	1.9	3.0260	2.4	0.2394	1.6	0.63	1383.4	19.3	1414.2	18.7	1460.9	36.0	1460.9	36.0	94.7
Trial-2_0520	58	42756	1.7	8.3743	2.7	5.9034	3.5	0.3585	2.3	0.65	1975.3	38.9	1961.7	30.7	1947.5	48.2	1947.5	48.2	101.4
Trial-2_0521	245	28411	1.5	18.6475	3.3	0.4995	4.2	0.0676	2.5	0.60	421.4	10.2	411.3	14.1	355.4	75.1	421.4	10.2	118.6
Trial-2_0522	52	24764	1.5	13.1522	4.3	1.9542	5.4	0.1864	3.4	0.62	1101.9	34.3	1099.9	36.6	1096.0	85.3	1096.0	85.3	100.5
Trial-2_0523	18	44279	1.9	13.0310	7.3	1.9755	9.3	0.1867	5.8	0.62	1103.5	58.6	1107.2	62.8	1114.4	145.7	1114.4	145.7	99.0
Trial-2_0524	159	338298	2.2	4.3341	1.2	17.9775	1.6	0.5651	1.1	0.67	2887.7	25.4	2988.5	15.6	3057.1	19.3	3057.1	19.3	94.5
Trial-2_0525	43	101582	1.8	13.2599	4.7	1.8882	6.0	0.1816	3.7	0.62	1075.7	36.8	1077.0	39.7	1079.6	93.9	1079.6	93.9	99.6
Trial-2_0526	97	72888	0.8	10.7321	2.5	3.3533	3.3	0.2610	2.1	0.63	1495.0	27.5	1493.6	25.4	1491.5	47.6	1491.5	47.6	100.2
Trial-2_0527	43	39901	1.3	11.1262	4.0	3.1111	5.1	0.2510	3.2	0.63	1443.9	41.9	1435.4	39.4	1422.9	75.9	1422.9	75.9	101.5
Trial-2_0528	83	42112	1.7	13.5771	3.5	1.7157	4.4	0.1689	2.7	0.62	1006.3	25.6	1014.4	28.4	1032.0	70.1	1032.0	70.1	97.5
Trial-2_0529	152	59702	2.4	12.8660	2.4	2.0692	3.1	0.1931	1.9	0.63	1138.1	20.3	1138.7	21.3	1139.9	48.3	1139.9	48.3	99.8
Trial-2_0530	79	130125	0.9	8.3318	2.4	5.7298	3.1	0.3462	2.0	0.65	1916.6	33.5	1935.9	27.0	1956.5	42.6	1956.5	42.6	98.0
Trial-2_0531	22	55042	1.1	13.2440	6.5	1.9050	8.2	0.1830	5.1	0.62	1083.3	51.1	1082.9	55.0	1082.0	129.7	1082.0	129.7	100.1
Trial-2_0532	117	38973	1.9	11.9055	2.6	2.4041	3.3	0.2076	2.1	0.63	1215.9	23.1	1243.8	23.8	1292.4	50.1	1292.4	50.1	94.1
Trial-2_0533	136	391429	1.2	13.4924	2.7	1.8135	3.4	0.1775	2.1	0.62	1053.1	20.5	1050.3	22.2	1044.6	53.6	1044.6	53.6	100.8
Trial-2_0534	783	97535	0.7	12.8297	1.0	1.8862	1.3	0.1755	0.9	0.66	1042.4	8.3	1076.2	8.6	1145.5	19.3	1145.5	19.3	91.0
Trial-2_0535	53	50938	1.6	13.1604	4.3	1.8239	5.5	0.1741	3.4	0.62	1034.6	32.4	1054.1	35.9	1094.7	85.9	1094.7	85.9	94.5
Trial-2_0536	75	34562	1.0	11.1790	3.0	3.1049	3.9	0.2517	2.5	0.63	1447.4	31.8	1433.9	29.8	1413.9	57.4	1413.9	57.4	102.4
Trial-2_0537	66	144165	0.9	9.5204	2.7	4.5248	3.6	0.3124	2.3	0.64	1752.7	35.1	1735.5	29.6	1714.9	50.3	1714.9	50.3	102.2
Trial-2_0538	144	136457	2.4	13.0751	2.5	2.0593	3.2	0.1953	2.0	0.62	1149.9	20.8	1135.4	21.6	1107.7	49.4	1107.7	49.4	103.8
Trial-2_0539	21	20034	1.3	11.1287	2.5	3.1003	5.1	0.2502	4.5	0.87	1439.7	58.0	1432.8	39.5	1422.5	47.9	1422.5	47.9	101.2
Trial-2_0540	444	534057	1.6	12.3259	1.4	2.0571	1.8	0.1839	1.1	0.63	1088.2	11.5	1134.7	12.5	1224.6	28.0	1224.6	28.0	88.9
Trial-2_0541	52	47512	1.6	12.4831	4.0	2.2045	5.2	0.1996	3.2	0.62	1173.1	34.6	1182.5	36.1	1199.6	79.7	1199.6	79.7	97.8
Trial-2_0542	95	61610	1.1	11.1766	2.5	3.1598	3.3	0.2561	2.1	0.63	1470.0	27.4	1447.4	25.4	1414.3	48.7	1414.3	48.7	103.9
Trial-2_0543	17	18259	1.5	12.4584	7.2	2.2951	9.3	0.2074	5.8	0.62	1214.8	64.0	1210.8	65.6	1203.5	142.8	1203.5	142.8	100.9
Trial-2_0544	37	23604	1.8	15.5791	6.1	1.1013	7.7	0.1244	4.8	0.61	756.0	33.9	754.0	41.2	747.8	129.2	756.0	33.9	101.1
Trial-2_0545	23	32266	0.9	7.9442	4.1	6.7849	5.4	0.3909	3.5	0.65	2127.1	63.9	2083.8	48.1	2041.2	73.1	2041.2	73.1	104.2
Trial-2_0546	432	90014	3.0	20.1207	3.4	0.2845	4.2	0.0415	2.5	0.60	262.3	6.5	254.2	9.5	181.0	79.3	262.3	6.5	NA
Trial-2_0547	141	72331	1.8	13.1280	2.5	2.0320	3.2	0.1935	2.0	0.62	1140.2	21.0	1126.3	21.9	1099.6	50.4	1099.6	50.4	103.7
Trial-2_0548	142	820941	1.5	7.9827	1.8	5.8054	2.4	0.3361	1.5	0.65	1867.9	24.8	1947.2	20.5	2032.6	31.8	2032.6	31.8	91.9
Trial-2_0549	78	58112	1.6	13.0567	3.3	2.0430	4.2	0.1935	2.7	0.62	1140.1	27.7	1130.0	29.0	1110.5	66.3	1110.5	66.3	102.7
Trial-2_0550	665	156859	2.1	17.7590	2.0	0.5298	2.5	0.0682	1.5	0.61	425.5	6.3	431.6	8.9	464.6	44.7	425.5	6.3	91.6
Trial-2_0551	70	219503	1.4	9.2191	2.7	4.8412	3.5	0.3237	2.3	0.64	1807.8	35.5	1792.1	29.6	1773.9	49.2	1773.9	49.2	101.9
Trial-2_0552	103	337803	0.9	11.1174	2.6	3.0314	3.4	0.2444	2.2	0.63	1409.7	27.3	1415.6	26.0	1424.4	50.4	1424.4	50.4	99.0
Trial-2_0553	89	42346	0.7	12.9935	3.2	1.9884	4.1	0.1874	2.6	0.62	1107.2	26.1	1111.6	27.8	1120.2	64.2	1120.2	64.2	98.8
Trial-2_0554	27	69940	1.0	13.4226	6.1	1.7947	7.7	0.1747	4.8	0.62	1038.0	46.0	1043.5	50.5	1055.1	122.4	1055.1	122.4	98.4
Trial-2_0555	590	126153	8.4	17.7478	2.2	0.5175	2.7	0.0666	1.7	0.60	415.7	6.6	423.4	9.5	466.0	48.1	415.7	6.6	89.2
Trial-2_0556	495	55583	1.2	19.4410	3.2	0.2869	4.0	0.0405	2.4	0.60	255.6	6.1	256.1	9.2	260.5	74.5	255.6	6.1	NA
Trial-2_0557	76	146764	1.4	2.8813	1.6	35.8641	2.1	0.7495	1.5	0.68	3605.5	40.4	3662.7	21.1	3694.2	23.8	3694.2	23.8	97.6
Trial-2_0558	227	545762	2.8	13.4072	2.1	1.8077	2.6	0.1758	1.6	0.62	1043.9	15.8	1048.2	17.2	1057.4	41.5	1057.4	41.5	98.7
Trial-2_0559	216	299223	2.0	11.0162	1.8	2.9220	2.3	0.2335	1.5	0.63	1352.6	18.1	1387.6	17.7	1441.9	34.5	1441.9	34.5	93.8
Trial-2_0560	111	6810	0.5	22.6554	7.9	0.1776	9.7	0.0292	5.7	0.59	185.4	10.5	166.0	14.9	-102.9	193.9	185.4	10.5	NA
Trial-2_0561	53	73468	1.0	13.7369	4.3	1.6742	5.5	0.1668	3.4	0.62	994.5	31.4	998.8	35.0	1008.3	87.5	1008.3	87.5	98.6
Trial-2_0562	110	19049	0.7	21.8739	7.0	0.2526	8.7	0.0401	5.1	0.59	253.3	12.8	228.7	17.8	-17.3	169.3	253.3	12.8	NA
Trial-2_0563	247	50205	0.9	20.7468	4.8	0.2797	5.9	0.0421	3.5	0.60	265.7	9.2	250.4	13.1	109.1	112.2	265.7	9.2	NA
Trial-2_0564	112	99922	7.9	13.8239	3.0	1.7423	3.9	0.1747	2.4	0.62	1037.9	23.1	1024.3	25.1	995.5	61.9	995.5	61.9	104.3
Trial-2_0565	149	455637	4.1	9.3895	1.9	4.3763	2.5	0.2980	1.6	0.64	1681.5	23.9	1707.9	20.8	1740.4	35.4	1740.4	35.4	96.6
Trial-2_0566	127	188493	1.8	9.8407	2.1	4.0352	2.7	0.2880	1.7	0.64	1631.5	24.9	1641.3	22.0	1653.9	38.4	1653.9	38.4	98.6
Trial-2_0567	65	27973	1.7	13.2173	3.9	1.8964	5.0	0.1818	3.1	0.62	1076.7	31.1	1079.8	33.5	1086.0	79.1	1086.0	79.1	99.1
Trial-2_0568	94	125365	1.2	13.3029	3.4	1.6733	4.3	0.1614	2.7	0.62	964.8	23.9	998.5	27.3	1073.1	67.9	1073.1	67.9	89.9
Trial-2_0569	117	44369	1.1	18.7931	5.0	0.4731	6.2	0.0645	3.8	0.60	402.9	14.7	393.4	20.3	337.8	112.9	402.9	14.7	119.3
Trial-2_0570	339	108942	1.0	18.3088	2.8	0.5221	3.5	0.0693	2.1	0.60	432.1	8.9	426.5	12.3	396.6	63.0	432.1	8.9	108.9
Trial-2_0571	235	82676	0.8	18.3383	3.5	0.5250	4.4	0.0698	2.6	0.60	435.1	11.1	428.5	15.2	393.0	77.9	435.1	11.1	110.7
Trial-2_0572	43	15695	1.2	19.6591	8.0	0.4924	10.1	0.0702	6.0	0.60	437.4	25.6	406.5	33.7	234.8	185.6	437.4	25.6	186.3
Trial-2_0573	96	166928	1.9	13.1982	3.1	1.9227	3.9	0.1840	2.4	0.62	1089.0	24.3	1089.0	26.1	1088.9	61.4	1088.9	61.4	100.0
Trial-2_0574	85	286273	0.6	5.2797	1.8	13.2754	2.4	0.5083	1.6	0.66	2649.5	35.1	2699.4	23.0	2737.0	29.9	2737.0	29.9	96.8
Trial-2_0575	507	291941	2.6	13.8005	1.5	1.4363	1.9	0.1438	1.2	0.62	865.9	9.8	904.2	11.7	998.9	31.1	998.9	31.1	86.7
Trial-2_0576	45	32306	1.3	11.1656	3.7	3.0378	4.8	0.2460	3.1	0.63	1417.8	38.9	1417.2	36.9	1416.2	71.7	1416.2	71.7	100.1
Trial-2_0577	27	37903	0.9	5.5724	3.2	12.6944	4.2	0.5130	2.8	0.66	2669.6	61.4	2657.2	39.9	2647.8	52.7	2647.8	52.7	100.8
Trial-2_0578	95	91049	2.6	12.2595	2.9	2.2861	3.8	0.2033	2.4	0.63	1192.8	25.8	1208.0	26.7	1235.2	57.8	1235.2	57.8	96.6
Trial-2_0579	225	193432	1.9	14.1467	2.2	1.4620	2.8	0.1500	1.8	0.62	901.0	14.8	914.8	17.2	948.4	45.8	948.4	45.8	95.0
Trial-2_0580	43	179537	0.9	11.0606	3.9	3.0680	5.0	0.2461	3.2	0.63	1418.4	40.3	1424.7	38.4	1434.2	74.1	1434.2	74.1	98.9
Trial-2_0581	22	26637	1.5	13.1152	6.5	1.9107	8.3	0.1817	5.1	0.62	1076.5	51.0	1084.8	55.2	1101.6	129.9	1101.6	129.9	97.7
Trial-2_0582	187	183133	1.0	18.1848	3.7	0.5322	4.6	0.0702	2.8	0.60	437.3	11.7	433.3	16.2	411.8	81.9	437.3	11.7	106.2
Trial-2_0583	53	30571	2.2	12.8691	4.0	2.0757	5.1	0.1937	3.2	0.62	1141.6	33.1	1140.8	34.7	1139.3	78.7	1139.3	78.7	100.2
Trial-2_0584	79	71155	1.3	11.2062	2.9	3.0452	3.7	0.2475	2.4	0.63	1425.6	30.1	1419.0	28.5	1409.2	55.2	1409.2	55.2	101.2
Trial-2_0585	65	58646	2.3	12.8553	3.7	2.1227	4.7	0.1979	2.9	0.62	1164.1	31.2	1156.2	32.4	1141.5	72.9	1141.5	72.9	102.0
Trial-2_0586	23	25800	0.3	5.5297	3.5	12.9602	4.6	0.5198	3.1	0.66	2698.2	67.6	2676.8	43.6	2660.6	57.3	2660.6	57.3	101.4
Trial-2_0587	425	563633	2.7	12.8017	1.5	1.8595	1.9	0.1726	1.2	0.62	1026.7	11.5	1066.8	12.9	1149.8	30.2	1149.8	30.2	89.3
Trial-2_0588	99	188258	2.7	9.7863	2.3	4.0633	3.1	0.2884	1.9	0.64	1633.5	28.1	1647.0	24.9	1664.1	43.5	1664.1	43.5	98.2
Trial-2_0589	260	75970	1.9	18.2252	3.1	0.5647	3.9	0.0746	2.4	0.60	464.1	10.5	454.6	14.3	406.9	69.5	464.1	10.5	114.1
Trial-2_0590	44	86555	1.4	8.7943	3.3	5.2203	4.3	0.3330	2.7	0.64	1852.7	44.3	1855.9	36.5	1859.5	59.2	1859.5	59.2	99.6
Trial-2_0591	154	265476	1.5	9.9060	1.9	3.8542	2.5	0.2769	1.6	0.64	1575.8	22.0	1604.1	19.9	1641.6	35.3	1641.6	35.3	96.0
Trial-2_0592	125	15551	1.5	20.5711	5.9	0.2922	7.3	0.0436	4.4	0.60	275.1	11.7	260.3	16.8	129.2	138.0	275.1	11.7	NA
Trial-2_0593	93	90983	2.6	13.8170	3.3	1.6771	4.2	0.1681	2.6	0.62	1001.4	24.3	999.9	26.9	996.5	67.3	996.5	67.3	100.5
Trial-2_0594	37	293054	1.9	9.8355	3.9	4.0027	5.1	0.2855	3.3	0.64	1619.1	46.7	1634.7	41.6	1654.9	73.1	1654.9	73.1	97.8
Trial-2_0595	43	46985	1.9	13.6144	4.7	1.7488	6.0	0.1727	3.7	0.62	1026.8	35.6	1026.7	39.0	1026.4	95.8	1026.4	95.8	100.0
Trial-2_0596	235	269514	2.9	13.6393	2.1	1.7099	2.7	0.1691	1.6	0.62	1007.4	15.4	1012.3	17.0	1022.7	42.0	1022.7	42.0	98.5
Trial-2_0597	136	45862	1.4	12.7467	2.5	2.0779	3.2	0.1921	2.0	0.63	1132.7	21.1	1141.5	22.2	1158.3	50.2	1158.3	50.2	97.8
Trial-2_0598	65	55888	0.3	8.9580	2.7	5.0495	3.6	0.3281	2.3	0.64	1829.0	36.4	1827.7	30.1	1826.1	49.4	1826.1	49.4	100.2
Trial-2_0599	122	484681	0.9	6.4165	1.7	9.2626	2.2	0.4311	1.5	0.66	2310.5	28.5	2364.3	20.5	2411.1	28.6	2411.1	28.6	95.8
Trial-2_0600	24	37274	0.8	13.5278	6.6	1.7321	8.4	0.1699	5.2	0.62	1011.8	48.4	1020.5	53.9	1039.3	132.9	1039.3	132.9	97.4
Trial-2_0601	38	25824	1.4	13.2076	4.9	1.8969	6.3	0.1817	3.9	0.62	1076.3	38.8	1080.0	42.0	1087.5	99.2	1087.5	99.2	99.0
Trial-2_0602	435	347562	1.8	13.2917	1.6	1.7366	2.0	0.1674	1.2	0.62	997.8	11.5	1022.2	12.9	1074.8	31.5	1074.8	31.5	92.8
Trial-2_0603	832	143047	1.7	21.3939	3.9	0.0986	4.8	0.0153	2.9	0.59	97.9	2.8	95.5	4.4	36.1	93.4	97.9	2.8	NA
Trial-2_0604	294	651647	2.0	13.1411	1.9	1.7818	2.4	0.1698	1.5	0.62	1011.1	13.8	1038.9	15.4	1097.6	37.2	1097.6	37.2	92.1
Trial-2_0605	112	49103	2.0	13.9245	3.1	1.6228	3.9	0.1639	2.4	0.62	978.4	22.0	979.1	24.6	980.7	62.5	980.7	62.5	99.8
Trial-2_0606	198	230511	2.5	12.5548	2.1	2.1062	2.7	0.1918	1.7	0.63	1131.0	17.7	1150.8	18.7	1188.4	41.9	1188.4	41.9	95.2
Trial-2_0607	49	21609	3.6	13.7193	4.8	1.5764	6.1	0.1569	3.8	0.62	939.3	33.3	961.0	38.2	1010.9	97.6	1010.9	97.6	92.9
Trial-2_0608	127	225811	2.7	13.7605	2.9	1.6857	3.6	0.1682	2.3	0.62	1002.4	21.0	1003.1	23.2	1004.8	58.0	1004.8	58.0	99.8
Trial-2_0609	91	603679	1.8	11.3202	2.7	2.9481	3.5	0.2420	2.2	0.63	1397.3	28.0	1394.4	26.8	1389.8	52.5	1389.8	52.5	100.5
Trial-2_0610	60	81240	2.0	13.9968	4.3	1.5679	5.4	0.1592	3.4	0.62	952.1	29.7	957.6	33.7	970.2	87.0	970.2	87.0	98.1
Trial-2_0611	17	14257	1.7	13.1438	7.6	1.9427	9.7	0.1852	6.1	0.62	1095.3	61.0	1095.9	65.4	1097.2	152.9	1097.2	152.9	99.8
Trial-2_0612	86	113592	1.7	13.4734	3.4	1.7184	4.3	0.1679	2.7	0.62	1000.6	24.9	1015.4	27.9	1047.5	68.7	1047.5	68.7	95.5

Trial-2_0613	13	6562	0.8	13.4408	8.8	1.8218	11.2	0.1776	7.0	0.62	1053.8	67.7	1053.3	73.7	1052.4	177.6	1052.4	177.6	100.1
Trial-2_0614	33	67025	0.5	11.5541	4.5	2.8876	5.7	0.2420	3.6	0.63	1397.0	45.6	1378.7	43.4	1350.5	86.1	1350.5	86.1	103.4
Trial-2_0615	139	72315	1.4	12.9315	2.6	1.9960	3.3	0.1872	2.0	0.62	1106.2	20.7	1114.2	22.1	1129.7	51.0	1129.7	51.0	97.9
Trial-2_0616	109	92791	2.3	19.2051	5.7	0.3870	7.1	0.0539	4.3	0.60	338.5	14.1	332.2	20.2	288.5	130.4	338.5	14.1	NA
Trial-2_0617	226	77191	0.8	20.6660	4.8	0.2832	6.0	0.0424	3.6	0.59	268.0	9.4	253.2	13.4	118.3	113.6	268.0	9.4	NA
Trial-2_0618	192	203882	0.7	18.3740	3.7	0.5292	4.6	0.0705	2.8	0.60	439.3	11.8	431.3	16.2	388.6	82.6	439.3	11.8	113.0
Trial-2_0619	11	5135	1.3	13.4380	9.4	1.8157	12.0	0.1770	7.4	0.62	1050.4	72.2	1051.2	78.7	1052.8	190.0	1052.8	190.0	99.8
Trial-2_0620	109	164643	1.8	13.8923	3.1	1.6645	4.0	0.1677	2.5	0.62	999.5	22.8	995.1	25.1	985.4	63.3	985.4	63.3	101.4
Trial-2_0621	409	100165	3.1	21.8717	5.7	0.0968	7.1	0.0154	4.2	0.59	98.3	4.1	93.8	6.4	-17.0	138.8	98.3	4.1	NA
Trial-2_0622	150	51964	1.1	17.2010	3.5	0.7816	4.4	0.0975	2.7	0.61	599.8	15.3	586.4	19.6	534.9	76.3	599.8	15.3	112.1
Trial-2_0623	331	61228	1.9	19.4326	3.2	0.3836	4.0	0.0541	2.4	0.60	339.5	7.9	329.7	11.2	261.5	73.4	339.5	7.9	NA
Trial-2_0624	306	542858	9.9	12.3139	1.7	2.0917	2.2	0.1868	1.4	0.63	1104.1	14.1	1146.1	15.2	1226.5	33.9	1226.5	33.9	90.0
Trial-2_0625	89	58260	3.5	13.1522	3.4	1.7262	4.4	0.1647	2.7	0.62	982.6	24.7	1018.3	28.1	1095.9	68.7	1095.9	68.7	89.7
Trial-2_0626	46	7033	1.0	22.8743	10.3	0.2366	12.8	0.0393	7.5	0.59	248.2	18.3	215.6	24.8	-126.6	255.1	248.2	18.3	NA
Trial-2_0627	18	4888	1.8	12.7006	7.4	2.0075	9.4	0.1849	5.8	0.62	1093.8	58.8	1118.1	64.0	1165.5	146.7	1165.5	146.7	93.8
Trial-2_0628	29	33937	2.0	9.3982	4.7	3.8094	6.1	0.2597	3.8	0.63	1488.1	51.0	1594.7	48.9	1738.7	86.4	1738.7	86.4	85.6
Trial-2_0629	21	9814	1.2	13.4747	6.8	1.7407	8.7	0.1701	5.4	0.62	1012.7	50.3	1023.7	56.1	1047.3	137.8	1047.3	137.8	96.7
Trial-2_0630	6	107878	2.0	13.4286	2.8	1.8007	3.0	0.1754	1.0	0.35	1041.7	10.0	1045.7	19.4	1054.2	56.1	1054.2	56.1	98.8
Trial-2_0631	99	61641	2.0	13.6687	3.2	1.7187	4.1	0.1704	2.5	0.62	1014.2	23.9	1015.5	26.4	1018.4	65.3	1018.4	65.3	99.6
Trial-2_0632	52	87554	2.7	9.1122	3.1	4.7365	4.1	0.3130	2.6	0.64	1755.6	40.0	1773.7	34.1	1795.1	56.8	1795.1	56.8	97.8
Trial-2_0633	41	82513	3.1	12.4938	4.5	2.0951	5.8	0.1898	3.6	0.62	1120.5	37.1	1147.2	39.8	1198.0	89.4	1198.0	89.4	93.5
Trial-2_0634	387	128144	2.2	17.4274	2.4	0.6505	3.0	0.0822	1.8	0.61	509.4	8.9	508.8	12.0	506.2	52.6	509.4	8.9	100.6
Trial-2_0635	13	75528	1.4	13.0569	8.4	1.9268	10.8	0.1825	6.7	0.62	1080.4	66.5	1090.4	72.0	1110.5	168.6	1110.5	168.6	97.3
Trial-2_0636	111	155455	2.1	9.4373	2.2	4.3983	2.8	0.3010	1.8	0.64	1696.5	27.0	1712.0	23.4	1731.1	39.8	1731.1	39.8	98.0
Trial-2_0637	97	17627	1.9	21.0092	7.4	0.2834	9.2	0.0432	5.5	0.59	272.6	14.6	253.4	20.6	79.3	175.3	272.6	14.6	NA
Trial-2_0638	76	112459	1.9	13.6363	3.6	1.7059	4.6	0.1687	2.8	0.62	1005.0	26.5	1010.7	29.3	1023.2	72.7	1023.2	72.7	98.2
Trial-2_0639	170	54189	1.0	13.3044	2.4	1.8576	3.0	0.1792	1.9	0.62	1062.8	18.5	1066.1	19.9	1072.9	47.5	1072.9	47.5	99.1
Trial-2_0640	221	437365	3.1	7.3960	1.4	6.7332	1.8	0.3612	1.2	0.65	1987.7	20.3	2077.0	16.1	2166.7	24.1	2166.7	24.1	91.7
Trial-2_0641	49	15722	0.8	21.7013	10.0	0.2456	12.5	0.0387	7.4	0.59	244.5	17.8	223.0	24.9	1.8	241.7	244.5	17.8	NA
Trial-2_0642	64	181852	0.7	5.4475	2.1	12.9614	2.8	0.5121	1.8	0.66	2665.6	40.4	2676.9	26.3	2685.4	34.5	2685.4	34.5	99.3
Trial-2_0643	37	35019	1.7	12.9217	4.9	2.1296	6.3	0.1996	3.9	0.63	1173.1	42.1	1158.5	43.4	1131.3	97.6	1131.3	97.6	103.7
Trial-2_0644	113	72881	1.6	12.3371	2.7	2.3291	3.5	0.2084	2.2	0.63	1220.3	24.3	1221.2	24.8	1222.8	53.4	1222.8	53.4	99.8
Trial-2_0645	222	713182	3.4	9.2747	1.6	4.2930	2.1	0.2888	1.3	0.64	1635.4	19.0	1692.0	16.9	1762.9	28.8	1762.9	28.8	92.8
Trial-2_0646	111	193815	2.1	9.6591	2.2	4.1290	2.8	0.2893	1.8	0.64	1637.8	26.1	1660.1	23.0	1688.3	39.9	1688.3	39.9	97.0
Trial-2_0647	100	192543	1.6	9.2482	2.3	4.6571	3.0	0.3124	1.9	0.64	1752.4	29.6	1759.6	25.1	1768.1	42.1	1768.1	42.1	99.1
Trial-2_0648	240	493804	3.3	9.1280	1.5	4.4221	2.0	0.2928	1.3	0.64	1655.3	18.5	1716.5	16.3	1792.0	27.5	1792.0	27.5	92.4
Trial-2_0649	46	5373	1.7	25.6385	13.1	0.1411	16.0	0.0262	9.3	0.58	166.9	15.3	134.0	20.1	-416.5	342.8	166.9	15.3	NA
Trial-2_0650	153	551040	2.3	5.1557	1.4	12.8811	1.9	0.4817	1.2	0.66	2534.5	26.0	2671.0	17.6	2776.0	22.9	2776.0	22.9	91.3
Trial-2_0651	64	101308	1.0	11.2898	3.2	2.9917	4.2	0.2450	2.6	0.63	1412.5	33.3	1405.5	31.7	1395.0	61.8	1395.0	61.8	101.3
Trial-2_0652	147	226079	2.2	13.2127	2.5	1.8662	3.3	0.1788	2.0	0.62	1060.6	19.8	1069.2	21.5	1086.7	51.0	1086.7	51.0	97.6
Trial-2_0653	196	70005	1.3	20.2197	4.8	0.2873	6.0	0.0421	3.6	0.60	266.1	9.3	256.5	13.5	169.5	111.6	266.1	9.3	NA
Trial-2_0654	153	858368	1.8	10.9550	2.1	3.0296	2.8	0.2407	1.7	0.63	1390.4	21.9	1415.1	21.1	1452.5	40.7	1452.5	40.7	95.7
Trial-2_0655	83	128919	1.2	11.0943	2.9	3.0844	3.7	0.2482	2.3	0.63	1429.1	29.9	1428.8	28.3	1428.4	54.6	1428.4	54.6	100.0
Trial-2_0656	1120	595132	1.1	6.5690	2.3	0.6055	3.0	0.0288	2.0	0.65	183.3	3.5	480.7	11.6	2371.1	39.5	183.3	3.5	NA
Trial-2_0657	372	13942	3.1	11.0734	1.4	2.9032	1.8	0.2332	1.1	0.63	1351.0	13.6	1382.7	13.3	1432.0	26.0	1432.0	26.0	94.3
Trial-2_0658	313	75115	1.7	16.1774	2.3	0.8423	3.0	0.0988	1.8	0.61	607.5	10.5	620.4	13.7	667.7	50.1	607.5	10.5	91.0
Trial-2_0659	149	211095	1.8	11.0524	2.1	3.0408	2.8	0.2437	1.7	0.63	1406.2	22.0	1417.9	21.0	1435.6	40.6	1435.6	40.6	97.9
Trial-2_0660	12	8387	1.2	12.6001	8.1	2.1518	10.4	0.1966	6.5	0.62	1157.2	68.6	1165.6	72.0	1181.3	160.4	1181.3	160.4	98.0
Trial-2_0661	42	17492	0.7	13.6101	5.0	1.6467	6.4	0.1625	4.0	0.62	970.9	35.6	988.3	40.4	1027.1	101.7	1027.1	101.7	94.5
Trial-2_0662	382	339533	3.9	9.9960	1.4	3.1116	1.8	0.2256	1.1	0.64	1311.3	13.6	1435.6	13.9	1624.8	26.0	1624.8	26.0	80.7
Trial-2_0663	226	476690	0.9	10.8341	1.7	3.0545	2.3	0.2400	1.4	0.63	1386.7	17.8	1421.4	17.2	1473.6	33.1	1473.6	33.1	94.1
Trial-2_0664	71	61137	1.6	11.1229	3.0	3.0854	3.9	0.2489	2.5	0.63	1432.8	31.7	1429.1	29.9	1423.5	57.6	1423.5	57.6	100.7
Trial-2_0665	31	18001	2.9	13.2004	5.5	1.9260	7.0	0.1844	4.3	0.62	1090.9	43.7	1090.1	46.9	1088.6	110.5	1088.6	110.5	100.2
Trial-2_0666	112	562073	1.1	13.4313	2.9	1.8302	3.7	0.1783	2.3	0.62	1057.6	22.7	1056.4	24.5	1053.8	58.9	1053.8	58.9	100.4
Trial-2_0667	147	60105	3.2	13.5792	2.6	1.7699	3.3	0.1743	2.1	0.62	1035.8	19.7	1034.5	21.5	1031.7	52.5	1031.7	52.5	100.4
Trial-2_0668	88	37205	1.8	17.4425	4.7	0.7535	5.9	0.0953	3.6	0.61	586.9	20.1	570.2	25.7	504.3	103.1	586.9	20.1	116.4
Trial-2_0669	26	93451	1.9	13.6044	8.0	1.3990	10.1	0.1380	6.3	0.62	833.6	48.9	888.5	60.0	1027.9	161.0	1027.9	161.0	81.1
Trial-2_0670	149	55272	2.3	13.3798	2.5	1.9265	3.2	0.1869	2.0	0.62	1104.8	20.2	1090.3	21.3	1061.5	50.1	1061.5	50.1	104.1
Trial-2_0671	74	6577	0.7	21.8671	8.7	0.2516	10.8	0.0399	6.4	0.59	252.3	15.7	227.9	22.0	-16.5	210.3	252.3	15.7	NA
Trial-2_0672	45	66049	1.0	5.3111	2.5	13.3374	3.3	0.5138	2.2	0.66	2672.6	48.2	2703.8	31.3	2727.2	40.8	2727.2	40.8	98.0
Trial-2_0673	229	64224	2.1	19.0616	3.6	0.4292	4.6	0.0593	2.7	0.60	371.6	9.9	362.6	13.9	305.6	83.1	371.6	9.9	NA
Trial-2_0674	95	119006	2.1	13.1362	3.2	1.9287	4.1	0.1838	2.5	0.62	1087.5	25.3	1091.1	27.1	1098.4	63.5	1098.4	63.5	99.0
Trial-2_0675	486	55264	1.8	20.1140	3.2	0.2780	4.0	0.0406	2.4	0.60	256.3	6.0	249.1	8.9	181.7	75.5	256.3	6.0	NA
Trial-2_0676	336	587436	2.0	9.9829	1.4	3.3882	1.8	0.2453	1.2	0.64	1414.3	14.8	1501.7	14.4	1627.2	26.4	1627.2	26.4	86.9
Trial-2_0677	152	114607	13.0	13.5620	2.6	1.7147	3.3	0.1687	2.1	0.62	1004.7	19.3	1014.0	21.5	1034.2	53.1	1034.2	53.1	97.1
Trial-2_0678	160	72868	1.4	16.9327	3.4	0.7923	4.3	0.0973	2.6	0.61	598.6	14.8	592.5	19.2	569.2	73.8	598.6	14.8	105.2
Trial-2_0679	153	21350	2.0	19.3275	5.6	0.2940	7.0	0.0412	4.2	0.60	260.4	10.6	261.7	16.1	273.9	128.5	260.4	10.6	NA
Trial-2_0680	219	23954	1.4	20.3672	4.9	0.2706	6.0	0.0400	3.6	0.59	252.7	8.9	243.2	13.1	152.6	113.9	252.7	8.9	NA
Trial-2_0681	238	1089406	2.7	9.1978	1.6	4.1739	2.1	0.2784	1.3	0.64	1583.5	18.6	1668.9	17.0	1778.1	29.0	1778.1	29.0	89.1
Trial-2_0682	462	570059	4.9	12.4853	1.4	2.0918	1.8	0.1894	1.1	0.63	1118.2	11.6	1146.1	12.4	1199.3	27.6	1199.3	27.6	93.2
Trial-2_0683	446	302685	5.7	12.4740	1.4	2.1622	1.8	0.1956	1.1	0.63	1151.7	11.8	1169.0	12.4	1201.1	27.3	1201.1	27.3	95.9
Trial-2_0684	128	43247	1.9	13.3698	2.8	1.8239	3.6	0.1769	2.2	0.62	1049.8	21.4	1054.1	23.3	1063.0	56.0	1063.0	56.0	98.8
Trial-2_0685	60	144832	1.3	11.2140	3.5	3.0774	4.5	0.2503	2.8	0.63	1440.0	36.8	1427.1	34.6	1407.9	67.0	1407.9	67.0	102.3
Trial-2_0686	109	54851	0.4	13.6943	3.1	1.7445	3.9	0.1733	2.4	0.62	1030.1	23.2	1025.1	25.3	1014.6	62.4	1014.6	62.4	101.5
Trial-2_0687	835	1597602	12.0	12.3818	1.2	2.1471	1.5	0.1928	0.9	0.63	1136.6	9.6	1164.1	10.2	1215.7	22.6	1215.7	22.6	93.5
Trial-2_0688	59	88190	4.1	11.0693	3.6	3.0315	4.6	0.2434	2.9	0.62	1404.2	36.5	1415.6	35.4	1432.7	69.2	1432.7	69.2	98.0
Trial-2_0689	112	189902	1.4	12.6774	2.9	2.0965	3.7	0.1928	2.3	0.62	1136.3	23.9	1147.7	25.3	1169.1	56.8	1169.1	56.8	97.2
Trial-2_0690	17	35288	1.6	13.7019	7.8	1.7972	9.9	0.1786	6.2	0.62	1059.3	60.1	1044.5	64.7	1013.4	157.6	1013.4	157.6	104.5
Trial-2_0691	127	589812	1.9	10.8556	2.3	3.1866	3.0	0.2509	1.9	0.63	1443.1	24.7	1453.9	23.3	1469.8	44.3	1469.8	44.3	98.2
Trial-2_0692	32	17048	1.0	12.3063	5.1	2.4188	6.5	0.2159	4.1	0.63	1260.1	46.8	1248.2	46.9	1227.7	99.9	1227.7	99.9	102.6
Trial-2_0693	129	121365	0.6	5.0549	1.5	14.0761	2.1	0.5161	1.4	0.67	2682.4	30.0	2754.8	19.5	2808.3	25.1	2808.3	25.1	95.5
Trial-2_0694	35	57886	1.2	13.0895	5.1	2.0336	6.5	0.1931	4.1	0.62	1137.9	42.3	1126.8	44.2	1105.5	101.4	1105.5	101.4	102.9
Trial-2_0695	28	50912	0.9	5.3340	3.2	13.4820	4.3	0.5216	2.8	0.66	2705.8	63.0	2714.0	40.5	2720.1	52.8	2720.1	52.8	99.5
Trial-2_0696	84	20146	2.2	18.0773	5.3	0.6415	6.7	0.0841	4.1	0.60	520.6	20.3	503.2	26.6	425.1	119.2	520.6	20.3	122.5
Trial-2_0697	44	51888	1.3	13.0829	4.6	2.0620	5.8	0.1957	3.6	0.62	1152.0	38.5	1136.3	40.0	1106.5	91.3	1106.5	91.3	104.1
Trial-2_0698	38	79764	2.3	13.6025	5.0	1.8422	6.3	0.1817	3.9	0.62	1076.5	39.1	1060.6	41.7	1028.2	100.4	1028.2	100.4	104.7
Trial-2_0699	71	32017	1.6	13.0882	3.8	1.9058	4.9	0.1809	3.0	0.62	1071.9	29.9	1083.1	32.4	1105.7	76.3	1105.7	76.3	96.9
Trial-2_0700	96	63270	2.5	12.6337	3.1	2.2192	4.0	0.2033	2.5	0.63	1193.2	27.1	1187.1	27.9	1176.0	61.5	1176.0	61.5	101.5
Trial-2_0701	362	189153	2.1	17.0635	2.5	0.6927	3.1	0.0857	1.9	0.61	530.2	9.6	534.4	12.9	552.4	53.6	530.2	9.6	96.0
Trial-2_0702	31	8057	1.1	13.2130	5.7	1.9246	7.3	0.1844	4.5	0.62	1091.2	45.2	1089.7	48.5	1086.7	114.0	1086.7	114.0	100.4
Trial-2_0703	147	103996	4.2	13.8781	2.7	1.6734	3.5	0.1684	2.2	0.62	1003.5	20.1	998.5	22.1	987.5	55.5	987.5	55.5	101.6
Trial-2_0704	84	387776	0.9	5.4422	1.9	12.7668	2.6	0.5039	1.7	0.66	2630.6	36.9	2662.6	24.3	2687.0	31.9	2687.0	31.9	97.9
Trial-2_0705	93	99558	3.8	13.8471	3.4	1.7156	4.4	0.1723	2.7	0.62	1024.7	25.7	1014.4	28.0	992.1	69.7	992.1	69.7	103.3
Trial-2_0706	161	166881	2.6	9.7887	1.9	3.9285	2.5	0.2789	1.6	0.64	1585.8	22.3	1619.6	20.1	1663.7	35.3	1663.7	35.3	95.3
Trial-2_0707	22	45189	2.1	13.0934	6.6	1.9708	8.5	0.1872	5.3	0.62	1106.0	53.5	1105.6	57.1	1104.9	133.0	1104.9	133.0	100.1
Trial-2_0708	100	496505	1.0	5.3482	1.8	13.1544	2.4	0.5102	1.6	0.66	2657.7	34.0	2690.8	22.2	2715.7	29.0	2715.7	29.0	97.9
Trial-2_0709	356	384256	3.6	13.2062	1.7	1.7520	2.2	0.1678	1.4	0.62	1000.0	12.9	1027.9	14.4	1087.7	35.0	1087.7	35.0	91.9
Trial-2_0710	583	154941	4.4	16.7945	1.9	0.7192	2.4	0.0876	1.5	0.61	541.4	7.7	550.2	10.3	587.0	41.9	541.4	7.7	92.2
Trial-2_0711	172	58974	2.2	13.2755	2.4	1.8813	3.1	0.1811	1.9	0.62	1073.2	18.9	1074.5	20.4	1077.3	48.2	1077.3	48.2	99.6
Trial-2_0712	300	83282	0.7	17.9946	3.0	0.5547	3.8	0.0724	2.3	0.60	450.6	9.9	448.1	13.7	435.3	67.1	450.6	9.9	103.5
Trial-2_0713	70	112715	0.7	5.9725	2.2	10.9042	2.9	0.4723	1.9	0.66	2493.8	39.9	2515.0	27.2	2532.2	36.9	2532.2	36.9	98.5
Trial-2_0714	589	1201924	2.0	9.2257	1.0	3.9675	1.4	0.2655	0.9	0.64	1517.8	11.8	1627.6	11.0	1772.6	19.0	1772.6	19.0	85.6
Trial-2_0715	163	219916	0.7	10.6281	2.0	3.3946	2.6	0.2617	1.6	0.64	1498.4	21.9	1503.2	20.2	1509.9	37.6	1509.9	37.6	99.2
Trial-2_0716	60	69979	1.2	11.2988	3.4	3.1045	4.4	0.2544	2.8	0.63	1461.2	36.3	1433.8	33.7	1393.4	65.2	1393.4	65.2	104.9
Trial-2_0717	38	40355	9.1	14.2668	5.7	1.4703	7.2	0.1521	4.5	0.62	912.9	37.9	918.3	43.5	931.1	116.1	931.1	116.1	98.1
Trial-2_0718	122	32410	1.3	18.7573	4.8	0.5229	6.0	0.0711	3.6	0.60	443.0	15.4	427.1	20.9	342.1	108.1	443.0	15.4	129.5
Trial-2_0719	85	56619	1.6	13.6653	3.5	1.7790	4.4	0.1763	2.7	0.62	1046.8	26.5	1037.8	28.7	1018.9	70.0	1018.9	70.0	102.7
Trial-2_0720	196	303498	2.4	9.1146	1.7	4.5102	2.2	0.2982	1.4	0.64	1682.1	21.1	1732.9	18.5	1794.6	31.1	1794.6	31.1	93.7
Trial-2_0721	111	169156	2.1	9.5129	2.3	4.2607	2.9	0.2940	1.9	0.64	1661.3	27.5	1685.8	24.1	1716.4	41.4	1716.4	41.4	96.8
Trial-2_0722	139	76519	1.8	13.8069	2.8	1.6936	3.5	0.1696	2.2	0.62	1009.9	20.4	1006.1	22.4	998.0	55.9	998.0	55.9	101.2
Trial-2_0723	1323	27508	3.5	15.9171	1.1	1.0414	1.4	0.1202	0.9	0.63	731.9	5.9	724.6	7.0	702.4	22.6	731.9	5.9	104.2
Trial-2_0724	25	14673	0.2	15.8915	8.2	0.9954	10.3	0.1147	6.3	0.61	700.1	41.8	701.5	52.5	705.7	175.0	700.1	41.8	99.2
Trial-2_0725	52	24259	1.1	21.8030	11.3	0.1881	14.1	0.0297	8.4	0.59	188.9	15.6	175.0	22.6	-9.4	274.6	188.9	15.6	NA
Trial-2_0726	310	1167613	2.3	8.9708	1.4	4.5478	1.8	0.2959	1.1	0.64	1670.9	16.9	1739.8	14.9	1823.5	24.8	1823.5	24.8	91.6
Trial-2_0727	63	114569	1.8	9.5974	3.0	4.3565	3.9	0.3032	2.5	0.64	1707.4	37.4	1704.1	32.1	1700.1	55.0	1700.1	55.0	100.4
Trial-2_0728	184	69666	1.7	18.9506	4.1	0.4796	5.1	0.0659	3.1	0.60	411.5	12.3	397.8	16.9	318.8	93.1	411.5	12.3	129.1
Trial-2_0729	42	48830	0.9	6.1741	2.8	10.4038	3.8	0.4659	2.5	0.66	2465.4	51.1	2471.4	35.0	2476.3	48.0	2476.3	48.0	99.6
Trial-2_0730	49	31825	1.0	11.2095	3.8	3.0700	4.9	0.2496	3.1	0.63	1436.3	39.9	1425.2	37.5	1408.7	72.5	1408.7	72.5	102.0
Trial-2_0731	113	153438	2.0	13.5970	3.0	1.8296	3.9	0.1804	2.4	0.62	1069.3	23.8	1056.1	25.5	1029.0	61.5	1029.0	61.5	103.9
Trial-2_0732	3	121201	5.6	13.2436	1.6	1.7860	3.5	0.1715	3.1	0.89	1020.6	29.3	1040.4	22.8	1082.1	32.3	1082.1	32.3	94.3
Trial-2_0733	48	49874	2.2	13.0255	4.7	1.8682	6.0	0.1765	3.7	0.62	1047.8	36.2	1069.9	39.9	1115.3	94.4	1115.3	94.4	93.9
Trial-2_0734	239	28180	1.3	20.3837	4.4	0.2834	5.5	0.0419	3.3	0.60	264.5	8.6	253.3	12.4	150.6	104.1	264.5	8.6	NA
Trial-2_0735	91	405131	1.3	9.7316	2.5	4.2189	3.3	0.2978	2.1	0.64	1680.2	30.8	1677.7	26.7	1674.5	46.2	1674.5	46.2	100.3
Trial-2_0736	380	349502	3.1	5.3837	1.0	10.9417	1.3	0.4272	0.9	0.66	2293.2	16.6	2518.2	12.1	2704.8	16.1	2704.8	16.1	84.8
Trial-2_0737	41	30549	2.0	13.2307	3.0	1.8058	5.0	0.1733	4.0	0.80	1030.2	37.6	1047.6	32.4	1084.0	60.1	1084.0	60.1	95.0
Trial-2_0738	143	10724	1.3	21.8986	7.2	0.1664	9.0	0.0264	5.3	0.59	168.1	8.8	156.3	13.0	-20.0	175.1	168.1	8.8	NA
Trial-2_0739	105	72335	0.5	11.1878	2.5	3.0058	3.2	0.2439	2.1	0.63	1406.9	26.0	1409.1	24.7	1412.4	48.0	1412.4	48.0	99.6
Trial-2_0740	432	179808	4.7	13.0575	1.7	1.6204	2.1	0.1535	1.3	0.62	920.3	11.4	978.2	13.4	1110.4	33.2	1110.4	33.2	82.9
Trial-2_0741	129	168139	1.6	8.8769	2.0	5.0301	2.6	0.3238	1.7	0.64	1808.5	26.7	1824.4	22.3	1842.6	36.4	1842.6	36.4	98.1
Trial-2_0742	308	289246	1.2	12.8460	1.9	1.6807	2.5	0.1566	1.6	0.62	937.8	13.6	1001.2	15.9	1143.0	38.7	1143.0	38.7	82.0
Trial-2_0743	188	52286	4.3	11.9665	2.3	2.0838	2.9	0.1808	1.8	0.63	1071.6	18.2	1143.5	20.1	1282.5	44.4	1282.5	44.4	83.6
Trial-2_0744	221	95844	2.6	9.3643	1.6	4.2436	2.1	0.2882	1.4	0.64	1632.6	19.6	1682.5	17.4	1745.3	29.8	1745.3	29.8	93.5
Trial-2_0745	24	50558	1.4	5.3584	3.3	13.8503	4.4	0.5383	3.0	0.67	2776.2	66.6	2739.5	42.0	2712.6	54.5	2712.6	54.5	102.3
Trial-2_0746	169	172513	1.4	20.0199	4.9	0.3486	6.2	0.0506	3.7	0.60	318.3	11.5	303.7	16.2	192.7	115.1	318.3	11.5	NA
Trial-2_0747	121	166024	3.4	7.7370	1.9	6.9045	2.5	0.3874	1.6	0.65	2110.9	29.2	2099.2	22.2	2087.8	33.4	2087.8	33.4	101.1
Trial-2_0748	165	199880	1.0	11.1062	2.1	2.9564	2.7	0.2381	1.7	0.63	1377.0	21.2	1396.5	20.5	1426.3	40.0	1426.3	40.0	96.5
Trial-2_0749	33	34199	1.6	13.4662	1.8	1.6312	5.0	0.1593	4.6	0.93	953.0	40.8	982.3	31.2	1048.6	37.1	1048.6	37.1	90.9
Trial-2_0750	40	136016	0.9	5.6297	2.7	12.5154	3.6	0.5110	2.4	0.66	2661.0	52.3	2643.9	34.0	2630.8	44.9	2630.8	44.9	101.1

Trial-2_0751	146	132985	0.6	5.1068	1.4	13.7206	1.9	0.5082	1.3	0.67	2648.9	28.1	2730.6	18.4	2791.6	23.7	2791.6	23.7	94.9
Trial-2_0752	99	100862	0.7	5.1936	1.7	13.6894	2.3	0.5156	1.6	0.67	2680.7	34.0	2728.5	22.0	2764.0	28.5	2764.0	28.5	97.0
Trial-2_0753	268	97901	4.3	17.1104	3.4	0.5677	4.3	0.0704	2.6	0.61	438.8	11.1	456.5	15.9	546.4	74.9	438.8	11.1	80.3
Trial-2_0754	188	151306	2.1	12.7246	2.2	2.0720	2.8	0.1912	1.8	0.63	1128.0	18.4	1139.6	19.5	1161.8	43.9	1161.8	43.9	97.1
Trial-2_0755	24	12639	2.2	12.2104	6.1	2.3621	7.9	0.2092	4.9	0.62	1224.5	54.7	1231.2	56.1	1243.1	120.3	1243.1	120.3	98.5
Trial-2_0756	400	407225	1.9	11.0862	1.4	2.8462	1.8	0.2289	1.1	0.63	1328.5	13.4	1367.8	13.3	1429.8	26.0	1429.8	26.0	92.9
Trial-2_0757	447	313561	4.3	16.9156	2.2	0.7235	2.8	0.0888	1.7	0.61	548.2	8.9	552.7	11.9	571.4	48.1	548.2	8.9	95.9
Trial-2_0758	122	74447	2.0	13.1865	2.9	1.8005	3.7	0.1722	2.3	0.62	1024.2	21.9	1045.7	24.2	1090.7	58.0	1090.7	58.0	93.9
Trial-2_0759	512	143253	4.4	16.7252	2.0	0.7322	2.6	0.0888	1.6	0.61	548.5	8.2	557.8	11.0	596.0	43.9	548.5	8.2	92.0
Trial-2_0760	5	8580	1.7	12.6636	2.2	2.0977	2.5	0.1927	1.2	0.47	1135.8	12.0	1148.1	17.0	1171.3	43.4	1171.3	43.4	97.0
Trial-2_0761	376	33392	0.4	19.9625	3.3	0.2054	4.2	0.0297	2.6	0.62	188.9	4.8	189.7	7.2	199.3	75.6	188.9	4.8	NA
Trial-2_0762	48	502626	0.9	5.2992	2.5	13.2687	3.4	0.5100	2.2	0.66	2656.5	48.6	2699.0	31.7	2730.9	41.3	2730.9	41.3	97.3
Trial-2_0763	108	305258	2.5	13.8084	3.1	1.7175	3.9	0.1720	2.4	0.62	1023.1	23.2	1015.1	25.3	997.7	62.6	997.7	62.6	102.5
Trial-2_0764	80	8176	1.4	21.9913	8.7	0.2690	10.8	0.0429	6.4	0.59	270.8	16.9	241.9	23.2	-30.2	211.0	270.8	16.9	NA
Trial-2_0765	102	36607	1.0	16.5238	4.3	0.8469	5.4	0.1015	3.3	0.61	623.2	19.7	623.0	25.3	622.2	93.0	623.2	19.7	100.2
Trial-2_0766	174	315479	2.3	8.8246	1.8	4.5124	2.4	0.2888	1.5	0.64	1635.5	22.3	1733.3	19.9	1853.3	33.2	1853.3	33.2	88.2
Trial-2_0767	266	108336	1.5	13.1984	1.9	1.8382	2.5	0.1760	1.6	0.62	1044.9	15.0	1059.2	16.4	1088.9	39.0	1088.9	39.0	96.0
Trial-2_0768	72	61863	1.4	9.4332	2.7	4.4268	3.5	0.3029	2.3	0.64	1705.5	33.8	1717.4	29.2	1731.9	49.5	1731.9	49.5	98.5
Trial-2_0769	116	46259	1.8	13.0208	2.9	1.9616	3.7	0.1852	2.3	0.62	1095.6	23.2	1102.4	24.8	1116.0	57.6	1116.0	57.6	98.2
Trial-2_0770	275	1612491	1.2	13.5394	2.0	1.6163	2.5	0.1587	1.6	0.62	949.6	14.0	976.5	16.0	1037.6	40.3	1037.6	40.3	91.5
Trial-2_0771	62	8756	1.8	19.2397	6.8	0.5560	8.5	0.0776	5.1	0.60	481.7	23.6	448.9	30.7	284.3	154.7	481.7	23.6	169.4
Trial-2_0772	83	187625	1.4	11.1063	3.0	3.0884	3.9	0.2488	2.4	0.63	1432.1	31.4	1429.8	29.7	1426.3	57.4	1426.3	57.4	100.4
Trial-2_0773	209	920402	2.1	9.1409	1.7	4.3537	2.2	0.2886	1.4	0.64	1634.7	20.2	1703.6	18.0	1789.4	30.5	1789.4	30.5	91.4
Trial-2_0774	337	249819	7.1	4.5654	1.0	13.8270	1.3	0.4578	0.9	0.67	2430.0	18.0	2737.9	12.6	2973.6	15.9	2973.6	15.9	81.7
Trial-2_0775	129	80769	3.0	9.1995	2.1	4.5151	2.7	0.3013	1.7	0.64	1697.5	26.1	1733.8	22.7	1777.8	38.2	1777.8	38.2	95.5
Trial-2_0776	38	91545	1.2	9.7405	4.2	3.7441	5.4	0.2645	3.4	0.64	1512.8	46.4	1580.8	43.4	1672.8	77.2	1672.8	77.2	90.4
Trial-2_0777	199	68490	2.1	18.2670	3.8	0.4941	4.8	0.0655	2.9	0.60	408.8	11.5	407.7	16.2	401.7	86.1	408.8	11.5	101.7
Trial-2_0778	38	13102	0.6	13.1888	5.2	1.8472	6.6	0.1767	4.1	0.62	1048.9	40.0	1062.4	43.8	1090.4	104.4	1090.4	104.4	96.2
Trial-2_0779	110	455630	2.2	13.7071	3.2	1.6826	4.1	0.1673	2.6	0.62	997.1	23.7	1002.0	26.4	1012.7	65.9	1012.7	65.9	98.5
Trial-2_0780	29	59164	1.3	8.3100	4.2	5.9417	5.5	0.3581	3.6	0.65	1973.2	60.4	1967.4	47.9	1961.2	75.1	1961.2	75.1	100.6
Trial-2_0781	390	38021	3.3	19.7133	3.9	0.2784	4.9	0.0398	2.9	0.60	251.6	7.2	249.4	10.9	228.5	91.0	251.6	7.2	NA
Trial-2_0782	59	10943	1.2	22.1200	10.2	0.2310	12.6	0.0371	7.5	0.59	234.6	17.3	211.0	24.1	-44.4	248.0	234.6	17.3	NA
Trial-2_0783	104	30246	1.6	13.3553	3.3	1.8686	4.2	0.1810	2.6	0.62	1072.4	25.7	1070.0	27.6	1065.2	65.8	1065.2	65.8	100.7
Trial-2_0784	141	19093	1.2	19.9249	5.3	0.3596	6.6	0.0520	4.0	0.60	326.6	12.6	311.9	17.8	203.7	123.3	326.6	12.6	NA
Trial-2_0785	164	70234	1.3	13.3515	2.6	1.8275	3.3	0.1770	2.0	0.62	1050.4	19.8	1055.4	21.5	1065.8	51.6	1065.8	51.6	98.6
Trial-2_0786	74	68422	1.2	11.1156	3.1	3.0595	4.0	0.2467	2.6	0.63	1421.2	32.6	1422.6	30.9	1424.7	59.8	1424.7	59.8	99.7
Trial-2_0787	67	64616	2.7	12.9660	3.9	2.0402	5.0	0.1919	3.1	0.62	1131.4	32.3	1129.0	34.1	1124.4	77.8	1124.4	77.8	100.6
Trial-2_0788	103	58600	1.6	13.9179	3.3	1.6703	4.2	0.1686	2.6	0.62	1004.4	24.5	997.3	27.0	981.7	67.8	981.7	67.8	102.3
Trial-2_0789	79	10607	1.0	23.8270	10.4	0.1592	12.9	0.0275	7.6	0.59	175.0	13.1	150.0	18.0	-228.4	263.7	175.0	13.1	NA
Trial-2_0790	183	168674	1.5	9.1161	1.8	4.6216	2.4	0.3056	1.5	0.64	1718.9	23.1	1753.2	19.9	1794.3	33.3	1794.3	33.3	95.8
Trial-2_0791	98	212782	0.7	11.2127	2.7	3.0063	3.5	0.2445	2.2	0.63	1409.9	28.4	1409.2	27.0	1408.1	52.6	1408.1	52.6	100.1
Trial-2_0792	77	58581	1.6	9.0188	2.8	4.9689	3.7	0.3250	2.4	0.64	1814.2	37.3	1814.1	31.1	1813.9	51.3	1813.9	51.3	100.0
Trial-2_0793	155	240876	2.3	13.1853	2.6	1.9410	3.3	0.1856	2.1	0.62	1097.6	20.9	1095.3	22.2	1090.9	52.0	1090.9	52.0	100.6
Trial-2_0794	6	107128	1.9	12.4554	2.7	2.2559	3.0	0.2038	1.1	0.38	1195.6	12.1	1198.6	20.8	1204.0	53.9	1204.0	53.9	99.3
Trial-2_0795	33	169217	1.2	13.1144	2.1	1.9494	5.3	0.1854	4.8	0.92	1096.5	48.7	1098.2	35.4	1101.7	42.3	1101.7	42.3	99.5
Trial-2_0796	96	80686	0.9	12.9754	3.2	2.0503	4.1	0.1929	2.6	0.62	1137.3	26.8	1132.4	28.1	1123.0	64.2	1123.0	64.2	101.3
Trial-2_0797	320	184369	3.1	13.1638	1.9	1.7545	2.5	0.1675	1.5	0.62	998.4	14.3	1028.8	16.1	1094.2	38.9	1094.2	38.9	91.2
Trial-2_0798	151	102926	2.3	9.7318	2.1	4.0259	2.7	0.2842	1.7	0.64	1612.3	24.5	1639.4	21.9	1674.5	38.2	1674.5	38.2	96.3
Trial-2_0799	30	31247	0.8	13.3121	6.3	1.8269	8.0	0.1764	5.0	0.62	1047.2	48.0	1055.2	52.5	1071.7	125.9	1071.7	125.9	97.7
Trial-2_0800	36	47089	2.1	12.7816	5.2	2.1103	6.7	0.1956	4.2	0.62	1151.8	43.9	1152.2	46.0	1152.9	103.4	1152.9	103.4	99.9
Trial-2_0801	164	80367	1.5	13.0299	2.6	1.8980	3.3	0.1794	2.1	0.62	1063.5	20.2	1080.4	22.0	1114.6	51.6	1114.6	51.6	95.4
Trial-2_0802	50	141914	0.7	5.4079	2.6	13.1777	3.4	0.5169	2.3	0.66	2685.8	50.1	2692.5	32.4	2697.4	42.4	2697.4	42.4	99.6
Trial-2_0803	74	88690	1.6	13.4092	3.8	1.8645	4.9	0.1813	3.0	0.62	1074.2	30.1	1068.6	32.3	1057.1	76.9	1057.1	76.9	101.6
Trial-2_0804	104	135129	1.2	11.0183	2.6	3.0666	3.4	0.2451	2.2	0.63	1412.9	27.5	1424.4	26.2	1441.5	50.4	1441.5	50.4	98.0
Trial-2_0805	132	25793	1.4	20.5417	6.6	0.2749	8.2	0.0410	4.9	0.59	258.8	12.4	246.6	18.0	132.5	155.8	258.8	12.4	NA
Trial-2_0806	80	4937	1.4	22.1792	8.8	0.2579	10.9	0.0415	6.4	0.59	262.0	16.5	232.9	22.6	-50.9	213.5	262.0	16.5	NA
Trial-2_0807	139	184734	2.1	12.9571	2.7	1.9829	3.5	0.1863	2.2	0.62	1101.5	22.2	1109.7	23.7	1125.8	54.7	1125.8	54.7	97.8
Trial-2_0808	36	391637	1.3	5.3544	3.0	13.6132	4.0	0.5287	2.6	0.66	2735.8	58.8	2723.2	37.6	2713.9	49.1	2713.9	49.1	100.8
Trial-2_0809	32	30441	2.5	8.9109	4.3	4.9890	5.6	0.3224	3.6	0.64	1801.6	56.3	1817.5	47.3	1835.7	77.6	1835.7	77.6	98.1
Trial-2_0810	43	28071	1.3	11.1330	4.1	3.1824	5.3	0.2570	3.3	0.63	1474.3	43.9	1452.9	40.6	1421.7	77.8	1421.7	77.8	103.7
Trial-2_0811	283	300285	2.9	9.1840	1.5	4.2994	1.9	0.2864	1.2	0.64	1623.4	17.8	1693.3	15.9	1780.8	27.0	1780.8	27.0	91.2
Trial-2_0812	59	33737	1.5	13.2616	4.5	1.7791	5.7	0.1711	3.5	0.62	1018.3	33.1	1037.9	36.9	1079.3	89.4	1079.3	89.4	94.3
Trial-2_0813	292	426652	1.4	10.9920	1.7	2.9114	2.2	0.2321	1.4	0.63	1345.5	16.8	1384.9	16.5	1446.1	32.2	1446.1	32.2	93.0
Trial-2_0814	193	330873	1.3	10.8122	2.0	3.1471	2.6	0.2468	1.6	0.63	1421.9	20.8	1444.3	19.8	1477.4	37.8	1477.4	37.8	96.2
Trial-2_0815	112	385069	0.8	5.2763	1.8	13.4995	2.4	0.5166	1.6	0.66	2684.7	34.6	2715.3	22.5	2738.1	29.2	2738.1	29.2	98.1
Trial-2_0816	46	133685	0.8	5.1609	2.6	14.2951	3.5	0.5351	2.4	0.66	2762.8	52.9	2769.5	33.6	2774.4	43.4	2774.4	43.4	99.6
Trial-2_0817	846	110226	1.4	19.8549	2.9	0.2120	3.6	0.0305	2.1	0.60	193.9	4.1	195.2	6.3	211.9	66.3	193.9	4.1	NA
Trial-2_0818	153	50482	1.9	13.4212	2.8	1.7702	3.6	0.1723	2.2	0.62	1024.8	20.9	1034.6	23.0	1055.3	56.0	1055.3	56.0	97.1
Trial-2_0819	23	32848	1.3	20.8904	15.9	0.2687	16.2	0.0407	2.6	0.16	257.3	6.5	241.7	34.8	92.8	379.8	257.3	6.5	NA
Trial-2_0820	132	55612	2.1	12.5956	2.8	2.1673	3.6	0.1980	2.2	0.63	1164.5	23.8	1170.6	24.8	1182.0	55.2	1182.0	55.2	98.5
Trial-2_0821	36	3487	0.9	21.0469	13.5	0.2610	16.7	0.0398	9.9	0.59	251.9	24.4	235.5	35.1	75.1	321.0	251.9	24.4	NA
Trial-2_0822	259	203761	1.6	12.6556	2.0	2.0282	2.6	0.1862	1.6	0.63	1100.5	16.2	1125.0	17.4	1172.6	39.4	1172.6	39.4	93.9
Trial-2_0823	48	6418	1.4	23.1878	11.2	0.2297	13.8	0.0386	8.2	0.59	244.3	19.6	209.9	26.3	-160.3	278.2	244.3	19.6	NA
Trial-2_0824	236	90965	2.0	13.1663	2.2	1.9316	2.8	0.1844	1.7	0.62	1091.2	17.4	1092.1	18.6	1093.8	43.5	1093.8	43.5	99.8
Trial-2_0825	190	158926	1.9	10.8217	1.9	3.2380	2.5	0.2541	1.6	0.64	1459.8	20.8	1466.3	19.5	1475.8	36.7	1475.8	36.7	98.9
Trial-2_0826	49	21351	1.1	12.8816	4.7	2.0224	6.0	0.1889	3.7	0.62	1115.7	38.0	1123.1	40.6	1137.4	93.0	1137.4	93.0	98.1
Trial-2_0827	140	87392	1.2	10.8795	2.4	3.1830	3.1	0.2512	2.0	0.63	1444.4	25.6	1453.0	24.2	1465.6	46.0	1465.6	46.0	98.6
Trial-2_0828	88	169324	0.9	11.0690	3.0	2.9933	3.8	0.2403	2.4	0.63	1388.3	30.3	1405.9	29.3	1432.8	56.9	1432.8	56.9	96.9
Trial-2_0829	37	13716	2.1	13.2845	6.0	1.7950	7.6	0.1729	4.7	0.62	1028.3	44.6	1043.7	49.5	1075.9	119.7	1075.9	119.7	95.6
Trial-2_0830	301	381644	1.6	12.9995	2.0	1.8202	2.5	0.1716	1.6	0.62	1021.0	14.8	1052.7	16.5	1119.3	39.2	1119.3	39.2	91.2
Trial-2_0831	22	36860	0.9	13.1926	3.0	1.8761	6.3	0.1795	5.5	0.88	1064.3	54.1	1072.7	41.5	1089.8	59.5	1089.8	59.5	97.7
Trial-2_0832	50	19617	0.8	18.4271	7.6	0.5656	9.6	0.0756	5.8	0.60	469.8	26.2	455.2	35.1	382.2	171.8	469.8	26.2	122.9
Trial-2_0833	204	206088	1.3	5.6042	1.3	11.2142	1.8	0.4558	1.2	0.66	2421.0	23.8	2541.1	16.6	2638.4	22.2	2638.4	22.2	91.8
Trial-2_0834	252	464318	2.6	13.0889	2.1	1.8564	2.6	0.1762	1.7	0.62	1046.4	16.0	1065.7	17.5	1105.6	41.4	1105.6	41.4	94.6
Trial-2_0835	257	464303	3.4	8.9036	1.5	4.6296	2.0	0.2990	1.3	0.64	1686.1	18.8	1754.6	16.5	1837.2	27.4	1837.2	27.4	91.8
Trial-2_0836	108	53078	0.9	13.9832	3.3	1.6585	4.2	0.1682	2.6	0.62	1002.2	24.3	992.8	26.8	972.1	67.7	972.1	67.7	103.1
Trial-2_0837	154	307083	1.9	13.6429	2.8	1.6932	3.6	0.1675	2.2	0.62	998.5	20.4	1006.0	22.7	1022.2	56.4	1022.2	56.4	97.7
Trial-2_0838	65	28905	1.2	12.1690	3.8	2.4145	4.9	0.2131	3.0	0.63	1245.3	34.5	1246.9	34.9	1249.7	74.1	1249.7	74.1	99.6
Trial-2_0839	231	171923	1.9	20.3777	5.6	0.2048	7.0	0.0303	4.2	0.59	192.2	7.9	189.1	12.1	151.3	131.9	192.2	7.9	NA
Trial-2_0840	754	574161	3.7	12.6935	1.3	1.7236	1.6	0.1587	1.0	0.62	949.4	8.9	1017.4	10.4	1166.6	24.9	1166.6	24.9	81.4
Trial-2_0841	76	34932	2.2	13.1560	3.8	1.9593	4.8	0.1870	3.0	0.62	1104.8	30.5	1101.7	32.5	1095.4	75.6	1095.4	75.6	100.9
Trial-2_0842	74	75165	1.1	10.9963	3.3	3.0923	4.2	0.2466	2.7	0.63	1421.0	34.3	1430.8	32.6	1445.3	62.7	1445.3	62.7	98.3
Trial-2_0843	33	32182	1.5	13.1153	5.8	1.9456	7.4	0.1851	4.6	0.62	1094.6	46.3	1096.9	49.6	1101.5	115.9	1101.5	115.9	99.4
Trial-2_0844	77	332674	2.7	12.8065	3.8	2.0097	4.9	0.1867	3.0	0.62	1103.3	30.9	1118.8	33.1	1149.1	75.7	1149.1	75.7	96.0
Trial-2_0845	13	5688	1.1	13.3780	9.5	1.8055	12.1	0.1752	7.5	0.62	1040.6	72.2	1047.4	79.3	1061.8	191.4	1061.8	191.4	98.0
Trial-2_0846	85	126652	2.0	13.6411	3.7	1.7168	4.7	0.1699	2.9	0.62	1011.3	27.2	1014.8	30.0	1022.5	74.1	1022.5	74.1	98.9
Trial-2_0847	170	120466	0.4	13.3270	2.6	1.8048	3.3	0.1744	2.0	0.62	1036.6	19.6	1047.2	21.5	1069.5	51.8	1069.5	51.8	96.9
Trial-2_0848	781	728505	13.8	12.5087	1.3	1.7749	1.6	0.1610	1.0	0.62	962.5	9.0	1036.3	10.5	1195.6	24.8	1195.6	24.8	80.5
Trial-2_0849	54	20499	2.3	13.0578	4.4	2.0017	5.6	0.1896	3.5	0.62	1119.1	36.1	1116.1	38.2	1110.3	88.1	1110.3	88.1	100.8
Trial-2_0850	49	22798	6.2	17.2224	7.1	0.7191	8.9	0.0898	5.4	0.61	554.5	28.7	550.1	37.8	532.2	154.8	554.5	28.7	104.2
Trial-2_0851	220	24153	1.9	19.8104	5.1	0.2790	6.4	0.0401	3.8	0.60	253.4	9.5	249.9	14.2	217.1	118.8	253.4	9.5	NA
Trial-2_0852	37	128347	0.9	5.3928	2.9	13.3782	3.9	0.5232	2.6	0.66	2712.9	57.6	2706.7	37.0	2702.1	48.4	2702.1	48.4	100.4
Trial-2_0853	17	57939	1.1	12.8152	7.5	2.1042	9.6	0.1956	6.0	0.62	1151.5	62.9	1150.2	66.0	1147.7	148.9	1147.7	148.9	100.3
Trial-2_0854	29	121784	0.3	8.4490	3.8	5.5962	5.3	0.3429	3.6	0.69	1900.7	59.5	1915.5	45.3	1931.6	68.3	1931.6	68.3	98.4
Trial-2_0855	23	40419	1.1	8.4948	4.7	5.7643	6.1	0.3551	3.9	0.64	1959.1	66.7	1941.1	53.0	1921.9	84.0	1921.9	84.0	101.9
Trial-2_0856	88	686500	1.5	5.1158	1.9	13.9928	2.5	0.5192	1.7	0.67	2695.7	37.1	2749.2	24.0	2788.8	30.9	2788.8	30.9	96.7
Trial-2_0857	68	46610	2.1	13.3132	4.0	1.8489	5.1	0.1785	3.2	0.62	1058.9	30.9	1063.0	33.5	1071.6	79.9	1071.6	79.9	98.8
Trial-2_0858	66	25899	1.1	13.6790	4.4	1.7043	5.6	0.1691	3.5	0.62	1007.0	32.3	1010.1	35.8	1016.8	88.8	1016.8	88.8	99.0
Trial-2_0859	45	117547	0.8	5.3362	2.7	13.0795	3.6	0.5062	2.4	0.66	2640.4	51.3	2685.4	33.6	2719.5	43.9	2719.5	43.9	97.1
Trial-2_0860	45	20613	1.8	13.3947	5.1	1.7379	6.6	0.1688	4.1	0.62	1005.7	37.9	1022.7	42.3	1059.3	103.6	1059.3	103.6	94.9
Trial-2_0861	59	150699	0.9	10.8746	3.4	3.1311	4.4	0.2469	2.8	0.63	1422.7	35.9	1440.4	34.2	1466.5	65.3	1466.5	65.3	97.0
Trial-2_0862	75	71978	1.6	11.5665	3.3	2.7427	4.3	0.2301	2.7	0.63	1334.9	32.6	1340.1	31.9	1348.4	64.3	1348.4	64.3	99.0
Trial-2_0863	22	44733	1.0	13.0820	7.4	1.9325	9.5	0.1834	5.9	0.62	1085.3	58.7	1092.4	63.5	1106.6	148.7	1106.6	148.7	98.1
Trial-2_0864	205	234152	3.2	18.3048	4.1	0.4540	5.2	0.0603	3.1	0.60	377.3	11.4	380.1	16.3	397.1	92.3	377.3	11.4	NA
Trial-2_0865	221	36797	1.6	20.3440	5.3	0.2738	6.6	0.0404	3.9	0.59	255.3	9.8	245.7	14.4	155.2	124.5	255.3	9.8	NA
Trial-2_0866	16	34659	0.9	13.3515	8.4	1.7915	10.7	0.1735	6.6	0.62	1031.3	63.2	1042.4	69.8	1065.8	168.9	1065.8	168.9	96.8
Trial-2_0867	5	63705	1.3	13.2415	2.8	1.8415	3.2	0.1768	1.6	0.50	1049.7	15.5	1060.4	21.1	1082.4	55.8	1082.4	55.8	97.0
Trial-2_0868	58	47906	9.6	17.8755	7.9	0.4759	9.9	0.0617	6.0	0.60	385.9	22.3	395.2	32.3	450.1	175.0	385.9	22.3	NA
Trial-2_0869	66	129369	1.8	11.8360	3.5	2.5814	4.6	0.2216	2.9	0.63	1290.3	33.5	1295.4	33.3	1303.8	68.7	1303.8	68.7	99.0
Trial-2_0870	144	533680	6.2	13.9746	2.9	1.5718	3.7	0.1593	2.3	0.62	952.9	20.5	959.1	23.1	973.4	59.5	973.4	59.5	97.9
Trial-2_0871	386	299195	2.5	12.3968	1.6	2.0002	2.1	0.1798	1.3	0.63	1066.1	13.0	1115.6	14.3	1213.3	32.5	1213.3	32.5	87.9
Trial-2_0872	51	5450	0.5	23.9796	12.7	0.1583	15.6	0.0275	9.1	0.58	175.1	15.8	149.2	21.7	-244.5	321.3	175.1	15.8	NA
Trial-2_0873	41	78685	1.3	5.9156	3.0	11.2328	3.9	0.4819	2.6	0.66	2535.7	54.7	2542.6	36.8	2548.2	49.7	2548.2	49.7	99.5
Trial-2_0874	53	22930	1.9	12.6706	4.4	2.1117	5.6	0.1941	3.5	0.62	1143.3	36.4	1152.6	38.4	1170.2	86.4	1170.2	86.4	97.7
Trial-2_0875	212	293003	1.7	12.8328	2.2	1.9233	2.9	0.1790	1.8	0.63	1061.5	17.8	1089.2	19.3	1145.0	44.6	1145.0	44.6	92.7
Trial-2_0876	200	316824	1.1	9.6009	1.8	4.0939	2.3	0.2851	1.5	0.64	1616.8	21.2	1653.1	18.9	1699.5	32.7	1699.5	32.7	95.1
Trial-2_0877	273	144363	3.4	11.9431	1.9	2.3489	2.4	0.2035	1.5	0.63	1193.9	16.5	1227.2	17.1	1286.3	36.5	1286.3	36.5	92.8
Trial-2_0878	56	46732	1.1	11.0532	3.7	3.1731	4.7	0.2544	3.0	0.63	1461.0	39.0	1450.6	36.4	1435.5	69.7	1435.5	69.7	101.8
Trial-2_0879	221	452854	1.3	8.6386	1.6	4.8619	2.1	0.3046	1.4	0.64	1714.1	20.6	1795.7	17.9	1891.7	29.3	1891.7	29.3	90.6
Trial-2_0880	148	64254	1.1	20.6065	6.3	0.2558	7.8	0.0382	4.6	0.60	241.8	11.0	231.3	16.1	125.1	147.5	241.8	11.0	NA
Trial-2_0881	119	63341	1.5	13.8091	3.1	1.7000	4.0	0.1703	2.5	0.62	1013.6	23.3	1008.5	25.6	997.6	63.8	997.6	63.8	101.6
Trial-2_0882	117	69709	2.4	13.1022	2.9	2.0566	3.7	0.1954	2.3	0.62	1150.7	24.5	1134.5	25.5	1103.6	58.2	1103.6	58.2	104.3
Trial-2_0883	229	11320	0.7	21.2967	5.8	0.1885	7.3	0.0291	4.3	0.59	185.0	7.8	175.3	11.7	47.0	139.7	185.0	7.8	NA
Trial-2_0884	170	112703	1.6	8.6493	1.8	4.9107	2.3	0.3081	1.5	0.64	1731.1	22.8	1804.1	19.6	1889.5	32.0	1889.5	32.0	91.6
Trial-2_0885	285	1564605	1.5	11.3533	1.8	2.6087	2.3	0.2148	1.4	0.63	1254.3	16.4	1303.1	16.7	1384.2	34.0	1384.2	34.0	90.6
Trial-2_0886	42	130163	0.5	11.3614	4.3	2.9860	5.5	0.2460	3.5	0.63	1418.1	44.6	1404.1	42.2	1382.8	82.6	1382.8	82.6	102.5
Trial-2_0887	113	312664	0.9	11.0871	2.5	3.1163	3.3	0.2506	2.1	0.63	1441.5	26.6	1436.7	25.0	1429.6	48.0	1429.6	48.0	100.8
Trial-2_0888	163	24202	1.8	21.1468	6.3	0.2830	7.8	0.0434	4.7	0.59	273.9	12.5	253.0	17.5	63.8	150.0	273.9	12.5	NA

Trial-2_0889	47	119822	1.6	13.0558	4.8	1.9667	6.1	0.1862	3.8	0.62	1100.9	38.4	1104.2	41.1	1110.6	95.6	1110.6	95.6	99.1
Trial-2_0890	79	66702	1.3	13.6200	4.1	1.4812	5.3	0.1463	3.3	0.62	880.3	27.0	922.7	32.0	1025.6	83.7	1025.6	83.7	85.8
Trial-2_0891	204	72433	1.3	10.8513	1.9	3.0665	2.5	0.2413	1.6	0.63	1393.6	19.9	1424.4	19.2	1470.6	36.8	1470.6	36.8	94.8
Trial-2_0892	110	82622	2.6	11.1699	2.7	3.0581	3.4	0.2477	2.2	0.63	1426.8	27.9	1422.3	26.4	1415.4	51.1	1415.4	51.1	100.8
Trial-2_0893	81	35635	2.1	13.2138	3.8	1.8248	4.9	0.1749	3.1	0.62	1038.9	29.3	1054.4	32.2	1086.6	77.0	1086.6	77.0	95.6
Trial-2_0894	224	279547	3.1	13.4257	2.2	1.7764	2.9	0.1730	1.8	0.62	1028.4	17.0	1036.9	18.6	1054.6	45.2	1054.6	45.2	97.5
Trial-2_0895	236	146199	2.3	10.7950	1.8	3.0505	2.4	0.2388	1.5	0.63	1380.6	18.7	1420.4	18.2	1480.4	34.9	1480.4	34.9	93.3
Trial-2_0896	186	75877	1.6	13.8528	2.6	1.6286	3.3	0.1636	2.0	0.62	976.9	18.3	981.3	20.5	991.3	51.9	991.3	51.9	98.6
Trial-2_0897	58	56043	1.9	5.2462	2.3	13.9140	3.1	0.5294	2.1	0.67	2739.0	45.9	2743.9	29.3	2747.4	38.0	2747.4	38.0	99.7
Trial-2_0898	95	89534	1.9	10.0586	2.6	3.8108	3.4	0.2780	2.2	0.64	1581.3	30.6	1595.0	27.5	1613.2	49.0	1613.2	49.0	98.0
Trial-2_0899	126	304226	2.2	13.7292	3.0	1.7178	3.8	0.1710	2.4	0.62	1017.9	22.5	1015.2	24.7	1009.4	61.1	1009.4	61.1	100.8
Trial-2_0900	115	63503	1.8	11.6387	2.7	2.6842	3.5	0.2266	2.2	0.63	1316.5	26.1	1324.1	25.7	1336.4	52.2	1336.4	52.2	98.5
Trial-2_0901	276	287825	2.3	9.5606	1.6	3.7716	2.1	0.2615	1.3	0.64	1497.6	17.6	1586.7	16.5	1707.2	29.1	1707.2	29.1	87.7
Trial-2_0902	163	149123	2.2	10.9428	2.2	3.0972	2.8	0.2458	1.8	0.63	1416.8	22.6	1432.0	21.5	1454.6	41.3	1454.6	41.3	97.4
Trial-2_0903	70	138150	1.5	11.3773	3.3	3.0365	4.2	0.2506	2.7	0.63	1441.3	34.5	1416.8	32.3	1380.2	62.9	1380.2	62.9	104.4
Trial-2_0904	61	45368	0.8	11.2982	3.6	2.8735	4.6	0.2355	2.9	0.63	1363.1	36.1	1375.0	35.0	1393.5	69.0	1393.5	69.0	97.8
Trial-2_0905	191	61760	2.4	12.5192	2.3	2.1072	3.0	0.1913	1.9	0.63	1128.6	19.4	1151.2	20.6	1193.9	46.1	1193.9	46.1	94.5
Trial-2_0906	163	152689	1.1	9.9400	2.0	3.9592	2.6	0.2854	1.7	0.64	1618.6	24.0	1625.9	21.3	1635.3	37.5	1635.3	37.5	99.0
Trial-2_0907	129	111213	1.4	13.6349	2.9	1.8163	3.8	0.1796	2.3	0.62	1064.9	23.0	1051.4	24.7	1023.4	59.7	1023.4	59.7	104.1
Trial-2_0908	94	37596	0.8	20.9292	7.6	0.2616	9.5	0.0397	5.7	0.60	251.0	14.0	236.0	20.0	88.4	181.3	251.0	14.0	NA
Trial-2_0909	125	130870	1.8	12.4754	2.8	2.2337	3.6	0.2021	2.2	0.63	1186.6	24.1	1191.7	24.9	1200.9	54.6	1200.9	54.6	98.8
Trial-2_0910	367	37324	1.3	21.0131	4.7	0.1756	5.9	0.0268	3.5	0.59	170.3	5.9	164.3	9.0	78.9	112.8	170.3	5.9	NA
Trial-2_0911	18	11824	0.6	12.2748	6.7	2.4829	8.6	0.2210	5.4	0.63	1287.4	63.0	1267.1	62.3	1232.8	131.3	1232.8	131.3	104.4
Trial-2_0912	710	84860	0.9	20.6656	3.4	0.1781	4.3	0.0267	2.5	0.59	169.9	4.2	166.5	6.5	118.4	80.8	169.9	4.2	NA
Trial-2_0913	586	452538	1.8	10.9733	1.3	2.6306	1.6	0.2094	1.0	0.63	1225.4	11.5	1309.2	12.0	1449.3	24.0	1449.3	24.0	84.6
Trial-2_0914	61	23903	2.0	19.0567	6.8	0.5463	8.6	0.0755	5.2	0.60	469.2	23.3	442.6	30.7	306.2	155.8	469.2	23.3	153.3
Trial-2_0915	95	63906	2.6	13.5885	3.6	1.6346	4.6	0.1611	2.8	0.62	962.9	25.4	983.6	28.8	1030.3	72.4	1030.3	72.4	93.5
Trial-2_0916	28	11790	1.8	19.8521	10.3	0.5217	12.9	0.0751	7.7	0.60	466.9	34.8	426.3	44.8	212.2	239.1	466.9	34.8	220.0
Trial-2_0917	217	279252	3.4	13.5858	2.3	1.7277	2.9	0.1702	1.8	0.62	1013.4	17.1	1018.9	18.9	1030.7	46.5	1030.7	46.5	98.3
Trial-2_0918	335	18123	1.9	12.4760	2.0	2.1764	2.5	0.1969	1.6	0.63	1158.8	16.6	1173.5	17.4	1200.8	38.5	1200.8	38.5	96.5
Trial-2_0919	75	62199	1.4	11.4100	3.4	2.7798	4.3	0.2300	2.7	0.63	1334.7	32.6	1350.1	32.2	1374.6	64.6	1374.6	64.6	97.1
Trial-2_0920	57	34766	0.9	12.5649	4.2	2.1638	5.4	0.1972	3.4	0.62	1160.2	35.7	1169.5	37.5	1186.8	83.3	1186.8	83.3	97.8
Trial-2_0921	27	19342	1.6	7.7456	3.9	7.1900	5.2	0.4039	3.4	0.65	2187.0	62.8	2135.3	46.4	2085.8	69.5	2085.8	69.5	104.9
Trial-2_0922	117	96327	1.1	12.3146	2.7	2.3541	3.5	0.2103	2.2	0.63	1230.2	24.7	1228.8	25.1	1226.4	53.8	1226.4	53.8	100.3
Trial-2_0923	120	79159	15.9	16.0485	6.3	0.8708	8.0	0.1014	4.9	0.61	622.4	28.8	636.0	37.8	684.8	135.6	622.4	28.8	90.9
Trial-2_0924	421	286225	3.0	10.9026	1.4	2.9564	1.8	0.2338	1.1	0.63	1354.3	13.7	1396.5	13.4	1461.6	25.9	1461.6	25.9	92.7
Trial-2_0925	37	44266	0.9	11.0923	4.4	3.1113	5.7	0.2503	3.6	0.63	1440.0	46.6	1435.5	44.0	1428.7	84.6	1428.7	84.6	100.8
Trial-2_0926	61	222906	0.7	7.9000	2.7	6.6329	3.5	0.3800	2.3	0.65	2076.5	40.6	2063.7	31.1	2051.0	47.3	2051.0	47.3	101.2
Trial-2_0927	90	62266	2.3	12.8806	3.2	2.0837	4.1	0.1947	2.5	0.63	1146.5	26.7	1143.5	27.9	1137.6	63.2	1137.6	63.2	100.8
Trial-2_0928	25	92032	0.8	5.3641	3.5	13.3737	4.6	0.5203	3.1	0.66	2700.4	67.7	2706.4	43.8	2710.9	57.4	2710.9	57.4	99.6
Trial-2_0929	72	62749	2.3	13.1367	3.7	1.9332	4.8	0.1842	3.0	0.62	1089.8	29.9	1092.7	32.0	1098.3	74.9	1098.3	74.9	99.2
Trial-2_0930	37	30196	2.7	9.0111	3.8	4.9989	4.9	0.3267	3.2	0.64	1822.4	50.2	1819.1	41.7	1815.4	68.5	1815.4	68.5	100.4
Trial-2_0931	59	119930	1.2	6.1248	2.4	10.4339	3.2	0.4635	2.1	0.66	2454.9	43.0	2474.1	29.6	2489.8	40.4	2489.8	40.4	98.6
Trial-2_0932	133	112049	1.1	13.2677	2.8	1.9314	3.5	0.1858	2.2	0.62	1098.8	22.2	1092.0	23.6	1078.4	55.4	1078.4	55.4	101.9
Trial-2_0933	6	6630	1.9	12.7653	3.4	2.0653	3.7	0.1912	1.5	0.40	1127.9	15.2	1137.4	25.1	1155.4	66.9	1155.4	66.9	97.6
Trial-2_0934	298	324564	3.1	12.7710	1.9	1.8531	2.4	0.1716	1.5	0.62	1021.1	14.0	1064.5	15.6	1154.5	36.7	1154.5	36.7	88.4
Trial-2_0935	81	466819	1.6	11.2119	3.0	2.9451	3.9	0.2395	2.5	0.63	1384.0	30.7	1393.6	29.6	1408.2	57.8	1408.2	57.8	98.3
Trial-2_0936	26	22838	1.5	13.1951	6.3	1.9324	8.0	0.1849	5.0	0.62	1093.8	50.1	1092.4	53.7	1089.4	125.9	1089.4	125.9	100.4
Trial-2_0937	42	13636	1.9	18.8082	7.7	0.6479	9.5	0.0884	5.6	0.59	546.0	29.5	507.2	38.0	336.0	173.9	546.0	29.5	162.5
Trial-2_0938	199	631000	2.1	10.8420	1.9	3.1731	2.5	0.2495	1.6	0.63	1436.0	20.1	1450.6	19.0	1472.2	36.1	1472.2	36.1	97.5
Trial-2_0939	40	7320	1.3	22.6549	11.6	0.2429	14.3	0.0399	8.5	0.59	252.3	21.0	220.8	28.5	-102.8	284.9	252.3	21.0	NA
Trial-2_0940	47	43037	0.9	12.5117	4.5	2.2301	5.8	0.2024	3.6	0.63	1188.0	39.0	1190.6	40.4	1195.1	88.7	1195.1	88.7	99.4
Trial-2_0941	117	19584	1.7	18.6840	5.0	0.5158	6.3	0.0699	3.8	0.60	435.5	16.0	422.3	21.8	351.0	113.6	435.5	16.0	124.1
Trial-2_0942	29	25788	2.1	13.1079	6.1	1.8299	7.8	0.1740	4.8	0.62	1033.9	46.0	1056.2	51.0	1102.7	121.9	1102.7	121.9	93.8
Trial-2_0943	293	44600	2.2	19.9691	4.1	0.2949	5.1	0.0427	3.0	0.60	269.7	8.0	262.4	11.7	198.6	94.4	269.7	8.0	NA
Trial-2_0944	101	36042	1.7	13.3996	3.1	1.8904	4.0	0.1837	2.5	0.62	1087.2	25.0	1077.7	26.6	1058.5	63.0	1058.5	63.0	102.7
Trial-2_0945	206	100962	2.1	13.4960	2.3	1.7238	3.0	0.1687	1.9	0.62	1005.1	17.3	1017.4	19.2	1044.1	47.2	1044.1	47.2	96.3
Trial-2_0946	46	96456	1.2	16.5165	6.4	0.8781	8.1	0.1052	4.9	0.61	644.7	30.2	639.9	38.5	623.1	138.9	644.7	30.2	103.5
Trial-2_0947	49	61149	1.4	11.8814	4.0	2.6690	5.2	0.2300	3.3	0.63	1334.5	39.4	1319.9	38.4	1296.4	78.4	1296.4	78.4	102.9
Trial-2_0948	226	516722	6.4	9.6249	1.7	3.9889	2.2	0.2785	1.4	0.64	1583.6	19.6	1631.9	17.7	1694.9	30.9	1694.9	30.9	93.4
Trial-2_0949	39	121673	1.9	12.0592	5.0	2.3717	6.4	0.2074	4.0	0.62	1215.1	44.1	1234.1	45.6	1267.5	97.3	1267.5	97.3	95.9
Trial-2_0950	83	217592	1.0	8.9470	2.5	5.1942	3.3	0.3370	2.1	0.64	1872.4	34.3	1851.7	27.9	1828.4	45.5	1828.4	45.5	102.4
Trial-2_0951	289	15788	2.1	22.7318	6.9	0.0965	8.5	0.0159	5.0	0.59	101.8	5.1	93.5	7.6	-111.2	169.3	101.8	5.1	NA
Trial-2_0952	80	330261	2.2	6.5925	2.2	9.2817	2.9	0.4438	1.9	0.66	2367.6	37.1	2366.2	26.2	2365.0	36.7	2365.0	36.7	100.1
Trial-2_0953	61	574717	1.8	9.0972	3.1	4.7232	4.0	0.3116	2.5	0.64	1748.8	39.1	1771.4	33.4	1798.1	55.8	1798.1	55.8	97.3
Trial-2_0954	38	7222	1.3	20.5747	9.1	0.4741	11.4	0.0708	6.8	0.60	440.7	29.0	394.0	37.2	128.8	215.1	440.7	29.0	342.2
Trial-2_0955	339	840802	4.7	13.6915	1.9	1.5793	2.4	0.1568	1.5	0.62	939.1	13.2	962.1	15.1	1015.0	38.7	1015.0	38.7	92.5
Trial-2_0956	109	168412	1.2	11.0771	2.6	3.2341	3.3	0.2598	2.1	0.63	1489.0	27.8	1465.4	25.6	1431.4	48.7	1431.4	48.7	104.0
Trial-2_0957	92	229084	2.2	13.2560	3.4	1.9069	4.3	0.1833	2.7	0.62	1085.2	26.7	1083.5	28.6	1080.2	67.5	1080.2	67.5	100.5
Trial-2_0958	20	18472	1.2	12.7027	7.1	2.1299	9.1	0.1962	5.7	0.62	1155.0	59.8	1158.6	62.9	1165.2	141.1	1165.2	141.1	99.1
Trial-2_0959	98	844266	0.6	5.4111	1.8	13.2051	2.4	0.5182	1.6	0.66	2691.7	35.0	2694.4	22.6	2696.5	29.6	2696.5	29.6	99.8
Trial-2_0960	79	152061	1.1	12.8080	3.6	2.0118	4.6	0.1869	2.9	0.62	1104.5	29.1	1119.5	31.2	1148.8	71.5	1148.8	71.5	96.1
Trial-2_0961	115	60955	2.3	9.2935	2.3	4.2200	3.0	0.2844	1.9	0.64	1613.7	27.3	1677.9	24.5	1759.2	41.8	1759.2	41.8	91.7
Trial-2_0962	39	267096	1.1	9.9514	4.3	3.7048	5.5	0.2674	3.5	0.63	1527.5	47.4	1572.4	44.0	1633.1	79.2	1633.1	79.2	93.5

Table _____. U-Pb geochronologic analyses.																									
Trial 3: CP-40 Element2 TC							Isotope ratios					Apparent ages (Ma)													
Analysis	U	206Pb	U/Th	206Pb*	±	207Pb*	±	206Pb*	±	error	206Pb*	±1s(a)	±1s(b)	±1s(c)	207Pb*	±1s(a)	±1s(b)	±1s(c)	206Pb*	±1s(a)	±1s(b)	±1s(c)	Best age	±	Conc
	(ppm)	204Pb		207Pb*	(%)	235U*	(%)	238U	(%)	corr.	238U*	(Ma)	(Ma)	(Ma)	235U	(Ma)	(Ma)	(Ma)	207Pb*	(Ma)	(Ma)	(Ma)	(Ma)	(Ma)	(%)
Trial3_001	488	NA()	4.1	16.9904	1.1	0.7355	1.7	0.0907	1.3	0.77	559.5	0.6	6.3	6.7	559.8	1.5	6.7	7.1	561.8	7.3	22.3	23.8	559.5	6.3	99.6
Trial3_002	100	NA()	1.8	10.5551	1.2	3.3716	2.2	0.2582	1.8	0.85	1480.7	8.8	17.8	18.7	1497.8	6.2	13.4	14.1	1522.9	8.0	19.9	21.2	1522.9	19.9	97.2
Trial3_003	62	NA()	2.9	8.0608	1.1	6.0005	2.2	0.3510	1.9	0.86	1939.2	10.7	22.2	23.4	1975.9	6.9	14.9	15.7	2015.4	8.5	19.1	20.3	2015.4	19.1	96.2
Trial3_004	196	NA()	1.6	6.7212	0.9	8.5365	1.2	0.4163	0.9	0.72	2243.7	7.4	23.3	24.7	2289.8	4.5	14.5	15.4	2332.0	5.3	17.4	18.6	2332.0	17.4	96.2
Trial3_005	103	NA()	1.9	13.1856	1.5	1.9212	2.5	0.1838	2.0	0.81	1087.8	7.4	13.8	14.5	1088.5	5.9	11.7	12.4	1090.8	10.0	21.8	23.1	1090.8	21.8	99.7
Trial3_006	239	NA()	1.6	9.7998	0.8	4.0712	1.3	0.2895	1.0	0.76	1639.0	6.1	17.9	19.0	1648.5	4.0	13.0	13.8	1661.6	4.9	18.5	19.8	1661.6	18.5	98.6
Trial3_007	25	NA()	1.7	13.0485	2.9	1.9102	3.8	0.1809	2.5	0.66	1071.7	9.2	14.7	15.4	1084.7	9.8	14.1	14.6	1111.7	22.9	30.0	30.9	1111.7	30.0	96.4
Trial3_008	188	NA()	3.3	12.5037	1.1	2.1418	1.9	0.1943	1.6	0.83	1144.7	6.0	13.6	14.4	1162.4	4.6	11.5	12.1	1196.4	6.9	20.3	21.6	1196.4	20.3	95.7
Trial3_009	171	NA()	1.8	11.0841	1.0	2.8645	1.8	0.2304	1.5	0.84	1336.5	6.9	15.7	16.5	1372.6	5.0	12.4	13.2	1430.2	6.4	19.5	20.8	1430.2	19.5	93.4
Trial3_010	194	NA()	1.7	11.3421	0.9	2.9692	1.6	0.2444	1.4	0.83	1409.3	6.6	16.2	17.1	1399.8	4.7	12.4	13.2	1386.1	6.2	19.6	20.9	1386.1	19.6	101.7
Trial3_011	45	NA()	0.9	16.6251	3.3	0.7842	4.1	0.0946	2.5	0.61	582.7	5.4	8.4	8.8	587.9	7.3	10.0	10.3	609.0	28.6	35.4	36.3	582.7	8.4	95.7
Trial3_012	77	NA()	0.6	10.7064	1.3	3.1594	2.4	0.2454	2.0	0.83	1414.9	9.1	17.4	18.3	1447.3	6.7	13.4	14.2	1496.0	9.1	20.4	21.7	1496.0	20.4	94.6
Trial3_013	146	NA()	1.4	17.4639	2.2	0.5495	3.2	0.0696	2.3	0.71	434.0	3.2	5.8	6.1	444.7	4.0	6.7	7.1	501.6	17.5	27.5	28.8	434.0	5.8	86.5
Trial3_014	53	NA()	1.1	8.9657	1.3	5.0662	2.4	0.3296	2.1	0.84	1836.3	11.1	21.7	22.8	1830.5	7.4	14.8	15.6	1824.6	9.5	19.9	21.1	1824.6	19.9	100.6
Trial3_015	70	NA()	1.9	11.6123	1.5	2.6648	2.6	0.2245	2.1	0.81	1305.8	9.2	16.5	17.3	1318.7	7.0	13.2	13.9	1340.8	10.5	21.4	22.6	1340.8	21.4	97.4
Trial3_016	82	NA()	2.6	8.8628	1.1	4.8689	2.2	0.3131	1.9	0.86	1756.0	10.2	20.6	21.7	1796.9	6.7	14.4	15.2	1845.5	8.0	19.2	20.4	1845.5	19.2	95.1
Trial3_017	211	NA()	2.8	13.7280	1.1	1.6922	2.0	0.1686	1.6	0.82	1004.2	5.3	12.0	12.7	1005.6	4.3	10.6	11.2	1009.6	7.5	21.0	22.3	1009.6	21.0	99.5
Trial3_018	237	NA()	3.3	11.7308	0.9	2.4277	1.6	0.2066	1.4	0.83	1210.9	5.5	14.0	14.8	1250.8	4.2	11.7	12.4	1321.1	5.8	19.6	21.0	1321.1	19.6	91.7
Trial3_019	28	NA()	1.3	13.2984	2.8	1.8750	3.8	0.1809	2.5	0.67	1072.0	9.3	14.8	15.4	1072.3	9.6	13.9	14.4	1073.8	22.3	29.6	30.6	1073.8	29.6	99.8
Trial3_020	61	NA()	0.8	9.9187	1.3	4.1543	2.4	0.2990	2.0	0.84	1686.3	10.4	20.2	21.2	1665.1	7.1	14.3	15.0	1639.2	9.5	20.3	21.5	1639.2	20.3	102.9
Trial3_021	102	NA()	0.9	13.5359	1.6	1.7495	2.6	0.1718	2.1	0.80	1022.2	7.1	13.1	13.7	1027.0	6.0	11.4	12.1	1038.1	10.8	22.3	23.6	1038.1	22.3	98.5
Trial3_022	72	NA()	2.0	13.6706	1.8	1.8083	2.9	0.1794	2.2	0.78	1063.5	7.9	13.9	14.5	1048.5	6.6	11.9	12.5	1018.1	12.3	23.1	24.3	1018.1	23.1	104.5
Trial3_023	260	NA()	0.9	9.0448	0.8	4.7174	1.2	0.3096	0.9	0.73	1738.7	5.8	18.7	19.8	1770.3	3.8	13.2	14.1	1808.6	4.5	18.1	19.4	1808.6	18.1	96.1
Trial3_024	233	NA()	3.2	9.3311	0.8	4.5441	1.3	0.3077	1.0	0.77	1729.2	6.4	18.8	19.9	1739.1	4.2	13.3	14.1	1751.8	4.9	18.4	19.6	1751.8	18.4	98.7
Trial3_025	58	NA()	1.8	11.4939	1.7	2.6972	2.8	0.2249	2.2	0.80	1308.0	9.3	16.7	17.5	1327.7	7.3	13.4	14.1	1360.5	11.4	21.8	23.0	1360.5	21.8	96.1
Trial3_026	64	NA()	0.8	13.6806	1.9	1.7545	3.0	0.1742	2.3	0.76	1035.0	8.0	13.7	14.3	1028.8	7.0	12.0	12.6	1016.6	13.9	24.0	25.2	1016.6	24.0	101.8
Trial3_027	67	NA()	1.3	5.4046	0.9	12.9904	1.9	0.5094	1.6	0.87	2654.2	10.4	27.4	29.0	2679.0	6.4	15.6	16.6	2698.4	8.0	17.9	18.9	2698.4	17.9	98.4
Trial3_028	112	NA()	2.3	13.7848	1.5	1.7615	2.5	0.1762	2.0	0.81	1046.1	7.0	13.2	13.9	1031.4	5.7	11.3	12.0	1001.2	10.2	22.1	23.4	1001.2	22.1	104.5
Trial3_029	462	NA()	1.9	6.1296	0.8	8.4889	1.3	0.3776	1.0	0.78	2064.8	1.7	20.6	22.1	2284.7	1.4	13.8	14.8	2488.5	1.9	16.4	17.6	2488.5	16.4	83.0
Trial3_030	256	NA()	3.7	13.6065	1.0	1.7195	1.8	0.1698	1.4	0.82	1010.8	4.5	11.8	12.5	1015.8	3.7	10.4	11.1	1027.6	6.6	20.6	22.0	1027.6	20.6	98.4
Trial3_031	120	NA()	1.2	11.0658	1.1	3.0860	2.1	0.2478	1.8	0.84	1427.1	8.4	17.1	18.0	1429.2	5.9	13.0	13.8	1433.3	7.7	20.0	21.3	1433.3	20.0	99.6
Trial3_032	664	NA()	5.1	5.1636	0.8	12.7754	1.3	0.4787	1.0	0.78	2521.4	3.6	24.6	26.2	2663.2	2.2	14.4	15.4	2773.5	2.6	16.0	17.2	2773.5	16.0	90.9
Trial3_033	116	NA()	2.2	12.1508	1.4	1.9657	2.4	0.1733	2.0	0.83	1030.3	7.0	13.1	13.8	1103.9	5.8	11.7	12.4	1252.6	9.0	20.9	22.2	1252.6	20.9	82.3
Trial3_034	61	NA()	0.8	10.6679	1.5	3.3046	2.6	0.2558	2.1	0.82	1468.3	10.3	18.4	19.3	1482.1	7.4	13.9	14.7	1502.8	10.2	20.9	22.1	1502.8	20.9	97.7
Trial3_035	148	NA()	0.7	18.3665	2.3	0.5051	3.2	0.0673	2.3	0.71	419.9	3.3	5.8	6.0	415.1	3.8	6.4	6.7	389.6	17.1	27.6	28.9	419.9	5.8	107.8
Trial3_036	31	NA()	1.6	13.7474	2.8	1.6626	3.7	0.1658	2.5	0.66	989.2	8.6	13.7	14.3	994.4	9.5	13.5	14.0	1006.7	23.7	30.7	31.7	1006.7	30.7	98.3
Trial3_037	86	NA()	0.8	13.8501	1.8	1.5588	2.8	0.1567	2.2	0.78	938.2	7.2	12.5	13.0	954.0	6.4	11.3	11.9	991.6	12.6	23.3	24.6	991.6	23.3	94.6
Trial3_038	83	NA()	0.4	14.3410	4.1	0.1222	5.2	0.0127	3.2	0.61	81.5	0.9	1.3	1.4	117.1	3.4	3.8	3.9	920.4	58.8	62.1	62.6	81.5	1.3	NA
Trial3_039	310	NA()	3.5	13.3424	0.9	1.7974	1.5	0.1740	1.2	0.80	1034.2	3.8	11.7	12.5	1044.5	3.2	10.4	11.0	1067.1	5.8	20.2	21.7	1067.1	20.2	96.9
Trial3_040	100	NA()	1.5	9.2163	1.1	4.3262	2.1	0.2893	1.8	0.86	1638.1	9.6	19.4	20.4	1698.4	6.4	14.0	14.8	1774.4	7.5	19.2	20.4	1774.4	19.2	92.3
Trial3_041	96	NA()	1.8	12.7002	1.4	2.1247	2.5	0.1958	2.0	0.82	1152.7	8.0	14.7	15.4	1156.9	6.3	12.2	12.8	1165.6	9.8	21.5	22.8	1165.6	21.5	98.9
Trial3_042	516	NA()	1.3	12.4571	0.8	1.3189	1.3	0.1192	1.0	0.78	726.0	0.6	8.0	8.6	854.0	1.5	8.9	9.5	1203.7	4.8	19.6	21.0	1203.7	19.6	60.3
Trial3_043	74	NA()	0.8	9.4346	1.2	4.6932	2.2	0.3213	1.9	0.85	1796.0	10.5	21.1	22.2	1766.0	6.9	14.4	15.2	1731.6	8.6	19.7	20.9	1731.6	19.7	103.7
Trial3_044	218	NA()	1.5	18.5970	1.9	0.5294	2.8	0.0714	2.1	0.75	444.8	2.8	5.8	6.1	431.4	3.2	6.2	6.5	361.5	14.1	25.9	27.4	444.8	5.8	123.1
Trial3_045	66	NA()	1.9	12.7391	1.7	2.2048	2.8	0.2038	2.2	0.79	1195.7	9.1	15.6	16.3	1182.6	7.2	12.8	13.4	1159.5	12.1	22.7	23.9	1159.5	22.7	103.1
Trial3_046	62	NA()	0.4	5.3380	1.0	13.1658	1.9	0.5099	1.7	0.87	2656.4	11.0	27.6	29.3	2691.6	6.9	15.8	16.8	2718.9	8.6	18.1	19.2	2718.9	18.1	97.7
Trial3_047	110	NA()	1.9	13.3970	1.5	1.8296	2.5	0.1779	2.0	0.81	1055.2	7.5	13.6	14.3	1056.1	6.1	11.6	12.3	1058.9	10.2	22.0	23.3	1058.9	22.0	99.6
Trial3_048	101	NA()	0.5	10.2480	1.1	3.6658	2.1	0.2726	1.8	0.85	1553.9	9.6	18.7	19.7	1564.0	6.5	13.7	14.5	1578.4	7.9	19.7	21.0	1578.4	19.7	98.5
Trial3_049	227	NA()	2.4	17.8513	1.8	0.5146	2.8	0.0667	2.1	0.75	416.0	2.6	5.4	5.7	421.5	3.1	6.1	6.4	453.1	14.0	25.6	27.0	416.0	5.4	91.8
Trial3_050	96	NA()	2.8	6.5335	0.9	9.0418	1.8	0.4286	1.5	0.86	2299.6	10.6	24.9	26.3	2342.2	6.4	15.2	16.2	2380.3	7.5	18.1	19.2	2380.3	18.1	96.6
Trial3_051	110	NA()	1.1	5.5409	0.9	11.9992	1.6	0.4824	1.3	0.81	2537.8	10.0	26.4	28.0	2604.3	6.1	15.4	16.4	2657.2	7.3	17.6	18.7	2657.2	17.6	95.5
Trial3_052	22	NA()	2.3	5.3566	1.4	12.4502	2.6	0.4839	2.2	0.85	2544.2	13.4	27.9	29.4	2639.0	8.5	16.6	17.5	2713.2	10.7	19.2	20.2	2713.2	19.2	93.8
Trial3_053	104	NA()	2.6	13.5645	1.5	1.8196	2.6	0.1791	2.1	0.81	1062.0	7.8	13.8	14.5	1052.5	6.3	11.7	12.3	1033.8	10.6	22.2	23.5	1033.8	22.2	102.7
Trial3_054	80	NA()	1.3	5.3610	0.9	12.8639	1.8	0.5004	1.5	0.85	2615.5	10.8	27.2	28.8	2669.7	6.6	15.7	16.7	2711.8	8.1	17.9	19.0	2711.8	17.9	96.4
Trial3_055	48	NA()	0.8	11.5729	1.9	2.5326	3.0	0.2127	2.3	0.78	1243.0	9.9	16.5	17.2	1281.5	8.0	13.6	14.3	1347.3	12.8	22.6	23.8	1347.3	22.6	92.3
Trial3_056	190	NA()	2.5	18.6917	2.3	0.4056	3.2	0.0550	2.3	0.71	345.2	2.5	4.7	4.9	345.7	3.3	5.5	5.8	350.0	19.2	29.1	30.3	345.2	4.7	NA
Trial3_057	112	NA()	0.8	15.7718	2.4	0.6211	3.4	0.0711	2.4	0.71	442.6	3.9	6.3	6.6	490.5	4.9	7.6	8.0	721.8	18.5	27.6	28.8	442.6	6.3	61.3
Trial3_058	38	NA()	2.3	12.4582	2.2	2.1948	3.3	0.1984	2.4	0.73	1166.7	9.8	15.8	16.5	1179.4	8.9	13.8	14.4	1203.6	17.3	25.7	26.8	1203.6	25.7	96.9

Trial3_059	75	NA()	1.7	17.1329	2.8	0.7016	3.7	0.0872	2.5	0.66	539.1	5.0	7.8	8.1	539.7	6.2	8.9	9.2	543.6	24.6	32.4	33.5	539.1	7.8	99.2
Trial3_060	35	NA()	0.7	10.2979	1.9	3.2977	3.0	0.2464	2.4	0.78	1419.9	11.0	18.5	19.3	1480.5	8.6	14.6	15.3	1569.3	12.9	22.2	23.4	1569.3	22.2	90.5
Trial3_061	28	NA()	0.8	10.7080	2.1	3.2525	3.2	0.2527	2.4	0.75	1452.4	11.6	19.1	19.9	1469.8	9.3	15.0	15.7	1495.8	15.2	23.8	24.9	1495.8	23.8	97.1
Trial3_062	20	NA()	1.6	9.6992	2.3	4.0454	3.3	0.2847	2.5	0.74	1615.0	12.7	20.9	21.8	1643.4	10.2	16.0	16.7	1680.7	16.2	24.1	25.1	1680.7	24.1	96.1
Trial3_063	15	NA()	1.0	7.5927	2.0	6.9453	3.1	0.3826	2.4	0.77	2088.6	14.2	25.1	26.3	2104.5	10.0	16.7	17.5	2120.8	13.9	21.9	22.9	2120.8	21.9	98.5
Trial3_064	125	NA()	1.2	10.7230	1.1	3.3890	2.0	0.2637	1.7	0.85	1508.7	9.2	18.1	19.1	1501.9	6.2	13.4	14.1	1493.1	7.6	19.8	21.1	1493.1	19.8	101.0
Trial3_065	78	NA()	2.0	12.8581	1.7	1.8493	2.8	0.1725	2.2	0.79	1026.1	8.2	13.7	14.4	1063.2	6.9	12.1	12.7	1141.1	11.9	22.6	23.8	1141.1	22.6	89.9
Trial3_066	196	NA()	2.8	13.8432	1.1	1.7282	2.0	0.1736	1.7	0.82	1031.9	6.4	12.8	13.5	1019.1	5.0	10.9	11.6	992.6	7.8	21.1	22.5	992.6	21.1	104.0
Trial3_067	92	NA()	2.5	11.3190	1.3	2.7423	2.4	0.2252	2.0	0.84	1309.4	9.4	16.7	17.5	1340.0	6.9	13.2	13.9	1390.0	9.2	20.7	21.9	1390.0	20.7	94.2
Trial3_068	220	NA()	3.3	12.7889	1.0	1.9647	1.8	0.1823	1.5	0.83	1079.6	6.0	13.0	13.8	1103.5	4.7	11.2	11.9	1151.8	6.7	20.3	21.7	1151.8	20.3	93.7
Trial3_069	115	NA()	1.9	13.2353	1.5	1.7194	2.5	0.1651	2.1	0.81	985.2	7.5	13.0	13.6	1015.8	6.2	11.5	12.1	1083.3	10.2	21.9	23.2	1083.3	21.9	90.9
Trial3_070	79	NA()	1.7	5.4380	1.0	12.9458	1.8	0.5108	1.5	0.84	2660.1	11.0	27.7	29.3	2675.7	6.9	15.8	16.8	2688.3	8.7	18.2	19.2	2688.3	18.2	99.0
Trial3_071	89	NA()	2.0	9.1263	1.1	4.9215	2.1	0.3259	1.8	0.86	1818.5	11.0	21.5	22.6	1805.9	7.0	14.6	15.4	1792.3	8.3	19.4	20.6	1792.3	19.4	101.5
Trial3_072	125	NA()	2.9	12.8630	1.3	2.0656	2.3	0.1928	1.9	0.83	1136.5	8.1	14.6	15.3	1137.5	6.2	12.0	12.7	1140.3	9.0	21.2	22.5	1140.3	21.2	99.7
Trial3_073	334	NA()	0.7	17.1125	1.4	0.6873	2.2	0.0853	1.7	0.78	527.9	2.4	6.4	6.7	531.2	2.7	6.8	7.2	546.2	9.8	23.3	24.7	527.9	6.4	96.7
Trial3_074	130	NA()	7.8	13.1301	1.3	1.9048	2.3	0.1815	1.9	0.83	1075.0	7.7	13.9	14.5	1082.8	6.0	11.7	12.3	1099.3	8.9	21.3	22.6	1099.3	21.3	97.8
Trial3_075	221	NA()	2.3	12.4404	1.0	1.8641	1.9	0.1683	1.6	0.83	1002.5	5.7	12.2	12.9	1068.4	4.6	11.0	11.7	1206.4	6.7	20.2	21.5	1206.4	20.2	83.1
Trial3_076	25	NA()	1.1	5.6411	1.3	12.3680	2.5	0.5062	2.2	0.86	2640.6	13.1	28.4	30.0	2632.7	8.4	16.5	17.4	2627.5	10.9	19.4	20.4	2627.5	19.4	100.5
Trial3_077	123	NA()	1.4	9.2577	0.9	4.6072	1.8	0.3095	1.6	0.86	1738.2	9.9	20.3	21.4	1750.6	6.3	14.1	14.9	1766.2	7.2	19.1	20.3	1766.2	19.1	98.4
Trial3_078	96	NA()	2.3	13.7009	1.6	1.7254	2.6	0.1715	2.1	0.80	1020.5	7.7	13.4	14.1	1018.0	6.3	11.6	12.2	1013.6	11.0	22.5	23.8	1013.6	22.5	100.7
Trial3_079	128	NA()	1.6	12.9803	1.4	1.8344	2.4	0.1728	2.0	0.82	1027.3	7.2	13.2	13.8	1057.8	5.8	11.5	12.1	1122.2	9.2	21.3	22.6	1122.2	21.3	91.5
Trial3_080	168	NA()	1.3	5.2838	0.9	13.6724	1.2	0.5242	0.8	0.67	2716.9	8.3	27.1	28.8	2727.3	5.2	15.2	16.2	2735.7	6.6	17.2	18.3	2735.7	17.2	99.3
Trial3_081	24	NA()	0.3	5.2961	1.3	13.2207	2.5	0.5080	2.2	0.86	2648.3	13.0	28.4	30.0	2695.5	8.3	16.5	17.4	2731.9	10.6	19.1	20.1	2731.9	19.1	96.9
Trial3_082	109	NA()	0.7	13.3009	1.5	1.8039	2.5	0.1741	2.1	0.81	1034.6	7.4	13.4	14.0	1046.9	6.0	11.6	12.2	1073.4	10.0	21.9	23.2	1073.4	21.9	96.4
Trial3_083	118	NA()	1.1	8.9805	0.9	4.9438	1.8	0.3221	1.6	0.86	1800.2	9.7	20.7	21.8	1809.8	6.1	14.2	15.0	1821.6	7.1	18.9	20.1	1821.6	18.9	98.8
Trial3_084	70	NA()	1.0	10.9224	1.4	3.0280	2.5	0.2400	2.1	0.83	1386.6	9.7	17.5	18.3	1414.7	7.1	13.6	14.3	1458.1	9.7	20.8	22.0	1458.1	20.8	95.1
Trial3_085	45	NA()	1.6	13.5317	2.2	1.8495	3.2	0.1816	2.4	0.74	1075.7	9.0	14.6	15.2	1063.3	7.9	12.7	13.3	1038.7	15.9	25.2	26.4	1038.7	25.2	103.6
Trial3_086	278	NA()	1.7	5.1124	0.8	13.9690	1.3	0.5182	1.0	0.77	2691.5	5.8	26.3	28.0	2747.6	3.6	14.8	15.8	2789.9	4.4	16.4	17.6	2789.9	16.4	96.5
Trial3_087	133	NA()	1.3	9.9443	1.0	3.9156	1.9	0.2825	1.6	0.86	1604.1	8.9	18.8	19.8	1616.9	5.9	13.6	14.4	1634.4	6.9	19.2	20.5	1634.4	19.2	98.1
Trial3_088	388	NA()	30.9	13.7708	0.8	1.7381	1.3	0.1737	1.0	0.77	1032.3	2.6	11.4	12.2	1022.8	2.4	10.0	10.7	1003.3	5.1	20.2	21.7	1003.3	20.2	102.9
Trial3_089	52	NA()	1.4	5.3373	1.0	13.0616	2.0	0.5058	1.8	0.88	2638.9	11.4	27.7	29.3	2684.1	7.1	15.9	16.9	2719.1	8.8	18.2	19.3	2719.1	18.2	97.0
Trial3_090	187	NA()	2.2	10.9442	0.9	3.1578	1.7	0.2508	1.4	0.84	1442.4	7.1	16.6	17.6	1446.9	4.9	12.7	13.4	1454.4	6.1	19.4	20.7	1454.4	19.4	99.2
Trial3_091	163	NA()	2.3	10.2727	0.9	3.6346	1.7	0.2709	1.5	0.85	1545.5	7.9	17.8	18.8	1557.1	5.3	13.2	14.0	1573.9	6.3	19.2	20.4	1573.9	19.2	98.2
Trial3_092	156	NA()	2.3	13.5262	1.2	1.8331	2.2	0.1799	1.8	0.82	1066.5	6.5	13.2	13.9	1057.4	5.1	11.2	11.8	1039.6	8.5	21.3	22.6	1039.6	21.3	102.6
Trial3_093	104	NA()	1.2	11.5069	1.2	2.7775	2.3	0.2319	1.9	0.84	1344.5	8.6	16.5	17.4	1349.5	6.2	12.9	13.6	1358.4	8.5	20.5	21.8	1358.4	20.5	99.0
Trial3_094	232	NA()	1.3	18.9449	2.5	0.2812	3.4	0.0387	2.3	0.68	244.5	1.5	3.2	3.3	251.6	2.5	4.2	4.4	319.6	20.7	30.2	31.4	244.5	3.2	NA
Trial3_095	151	NA()	1.3	10.9783	1.0	3.1217	1.9	0.2487	1.6	0.85	1431.6	7.6	16.8	17.7	1438.1	5.3	12.8	13.6	1448.4	6.9	19.7	21.0	1448.4	19.7	98.8
Trial3_096	63	NA()	0.5	16.5135	3.8	0.3039	4.9	0.0364	3.2	0.64	230.6	2.4	3.5	3.7	269.5	5.9	6.9	7.1	623.5	49.1	53.4	54.0	230.6	3.5	NA
Trial3_097	179	NA()	1.6	18.2610	2.1	0.5090	3.1	0.0674	2.2	0.72	420.7	2.8	5.5	5.8	417.8	3.6	6.3	6.6	402.5	17.4	27.8	29.1	420.7	5.5	104.5
Trial3_098	151	NA()	1.0	18.5793	2.4	0.4705	3.3	0.0634	2.3	0.69	396.4	2.9	5.4	5.6	391.5	3.8	6.2	6.5	363.6	20.2	29.7	30.9	396.4	5.4	NA
Trial3_099	162	NA()	2.0	5.3801	0.9	12.8326	1.2	0.5010	0.9	0.70	2617.9	8.0	26.3	28.0	2667.4	4.9	15.1	16.0	2705.9	5.9	17.0	18.1	2705.9	17.0	96.7
Trial3_100	100	NA()	1.8	17.7851	2.8	0.5093	3.7	0.0657	2.4	0.66	410.3	3.5	5.8	6.0	418.0	4.7	7.0	7.3	461.3	23.9	32.1	33.2	410.3	5.8	88.9
Trial3_101	89	NA()	1.6	12.0151	1.5	2.2653	2.6	0.1975	2.1	0.82	1161.8	7.9	14.7	15.4	1201.5	6.4	12.4	13.1	1274.6	10.1	21.4	22.6	1274.6	21.4	91.2
Trial3_102	25	NA()	0.9	13.4668	3.4	1.3211	4.6	0.1291	3.2	0.68	782.7	7.4	11.4	11.8	855.0	9.5	12.9	13.4	1048.5	26.2	32.6	33.5	1048.5	32.6	74.7
Trial3_103	101	NA()	5.0	17.8980	2.7	0.5370	3.6	0.0697	2.4	0.67	434.6	3.6	6.1	6.4	436.4	4.8	7.2	7.5	447.3	22.7	31.3	32.4	434.6	6.1	97.2
Trial3_104	394	NA()	2.0	9.1710	0.8	4.7202	1.3	0.3141	1.0	0.78	1760.9	2.8	18.2	19.4	1770.8	2.1	12.8	13.7	1783.4	3.1	17.9	19.2	1783.4	17.9	98.7
Trial3_105	52	NA()	1.9	12.8916	2.0	1.9904	3.1	0.1862	2.3	0.76	1100.7	8.3	14.4	15.1	1112.3	7.4	12.6	13.2	1135.9	14.3	24.0	25.1	1135.9	24.0	96.9
Trial3_106	84	NA()	0.7	18.0286	3.0	0.5273	3.8	0.0690	2.5	0.64	430.0	3.7	6.1	6.4	430.0	5.0	7.3	7.6	431.1	25.2	33.1	34.2	430.0	6.1	99.7
Trial3_107	149	NA()	1.7	13.9990	1.3	1.6345	2.3	0.1660	1.9	0.81	990.2	5.9	12.2	12.9	983.6	4.9	10.7	11.3	969.8	8.9	21.6	23.0	969.8	21.6	102.1
Trial3_108	32	NA()	1.2	19.4448	5.6	0.2096	6.5	0.0296	3.2	0.50	187.9	2.3	3.1	3.2	193.2	8.7	9.1	9.1	260.1	110.0	112.2	112.6	187.9	3.1	NA
Trial3_109	35	NA()	0.9	10.9861	1.9	3.3172	3.0	0.2644	2.3	0.78	1512.5	10.6	19.0	19.9	1485.1	8.2	14.4	15.1	1447.1	13.3	22.7	23.8	1447.1	22.7	104.5
Trial3_110	48	NA()	1.1	19.6639	4.1	0.2610	5.2	0.0372	3.2	0.62	235.7	2.6	3.7	3.9	235.5	6.2	7.0	7.1	234.3	63.6	67.4	68.0	235.7	3.7	NA
Trial3_111	58	NA()	1.3	12.2205	1.8	2.3096	2.9	0.2048	2.3	0.79	1201.0	9.0	15.6	16.3	1215.2	7.3	13.0	13.6	1241.4	12.4	22.6	23.8	1241.4	22.6	96.7
Trial3_112	57	NA()	1.7	13.7146	2.1	1.5387	3.2	0.1531	2.4	0.74	918.5	7.5	12.5	13.0	946.0	7.1	11.7	12.3	1011.6	15.3	24.8	26.0	1011.6	24.8	90.8
Trial3_113	250	NA()	1.9	11.1774	0.8	2.9236	1.4	0.2371	1.1	0.81	1371.6	5.5	15.4	16.3	1388.0	4.0	12.1	12.9	1414.1	5.2	19.2	20.6	1414.1	19.2	97.0
Trial3_114	99	NA()	3.3	13.2729	1.6	1.7274	2.6	0.1664	2.1	0.80	992.0	7.3	12.9	13.6	1018.8	6.2	11.5	12.1	1077.6	10.9	22.2	23.5	1077.6	22.2	92.1
Trial3_115	217	NA()	18.2	13.3942	1.1	1.8104	1.9	0.1759	1.6	0.82	1044.8	5.4	12.5	13.2	1049.2	4.3	10.8	11.4	1059.4	7.0	20.7	22.1	1059.4	20.7	98.6
Trial3_116	74	NA()	1.2	10.6963	1.4	3.3266	2.5	0.2582	2.0	0.83	1480.5	9.8	18.2	19.2	1487.3	7.0	13.7	14.5	1497.8	9.6	20.6	21.9	1497.8	20.6	98.8
Trial3_117	58	NA()	2.1	14.0323	2.1	1.5480	3.2	0.1576	2.4	0.74	943.5	7.8	12.8	13.4	949.7	7.4	11.9	12.5	965.0	16.6	25.8	26.9	965.0	25.8	97.8
Trial3_118	127	NA()	1.5	10.6012	1.1	3.2225	2.0	0.2479	1.7	0.85	1427.5	8.4	17.1	18.0	1462.6	5.9	13.1	13.9	1514.7	7.3	19.6	20.9	1514.7	19.6	94.2
Trial3_119	117	NA()	2.0	10.9736	1.1	3.0212	2.1	0.2406	1.8	0.84	1389.6	8.6	16.9	17.8	1413.0	6.1	13.1	13.8	1449.3	7.8	20.0	21.3	1449.3	20.0	95.9
Trial3_120	23	NA()	2.6	13.7573	3.3	1.6860	4.1	0.1683	2.6	0.62	1002.8	9.2	14.2	14.8	1003.3	11.1	14.7	15.2	1005.3	28.9	34.9	35.8	1005.3	34.9	99.8
Trial3_121	67	NA()	2.4	17.0566	3.0	0.6598	3.9	0.0817	2.5	0.63	506.0	4.7	7.4	7.7	514.5	6.6	9.0	9.3	553.3	28.4	35.4	36.4	506.0	7.4	91.5
Trial3_122	85	NA()	1.8	12.8528	1.6	1.9784	2.7	0.1845	2.2	0.80	1091.5	8.1	14.2	14.9	1108.2	6.6	12.2	12.8	1141.9	11.1	22.2	23.4	1141.9	22.2	95.6
Trial3_123	101	NA()	3.1	11.1053	1.2	2.9254	2.3	0.2357	1.9	0.84	1364.5	9.1	17.0	17.8	1388.5	6.5	13.2	13.9	1426.5	8.5	20.3	21.6	1426.5	20.3	95.6
Trial3_124	129	NA()	3.7	9.3001	0.9	4.6709	1.8	0.3152	1.5	0.86	1766.2	9.0	20.1	21.3	1762.0	5.8	13.9	14.8	1757.9	6.9	19.0	20.2	1757.9	19.0	100.5
Trial3_125	78	NA()	2.4	13.4671	1.8	1.6508	2.9	0.1613	2.2	0.78	964.0	7.5	12.8	13.4	989.8	6.6	11.6	12.2	1048.4	12.9	23.4	24.6	1048.4	23.4	92.0
Trial3_126	41	NA()	1.4	16.9073	3.5	0.6633	4.7	0.0814	3.2	0.68	504.3	5.0	7.6	7.9	516.6	8.1	10.1	10.4	572.5	37.0	42.6	43.4	504.3	7.6	88.1
Trial3_127	113	NA()	2.6	13.7336	1.5	1.6995	2.5	0.1694	2.0	0.80	1008.6	7.0	12.9	13.6	1008.3	5.8	11.3	11.9	1008.8	10.4	22.2	23.5	1008.8	22.2	100.0
Trial3_128	217	NA()	1.7	14.2277	1.2	1.4126	2.1	0.1458	1.7	0.81	877.5	4.7	10.6	11.2	894.3	4.1	9.9	10.5	936.7	7.9	21.3	22.7	936.7	21.3	93.7
Trial3_129	23	NA()	1.5	12.8283	3.0	2.0431	3.9	0.1902	2.5	0.65	1122.3	9.8	15.5	16.1	1130.0	10.6	14.8	15.3	1145.7	24.4	31.1	32.0	1145.7	31.1	98.0
Trial3_130	67	NA()	1.1	18.7095	3.8	0.2720	5.0	0.0369	3.2	0.64	233.7	2.4	3.6	3.7	244.3	5.7	6.6	6.7	347.9	54.5	58.7	59.3	233.7	3.6	NA
Trial3_131	328	NA()	116.3	13.3363	0.9	1.8369	1.4	0.1778	1.1	0.79	1054.7	3.4	11.8	12.6	1058.8	2.9	10.4	11.1	1068.1	5.5	20.2	21.6	1068.1	20.2	98.7
Trial3_132	60	NA()	1.0	10.1127	1.5	3.5313	2.6	0.2591	2.1	0.83	1485.3	10.1	18.5	19.4	1534.2	7.3	14.0	14.8	1603.2	9.8	20.5	21.7	1603.2	20.5	92.6
Trial3_133	195	NA()	1.0	13.1363	1.2	1.7308	2.1	0.1650	1.7	0.83	984.4	5.5	12.0	12.6	1020.1	4.6	10.8	11.4	1098.4	7.5	20.7	22.1	1098.4	20.7	89.6
Trial3_134	25	NA()	1.8	13.2059	3.0	1.8215	3.9	0.1745	2.5	0.64	1037.1	9.3	14.5	15.1	1053.2	10.0	14.1	14.6	1087.8	23.8	30.7	31.6	1087.8	30.7	95.3
Trial3_135	51	NA()	2.0	11.3952	1.8	2.7472	2.9	0.2271	2.3	0.79	1319.5	9.8	17.0	17.8	1341.3	7.7	13.7	14.3	1377.1	12.3	22.3	23.4	1377.1	22.3	95.8
Trial3_136	47	NA()	1.2	10.9638	1.7	3.2464	2.8	0.2583	2.2	0.80	1480.9	10.6	18.7	19.6	1468.3	7.9	14.2	14.9	1451.0	12.0	21.9	23.1	1451.0	21.9	102.1
Trial3_137	34	NA()	1.7	13.3545	2.6	1.7766	3.6	0.1722	2.5	0.69	1023.9	8.9	14.2	14.8	1036.9	9.1	13.4	13.9	1065.3	20.7	28.4	29.4	1065.3	28.4	96.1
Trial3_138	518	NA()	2.9	16.3713	1.2	0.5721	2.0	0.0680	1.5	0.78	423.8	0.5	4.8	5.1	459.3	1.5	5.8	6.2	642.2	8.1	22.3	23.8	423.8	4.8	66.0
Trial3_139	459	NA()	3.6	18.1924	2.6	0.1244	3.5	0.0164	2.3	0.65	105.0	0.3	1.2	1.3	119.0	1.1	2.0	2.1	410.9	20.9	30.1	31.3	105.0	1.2	NA
Trial3_140	390	NA()	1.0	19.1917	1.9	0.2830	2.8	0.0394	2.0	0.72	249.1	0.8	3.0	3.1	253.0	1.7	3.8	4.0	290.1	15.6	27.0	28.4	249.1	3.0	NA
Trial3_141	69	NA()	0.6	13.6497	1.9	1.6297	3.0	0.1614	2.3	0.77	964.6	7.7	13.0	13.5	981.7	6.8	11.7	12.3	1021.2	13.4	23.7	24.9	1021.2	23.7	94.5
Trial3_142	88	NA()	0.9	9.3841	1.1	4.5389	2.1	0.3091	1.8	0.86	1736.1	10.2	20.5	21.5	1738.1	6.7	14.3	15.1	1741.4	8.1	19.5	20.7	1741.4	19.5	99.7
Trial3_143	57	NA()	0.7	10.6527	1.5	3.2649	2.7	0.2524	2.2	0.82	1450.6	10.1	18.2	19.1	1472.7	7.4	13.9	14.6	1505.5	10.3	21.0	22.2	1505.5	21.0	96.4
Trial3_144	91	NA()	2.2	13.8106	1.7	1.6966	2.7	0.1700	2.2	0.79	1012.2	7.5	13.2	13.9	1007.2	6.2	11.5	12.1	997.4	11.4	22.7	24.0	997.4	22.7	101.5
Trial3_145	131	NA()	0.9	18.6936	3.2	0.2915	4.1	0.0395	2.5	0.61	250.0	2.1	3.6	3.7	259.7	3.6	5.0	5.2	349.8	29.8	37.0	38.0	250.0	3.6	NA
Trial3_146	243	NA()	2.2	18.2285	1.9	0.4856	2.8	0.0642	2.1	0.74	401.3	2.3	5.1	5.4	401.9	2.9	5.8	6.1	406.5	14.3	25.9	27.3	401.3	5.1	98.7
Trial3_147	53	NA()	3.3	11.7967	1.8	2.5229	2.9	0.2160	2.3	0.79	1260.5	9.5	16.4	17.1	1278.7	7.6	13.4	14.0	1310.2	12.3	22.4	23.6	1310.2	22.4	96.2
Trial3_148	67	NA()	0.8	14.1422	2.0	1.6062	3.0	0.1648	2.3	0.76	983.5	7.6	13.1	13.7	972.6	6.7	11.6	12.2	949.1	13.8	24.1	25.3	949.1	24.1	103.6
Trial3_149	32	NA()	0.8	5.4117	1.2	13.1948	2.4	0.5181	2.1	0.87	2691.2	12.0	28.3	29.9	2693.7	7.5	16.1	17.1	2696.3	9.6	18.6	19.7	2696.3	18.6	99.8
Trial3_150	581	NA()	1.1	11.1296	0.8	3.0516	1.3	0.2464	1.0	0.78	1420.1	1.8	14.9	16.0	1420.6	1.6	11.7	12.5	1422.3	3.1	18.7	20.1	1422.3	18.7	99.8
Trial3_151	124	NA()	1.7	17.1480	2.2	0.6789	3.2	0.0845	2.3	0.72	522.7	4.2	7.2	7.5	526.1	4.8	7.9	8.2	541.7	18.1	27.8	29.0	522.7	7.2	96.5
Trial3_152	31	NA()	2.0	13.2948	2.6	1.9031	3.6	0.1836	2.5	0.69	1086.5	9.4	15.0	15.6	1082.2	8.9	13.5	14.0	1074.3	19.2	27.3	28.3	1074.3	27.3	101.1
Trial3_153	189	NA()	1.5	11.2185	0.9	3.0477	1.7	0.2481	1.4	0.84	1428.6	7.2	16.6	17.5	1419.7	4.9	12.6	13.3	1407.1	6.2	19.5	20.8	1407.1	19.5	101.5
Trial3_154	143	NA()	2.1	13.2923	1.3	1.9109	2.3	0.1843	1.9	0.82	1090.4	7.0	13.6	14.3	1084.9	5.4	11.5	12.1	1074.7	8.6	21.2	22.6	1074.7	21.2	101.5
Trial3_155	51	NA()	0.7	19.4608	4.9	0.1984	5.9	0.0280	3.2	0.55	178.1	2.1	2.9	3.0	183.7	6.9	7.4	7.4	258.2	90.5	93.2	93.6	178.1	2.9	NA
Trial3_156	205	NA()	2.4	10.8939	0.9	3.2624	1.6	0.2579	1.3	0.84	1479.0	7.0	16.9	17.9	1472.1	4.8	12.7	13.5	1463.1	5.9	19.3	20.6	1463.1	19.3	101.1
Trial3_157	1298	NA()	6.2	18.8644	0.9	0.4136	2.1	0.0566	1.8	0.89	355.0	5.5	6.8	7.0	351.5	4.8	6.6	6.8	329.2	6.6	22.9	24.5	355.0	6.8	NA
Trial3_158	149	NA()	1.4	8.8202	0.9	5.2544	1.6	0.3363	1.4	0.84	1868.7	9.1	21.0	22.2	1861.5	5.7	14.1	14.9	1854.2	6.4	18.6	19.8	1854.2	18.6	100.8
Trial3_159	81	NA()	1.2	9.8962	1.2	4.2440	2.2	0.3047	1.9	0.85	1714.8	10.5	20.4	21.5	1682.6	6.9	14.2	15.0	1643.4	8.6	19.9	21.1	1643.4	19.9	104.3
Trial3_160	72	NA()	1.9	19.0336	3.4	0.4870	4.3	0.0673	2.5	0.59	419.6	4.0	6.2	6.5	402.9	5.9	7.8	8.0	308.9	33.7	40.2	41.1	419.6	6.2	135.8
Trial3_161	101	NA()	1.9	19.6712	3.8	0.2734	4.6	0.0390	2.5	0.55	246.8	2.3	3.7	3.8	245.4	5.6	6.5	6.6	233.4	55.3	59.6	60.3	246.8	3.7	NA
Trial3_162	55	NA()	1.0	14.0314	2.1	1.6512	3.2	0.1681	2.4	0.74	1001.7	8.3	13.6	14.2	990.0	7.3	12.1	12.6	965.2	15.2	24.9	26.1	965.2	24.9	103.8
Trial3_163	213	NA()	1.7	13.8490	1.1	1.7164	2.0	0.1725	1.6	0.82	1025.8	5.5	12.3	13.0	1014.7	4.4	10.6	11.3	991.8	7.4	21.0	22.4	991.8	21.0	103.4
Trial3_164	113	NA()	2.4	12.3946	1.3	2.3098	2.3	0.2077	1.9	0.83	1216.7	8.2	15.3	16.0	1215.3	6.1	12.4	13.0	1213.7	9.0	21.1	22.3	1213.7	21.1	100.3
Trial3_165	100	NA()	1.9	13.2425	1.5	1.9482	2.6	0.1872	2.1	0.81	1106.2	7.8	14.2	14.9	1097.8	6.3	11.9	12.6	1082.2	10.5	22.1	23.3	1082.2	22.1	102.2
Trial3_166	139	NA()	1.9	9.3975	0.9	4.6545	1.7	0.3174	1.5	0.86	1776.9	9.0	20.2	21.3	1759.1	5.7	13.9	14.7	1738.8	6.6	18.9	20.2	1738.8	18.9	102.2
Trial3_167	40	NA()	0.9	13.9411	2.5	1.6964	3.5	0.1716	2.5	0.70	1020.9	8.7	14.0	14.6	1007.2	8.5	12.9	13.4	978.3	19.7	27.8	28.9	978.3	27.8	104.4
Trial3_168	18	NA()	0.5	5.1430	1.4	13.6832	2.7	0.5106	2.3	0.85	2659.3	13.5	28.8	30.3	2728.0	8.6	16.7	17.6	2780.1	10.8	19.2	20.2	2780.1	19.2	95.7
Trial3_169	202	NA()	1.6	11.2414	0.9	3.0343	1.6	0.2475	1.3	0.83	1425.6	6.8	16.4	17.3	1416.3	4.7	12.5	13.3	1403.2	6.0	19.5	20.8	1403.2	19.5	101.6
Trial3_170	55	NA()	1.2	11.2716	1.6	3.0854	2.7	0.2523	2.2	0.81	1450.5	10.0	18.2	19.0	1429.1	7.4	13.8	14.5	1398.1	11.0	21.5	22.7	1398.1	21.5	103.8
Trial3_171	50	NA()	0.6	8.6394	1.3	5.4560	2.5	0.3420	2.1	0.85	1896.4	11.3	22.2	23.4	1893.7	7.4	15.0	15.8	1891.6	9.5	19.8	21.0	1891.6	19.8	100.3
Trial3_172	97	NA()	1.4	19.1491	3.4	0.2908	4.2	0.0404	2.5	0.60	255.3	2.3	3.7	3.9	259.2	4.1	5.4	5.5	295.2	34.6	41.1	42.0	255.3	3.7	NA
Trial3_173	233	NA()	1.4	11.2495	0.9	2.8012	1.5	0.2287	1.3	0.83	1327.4	5.8	15.1	16.0	1355.9	4.2	12.1	12.8	1401.8	5.5	19.3	20.7	1401.8	19.3	94.7
Trial3_174	138	NA()	1.1	19.6086	3.1	0.3031	3.9	0.0431	2.4	0.62	272.1	2.2	3.8	4.0	268.8	3.5	5.0	5.2	240.7	28.4	36.1	37.2	272.1	3.8	NA
Trial3_175	61	NA()	0.9	6.0695	0.9	10.8366	2.0	0.4772	1.7	0.88	2515.2	11.0	26.6	28.2	2509.2	6.8	15.6	16.5	2505.1	8.4	18.3	19.4	2505.1	18.3	100.4
Trial3_176	148	NA()	1.5	9.2889	0.9	4.5338	1.7	0.3056	1.4	0.86	1718.9	8.7	19.6	20.7	1737.2	5.6	13.8	14.6	1760.1	6.4	18.8	20.0	1760.1	18.8	97.7
Trial3_177	582	NA()	3.7	19.9095	1.6	0.2647	2.5	0.0382	1.8	0.74	241.9	0.4	2.8	3.0	238.4	1.2	3.4	3.6	205.5	12.0	25.4	26.9	241.9	2.8	NA
Trial3_178	56	NA()	0.8	9.9772	1.4	4.0241	2.5	0.2913	2.1	0.83	1648.1	10.6	20.0	21.0	1639.1	7.3	14.3	15.1	1628.3	9.9	20.5	21.7	1628.3	20.5	101.2
Trial3_179	143	NA()	2.5	10.7147	1.0	3.1265	1.9	0.2431	1.6	0.85	1402.7	7.8	16.6	17.5	1439.2	5.5	12.9	13.6	1494.6	6.9	19.5	20.8	1494.6	19.5	93.9
Trial3_180	24	NA()	0.7	7.4101	1.7	5.9890	2.9	0.3220	2.4	0.81	1799.6	12.3	22.0	23.1	1974.2	8.9	15.9	16.7	2163.4	11.5	20.4	21.4	2163.4	20.4	83.2
Trial3_181	155	NA()	1.0	9.6590	0.9	4.4244	1.6	0.3101	1.4	0.85	1741.1	8.3	19.6	20.7	1716.9	5.3	13.6	14.4	1688.3	6.3	18.9	20.2	1688.3	18.9	103.1
Trial3_182	178	NA()	1.2	17.8948	2.0	0.5569	3.0	0.0723	2.2	0.74	450.1	3.1	5.9	6.2	449.5	3.5	6.5	6.9	447.7	14.8	26.1	27.4	450.1	5.9	100.5
Trial3_183	439	NA()	0.1	18.9873	1.5	0.3915	2.4	0.0539	1.8	0.76	338.7	0.7	3.9	4.2	335.5	1.5	4.6	4.9	314.5	11.4	24.8	26.3	338.7	3.9	NA
Trial3_184	73	NA()	1.5	11.8806	1.5	2.6060	2.6	0.2247	2.1	0.81	1306.4	9.0	16.4	17.3	1302.3	6.8	13.0	13.7	1296.5	10.4	21.5	22.7	1296.5	21.5	100.8
Trial3_185	12	NA()	1.3	13.6260	3.6	1.7489	4.8	0.1729	3.2	0.67	1028.1	10.1	14.9	15.5	1026.7	14.1	17.2	17.6	1024.7	38.8	43.4	44.1	1024.7	43.4	100.3
Trial3_186	53	NA()	1.6	12.7929	1.9	2.1575	3.0	0.2003	2.3	0.77	1176.8	8.9	15.4	16.1	1167.5	7.5	12.9	13.5	1151.1	13.9	23.7	24.9	1151.1	23.7	102.2
Trial3_187	43	NA()	1.9	11.9709	1.9	2.6695	3.0	0.2319	2.3	0.77	1344.3	9.9	17.3	18.1	1320.1	7.8	13.6	14.3	1281.8	13.0	22.9	24.1	1281.8	22.9	104.9
Trial3_188	267	NA()	0.9	19.4249	2.1	0.3425	3.0	0.0483	2.1	0.71	303.9	1.6	3.8	4.0	299.0	2.3	4.6	4.8	262.4	16.8	27.8	29.2	303.9	3.8	NA
Trial3_189	341	NA()	8.8	19.3517	1.9	0.3130	2.8	0.0440	2.1	0.73	277.3	1.1	3.3	3.5	276.5	1.8	4.1	4.3	271.0	14.5	26.5	27.9	277.3	3.3	NA
Trial3_190	48	NA()	1.3	12.5857	2.1	2.0298	3.1	0.1854	2.4	0.75	1096.2	8.6	14.6	15.2	1125.5	7.6	12.8	13.4	1183.5	14.0	23.7	24.8	1183.5	23.7	92.6
Trial3_191	94	NA()	2.1	14.0257	1.7	1.6687	2.7	0.1698	2.1	0.79	1011.1	7.2	13.1	13.7	996.7	6.2	11.4	12.0	966.0	12.2	23.2	24.5	966.0	23.2	104.7
Trial3_192	252	NA()	1.6	19.7734	2.5	0.2534	3.4	0.0364	2.3	0.67	230.2	1.3	2.9	3.1	229.3	2.3	3.9	4.1	221.4	22.7	31.9	33.1	230.2	2.9	NA
Trial3_193	389	NA()	3.7	9.2324	0.8	4.8023	1.3	0.3217	1.0	0.78	1798.0	2.9	18.5	19.7	1785.3	2.1	12.9	13.8	1771.2	3.1	17.9	19.2	1771.2	17.9	101.5
Trial3_194	346	NA()	7.7	8.9924	0.8	4.7874	1.3	0.3124	1.0	0.78	1752.4	3.8	18.3	19.5	1782.7	2.6	13.0	13.9	1819.2	3.4	17.9	19.2	1819.2	17.9	96.3
Trial3_195	101	NA()	1.0	20.1452	3.4	0.2579	4.2	0.0377	2.5	0.60	238.5	2.1	3.5	3.6	233.0	3.8	4.9	5.1	178.1	36.5	42.9	43.9	238.5	3.5	NA
Trial3_196	56	NA()	1.6	13.3711	1.9	1.7715	3.7	0.1719	3.2	0.86	1022.4	8.6	17.7	18.2	1035.1	7.9	14.3	14.8	1062.8	16.1	23.5	24.7	1062.8	23.5	96.2

Trial3_197	144	NA()	1.6	8.6462	0.8	5.0863	2.1	0.3191	2.0	0.93	1785.3	9.2	27.2	28.1	1833.8	5.9	16.8	17.5	1890.1	6.6	16.7	18.0	1890.1	16.7	94.5
Trial3_198	106	NA()	1.1	18.0095	2.4	0.5375	4.1	0.0702	3.3	0.81	437.6	3.9	8.0	8.2	436.8	5.0	8.3	8.5	433.4	23.7	30.4	31.5	437.6	8.0	101.0
Trial3_199	91	NA()	0.8	17.0757	2.3	0.6642	4.0	0.0823	3.3	0.82	509.8	4.6	9.3	9.5	517.2	5.3	9.2	9.5	550.9	20.1	27.3	28.6	509.8	9.3	92.5
Trial3_200	129	NA()	1.2	17.1162	2.1	0.5957	3.8	0.0740	3.2	0.84	460.1	3.9	8.3	8.5	474.5	4.5	8.3	8.6	545.7	17.9	25.8	27.1	460.1	8.3	84.3
Trial3_201	60	NA()	1.0	19.9814	4.5	0.1865	5.5	0.0270	3.2	0.58	172.0	2.0	3.4	3.5	173.7	5.6	6.4	6.4	197.2	77.7	80.2	80.7	172.0	3.4	NA
Trial3_202	139	NA()	2.2	13.4109	1.2	1.7950	2.9	0.1747	2.6	0.91	1037.8	7.1	17.3	17.8	1043.6	5.7	13.3	13.9	1056.8	9.1	19.4	20.9	1056.8	19.4	98.2
Trial3_203	49	NA()	2.6	9.2731	1.3	4.4832	3.2	0.3016	3.0	0.92	1699.5	11.5	27.1	27.9	1727.9	7.9	17.3	17.9	1763.2	10.2	18.6	19.9	1763.2	18.6	96.4
Trial3_204	30	NA()	0.7	5.6135	1.1	11.9929	3.1	0.4885	2.9	0.93	2564.1	12.7	37.0	38.1	2603.9	8.0	19.1	19.9	2635.7	10.2	17.4	18.6	2635.7	17.4	97.3
Trial3_205	83	NA()	2.3	13.5189	1.6	1.6244	3.4	0.1593	3.0	0.88	953.1	7.7	16.5	16.9	979.7	6.8	13.5	14.0	1040.6	13.3	21.7	23.1	1040.6	21.7	91.6
Trial3_206	339	NA()	1.6	9.1493	0.8	4.0654	1.4	0.2699	1.1	0.79	1540.2	4.7	23.0	23.8	1647.4	3.3	15.4	16.2	1787.7	3.9	16.0	17.4	1787.7	16.0	86.2
Trial3_207	114	NA()	2.0	13.5029	1.3	1.7628	3.1	0.1727	2.8	0.90	1027.0	7.7	17.4	17.9	1031.9	6.2	13.5	14.0	1043.0	10.3	20.0	21.4	1043.0	20.0	98.5
Trial3_208	144	NA()	2.1	9.1836	0.8	4.7763	2.1	0.3183	2.0	0.93	1781.3	9.6	27.3	28.2	1780.7	6.1	16.7	17.4	1780.9	6.9	17.0	18.3	1780.9	17.0	100.0
Trial3_209	83	NA()	1.8	13.2601	1.5	1.8338	3.4	0.1764	3.0	0.89	1047.5	8.3	17.9	18.4	1057.6	6.8	13.9	14.5	1079.5	11.9	20.8	22.2	1079.5	20.8	97.0
Trial3_210	44	NA()	1.1	12.4672	1.9	2.0645	3.8	0.1868	3.3	0.86	1103.8	9.4	19.1	19.7	1137.1	8.2	15.1	15.6	1202.2	14.9	22.4	23.6	1202.2	22.4	91.8
Trial3_211	40	NA()	2.6	11.8592	1.9	2.2827	3.8	0.1964	3.3	0.86	1156.1	9.5	19.8	20.4	1206.9	8.3	15.5	16.0	1300.0	14.6	22.1	23.3	1300.0	22.1	88.9
Trial3_212	65	NA()	1.3	14.1225	1.9	1.5451	3.7	0.1583	3.2	0.86	947.5	7.8	16.4	16.9	948.5	7.1	13.4	13.9	951.9	15.1	23.0	24.3	951.9	23.0	99.5
Trial3_213	329	NA()	3.1	11.1476	0.8	2.9021	1.5	0.2347	1.3	0.84	1359.3	4.4	20.6	21.3	1382.5	3.3	14.3	15.0	1419.2	4.6	16.9	18.4	1419.2	16.9	95.8
Trial3_214	161	NA()	1.2	11.4941	1.0	2.5115	2.5	0.2095	2.3	0.92	1226.0	7.3	19.7	20.3	1275.4	5.4	14.5	15.1	1360.5	7.1	17.9	19.3	1360.5	17.9	90.1
Trial3_215	413	NA()	1.1	10.8444	0.8	2.9810	1.3	0.2346	1.0	0.76	1358.4	2.8	20.3	21.0	1402.8	2.3	14.3	14.9	1471.8	3.7	16.6	18.1	1471.8	16.6	92.3
Trial3_216	101	NA()	1.8	12.5429	1.3	2.2007	3.0	0.2003	2.8	0.91	1176.9	8.6	19.6	20.2	1181.3	6.5	14.5	15.0	1190.2	9.6	19.3	20.8	1190.2	19.3	98.9
Trial3_217	120	NA()	2.1	13.2425	1.3	1.8015	3.0	0.1731	2.8	0.91	1029.2	7.5	17.3	17.8	1046.0	6.1	13.5	14.0	1082.2	9.7	19.6	21.1	1082.2	19.6	95.1
Trial3_218	188	NA()	19.8	13.6231	1.1	1.7277	2.6	0.1708	2.4	0.91	1016.4	6.3	16.7	17.2	1018.9	5.0	12.9	13.4	1025.1	8.0	19.0	20.5	1025.1	19.0	99.2
Trial3_219	56	NA()	1.5	16.8344	2.8	0.7582	4.4	0.0926	3.4	0.78	571.0	5.6	10.6	10.9	573.0	6.8	10.5	10.9	581.9	25.0	31.1	32.1	571.0	10.6	98.1
Trial3_220	114	NA()	0.1	13.1719	1.4	1.5547	3.2	0.1486	2.9	0.90	893.1	6.9	15.3	15.8	952.4	6.1	12.9	13.4	1092.9	10.6	20.0	21.4	1092.9	20.0	81.7
Trial3_221	334	NA()	4.5	14.0550	0.9	1.5442	2.0	0.1575	1.8	0.90	942.8	3.7	14.9	15.4	948.2	3.2	11.8	12.4	961.7	6.1	18.4	20.0	961.7	18.4	98.0
Trial3_222	58	NA()	0.9	5.5602	0.9	11.7871	2.6	0.4755	2.5	0.94	2507.8	11.8	36.1	37.2	2587.6	7.2	18.8	19.5	2651.5	8.6	16.5	17.7	2651.5	16.5	94.6
Trial3_223	84	NA()	1.3	18.2108	2.8	0.5369	4.4	0.0709	3.4	0.77	441.8	4.2	8.2	8.4	436.4	6.2	9.0	9.3	408.6	32.5	37.6	38.6	441.8	8.2	108.1
Trial3_224	58	NA()	1.7	9.9066	1.3	4.0721	3.2	0.2927	2.9	0.92	1655.1	11.3	26.5	27.3	1648.7	7.7	16.9	17.6	1641.5	9.9	18.6	19.9	1641.5	18.6	100.8
Trial3_225	100	NA()	0.8	13.3878	1.4	1.8611	3.2	0.1808	2.9	0.90	1071.3	8.0	18.1	18.6	1067.4	6.4	13.8	14.3	1060.3	10.7	20.2	21.6	1060.3	20.2	101.0
Trial3_226	169	NA()	2.0	10.3977	0.8	3.5309	2.2	0.2664	2.0	0.92	1522.4	8.2	23.7	24.5	1534.2	5.5	15.6	16.3	1551.2	6.5	17.2	18.7	1551.2	17.2	98.1
Trial3_227	46	NA()	2.0	15.7059	2.7	0.9868	4.3	0.1125	3.4	0.79	687.0	6.5	12.5	12.9	697.1	7.6	12.0	12.4	730.7	23.9	29.9	31.0	687.0	12.5	94.0
Trial3_228	22	NA()	1.8	13.1328	2.9	1.9681	4.5	0.1875	3.5	0.77	1108.1	10.1	19.6	20.1	1104.7	10.8	16.5	17.0	1098.9	25.3	30.5	31.4	1098.9	30.5	100.8
Trial3_229	91	NA()	1.3	13.1753	1.4	1.8583	3.3	0.1777	2.9	0.90	1054.1	8.0	17.9	18.4	1066.4	6.6	13.9	14.4	1092.4	11.4	20.5	21.8	1092.4	20.5	96.5
Trial3_230	65	NA()	1.5	13.6174	1.8	1.6330	3.6	0.1614	3.2	0.87	964.3	8.0	16.8	17.2	983.0	7.1	13.6	14.1	1026.0	13.9	22.1	23.4	1026.0	22.1	94.0
Trial3_231	71	NA()	1.3	11.0012	1.3	3.0180	3.2	0.2409	2.9	0.91	1391.4	9.7	22.7	23.4	1412.2	7.1	15.8	16.4	1444.5	10.0	19.0	20.4	1444.5	19.0	96.3
Trial3_232	87	NA()	2.2	13.4763	1.5	1.6913	3.4	0.1654	3.0	0.89	986.6	7.5	16.8	17.3	1005.2	6.4	13.4	13.9	1047.0	11.6	20.7	22.1	1047.0	20.7	94.2
Trial3_233	210	NA()	1.6	12.9992	1.0	1.7665	2.5	0.1666	2.3	0.91	993.5	5.5	16.1	16.6	1033.2	4.5	12.8	13.4	1119.3	7.2	18.4	19.9	1119.3	18.4	88.8
Trial3_234	106	NA()	2.2	17.3933	2.7	0.4311	4.3	0.0544	3.4	0.79	341.5	3.1	6.3	6.4	364.0	4.5	7.2	7.4	510.5	24.8	31.1	32.2	341.5	6.3	NA
Trial3_235	99	NA()	0.8	10.3971	1.1	3.5194	2.8	0.2655	2.6	0.92	1517.9	9.4	24.1	24.8	1531.6	6.5	16.0	16.6	1551.3	8.1	17.9	19.3	1551.3	17.9	97.9
Trial3_236	157	NA()	1.9	13.5323	1.2	1.6421	2.8	0.1612	2.6	0.91	963.6	6.2	16.0	16.4	986.5	5.2	12.8	13.3	1038.6	8.7	19.3	20.8	1038.6	19.3	92.8
Trial3_237	108	NA()	1.3	16.8981	1.9	0.8334	3.7	0.1022	3.1	0.85	627.2	5.1	11.0	11.4	615.5	5.3	10.0	10.4	573.7	16.8	24.9	26.3	627.2	11.0	109.3
Trial3_238	134	NA()	2.3	13.4984	1.2	1.7938	2.9	0.1757	2.7	0.91	1043.4	6.9	17.3	17.8	1043.2	5.6	13.3	13.8	1043.7	9.5	19.6	21.1	1043.7	19.6	100.0
Trial3_239	108	NA()	1.1	11.0504	1.1	3.0199	2.8	0.2421	2.6	0.92	1397.8	8.8	22.4	23.1	1412.6	6.3	15.5	16.1	1436.0	8.3	18.2	19.6	1436.0	18.2	97.3
Trial3_240	240	NA()	1.0	19.9587	2.6	0.1950	4.2	0.0282	3.3	0.78	179.5	1.2	3.1	3.2	180.9	2.1	3.7	3.9	199.8	25.4	32.1	33.4	179.5	3.1	NA
Trial3_241	106	NA()	1.4	11.0265	1.1	3.1067	2.8	0.2486	2.6	0.92	1431.0	8.9	22.9	23.6	1434.4	6.4	15.6	16.2	1440.1	8.5	18.3	19.7	1440.1	18.3	99.4
Trial3_242	133	NA()	0.6	17.6898	2.2	0.5496	3.9	0.0706	3.2	0.83	439.5	3.5	7.8	8.0	444.8	4.5	8.0	8.3	473.2	20.4	27.7	29.0	439.5	7.8	92.9
Trial3_243	36	NA()	1.1	6.1813	1.1	10.2935	3.1	0.4617	2.9	0.93	2447.0	12.2	35.6	36.7	2461.5	7.7	18.8	19.5	2474.3	9.7	17.3	18.5	2474.3	17.3	98.9
Trial3_244	52	NA()	2.2	11.1255	1.6	2.8640	3.5	0.2312	3.1	0.89	1340.8	9.9	22.2	22.8	1372.5	7.8	16.0	16.6	1423.0	12.2	20.3	21.6	1423.0	20.3	94.2
Trial3_245	283	NA()	3.5	12.4748	0.8	2.0503	1.9	0.1856	1.8	0.91	1097.4	4.5	17.2	17.8	1132.4	3.7	13.1	13.7	1200.9	5.7	17.7	19.2	1200.9	17.7	91.4
Trial3_246	106	NA()	0.8	19.5548	3.4	0.2659	4.8	0.0377	3.4	0.72	238.7	2.2	4.4	4.5	239.4	3.8	5.5	5.6	247.1	35.2	40.3	41.2	238.7	4.4	NA
Trial3_247	86	NA()	2.1	13.4164	1.5	1.8056	3.3	0.1758	3.0	0.89	1043.8	7.6	17.6	18.1	1047.5	6.4	13.7	14.2	1056.0	11.5	20.6	22.0	1056.0	20.6	98.8
Trial3_248	129	NA()	2.4	12.6713	1.2	2.0841	2.9	0.1916	2.6	0.91	1130.1	7.2	18.5	19.0	1143.6	5.6	13.9	14.5	1170.1	8.7	19.0	20.4	1170.1	19.0	96.6
Trial3_249	93	NA()	1.4	13.1401	1.5	1.7653	3.3	0.1683	2.9	0.89	1002.8	7.3	16.9	17.4	1032.8	6.4	13.6	14.1	1097.8	12.1	20.9	22.3	1097.8	20.9	91.3
Trial3_250	57	NA()	1.2	13.5113	1.8	1.7179	3.7	0.1684	3.2	0.86	1003.4	8.0	17.2	17.7	1015.2	7.2	13.9	14.4	1041.8	14.7	22.6	23.9	1041.8	22.6	96.3
Trial3_251	76	NA()	1.5	18.5329	3.0	0.4958	4.5	0.0667	3.4	0.75	416.1	3.7	7.6	7.8	408.9	5.5	8.3	8.6	369.3	30.5	36.0	37.0	416.1	7.6	112.7
Trial3_252	114	NA()	1.7	12.2815	1.2	2.1535	2.9	0.1919	2.7	0.91	1131.7	7.2	18.5	19.0	1166.2	5.8	14.1	14.6	1231.7	9.1	19.0	20.4	1231.7	19.0	91.9
Trial3_253	111	NA()	1.2	8.7964	0.9	4.9645	2.4	0.3169	2.2	0.93	1774.4	9.0	27.0	27.9	1813.3	5.9	16.7	17.4	1859.1	7.0	16.9	18.2	1859.1	16.9	95.4
Trial3_254	198	NA()	1.7	12.5762	1.5	0.7587	3.4	0.0692	3.0	0.89	431.5	2.7	7.4	7.6	573.2	3.8	8.9	9.3	1185.0	11.2	20.2	21.6	431.5	7.4	36.4
Trial3_255	394	NA()	3.9	10.8727	1.0	1.0039	2.6	0.0792	2.4	0.92	491.4	1.2	7.9	8.1	705.8	2.2	9.7	10.1	1466.8	6.7	17.5	18.9	1466.8	17.5	33.5
Trial3_256	26	NA()	1.1	5.3248	1.2	12.7665	3.2	0.4933	3.0	0.93	2584.7	12.0	37.0	38.2	2662.6	7.8	19.1	19.9	2723.0	9.9	17.2	18.3	2723.0	17.2	94.9
Trial3_257	67	NA()	2.5	13.6316	1.8	1.6763	3.6	0.1658	3.1	0.87	989.0	7.4	16.8	17.3	999.6	6.7	13.5	14.0	1023.9	13.8	22.1	23.4	1023.9	22.1	96.6
Trial3_258	62	NA()	1.6	13.0544	1.7	1.8815	3.6	0.1782	3.1	0.87	1057.2	7.7	17.8	18.3	1074.6	6.9	14.1	14.6	1110.8	13.5	21.7	23.0	1110.8	21.7	95.2
Trial3_259	78	NA()	1.3	19.3655	3.6	0.2599	4.8	0.0365	3.1	0.66	231.2	2.2	4.3	4.4	234.6	4.6	6.0	6.1	269.4	45.1	49.1	49.9	231.2	4.3	NA
Trial3_260	73	NA()	2.2	13.3841	1.7	1.6839	3.5	0.1635	3.1	0.88	976.4	7.1	16.5	17.0	1002.5	6.4	13.4	13.9	1060.9	12.8	21.4	22.7	1060.9	21.4	92.0
Trial3_261	301	NA()	0.7	18.0781	1.5	0.5055	3.1	0.0663	2.7	0.87	413.9	1.6	6.8	7.0	415.5	2.4	6.7	7.0	425.0	12.5	22.7	24.3	413.9	6.8	97.4
Trial3_262	23	NA()	1.6	6.0373	1.3	10.3457	3.4	0.4532	3.1	0.92	2409.5	12.0	35.1	36.2	2466.2	7.9	18.9	19.6	2514.0	10.3	17.6	18.8	2514.0	17.6	95.8
Trial3_263	154	NA()	1.2	19.6123	2.9	0.2518	4.4	0.0358	3.3	0.76	227.0	1.6	4.0	4.1	228.1	3.0	4.8	5.0	240.3	29.1	35.1	36.2	227.0	4.0	NA
Trial3_264	95	NA()	1.5	17.2357	2.3	0.6541	4.0	0.0818	3.3	0.82	506.9	3.9	8.9	9.2	511.0	4.8	8.9	9.2	530.5	19.7	27.1	28.3	506.9	8.9	95.5
Trial3_265	239	NA()	0.7	13.2678	0.9	1.9236	2.2	0.1852	2.0	0.91	1095.2	4.4	17.1	17.7	1089.3	3.6	12.9	13.5	1078.4	6.5	18.2	19.8	1078.4	18.2	101.6
Trial3_266	151	NA()	0.9	18.9227	2.4	0.3963	4.0	0.0544	3.2	0.80	341.6	2.3	5.9	6.1	339.0	3.7	6.5	6.7	322.2	24.6	31.3	32.5	341.6	5.9	NA
Trial3_267	72	NA()	1.0	19.3190	3.8	0.2631	4.9	0.0369	3.1	0.64	233.5	2.1	4.3	4.4	237.2	5.2	6.5	6.6	274.9	52.5	56.0	56.7	233.5	4.3	NA
Trial3_268	87	NA()	1.8	12.5762	1.4	2.0523	3.2	0.1873	2.9	0.90	1106.6	7.1	18.1	18.7	1133.1	6.1	14.0	14.6	1185.0	10.9	20.0	21.4	1185.0	20.0	93.4
Trial3_269	214	NA()	1.3	11.7977	0.8	2.6404	2.0	0.2260	1.9	0.91	1313.6	5.5	20.3	21.0	1312.0	4.1	14.2	14.9	1310.1	6.0	17.5	19.0	1310.1	17.5	100.3
Trial3_270	111	NA()	1.1	19.7164	3.6	0.1944	4.8	0.0278	3.1	0.65	176.8	1.5	3.2	3.3	180.4	3.7	4.8	4.9	228.1	47.9	51.8	52.5	176.8	3.2	NA
Trial3_271	41	NA()	3.8	13.4537	2.1	1.8270	3.9	0.1784	3.3	0.84	1058.0	8.2	18.0	18.5	1055.2	8.0	14.5	15.0	1050.4	17.8	24.7	25.9	1050.4	24.7	100.7
Trial3_272	84	NA()	0.9	5.2746	0.9	13.5077	2.2	0.5170	2.0	0.91	2686.3	9.3	37.3	38.5	2715.8	5.9	18.4	19.3	2738.6	7.4	15.8	17.0	2738.6	15.8	98.1
Trial3_273	27	NA()	1.6	13.7552	2.7	1.6485	4.3	0.1645	3.4	0.79	981.9	8.3	17.1	17.6	989.0	9.0	14.8	15.2	1005.6	22.2	28.1	29.1	1005.6	28.1	97.6
Trial3_274	99	NA()	2.6	12.5934	1.3	2.1485	3.1	0.1963	2.8	0.91	1155.6	7.4	18.9	19.5	1164.6	5.9	14.1	14.7	1182.3	9.7	19.4	20.8	1182.3	19.4	97.7
Trial3_275	23	NA()	1.5	13.3883	2.8	1.8945	4.4	0.1840	3.4	0.78	1089.0	9.1	18.8	19.3	1079.1	10.2	16.0	16.4	1060.2	24.9	30.2	31.2	1060.2	30.2	102.7
Trial3_276	187	NA()	0.7	20.5323	2.9	0.2066	4.4	0.0308	3.3	0.75	195.4	1.3	3.4	3.5	190.7	2.5	4.1	4.2	133.6	30.2	36.2	37.3	195.4	3.4	NA
Trial3_277	26	NA()	1.5	13.0624	2.6	1.9214	4.3	0.1821	3.4	0.80	1078.5	8.8	18.5	19.0	1088.5	9.4	15.5	16.0	1109.6	21.8	27.6	28.7	1109.6	27.6	97.2
Trial3_278	342	NA()	2.8	12.4252	0.8	1.8190	1.9	0.1640	1.7	0.91	979.0	2.6	15.1	15.7	1052.3	2.6	12.4	13.0	1208.8	5.3	17.5	19.1	1208.8	17.5	81.0
Trial3_279	90	NA()	1.5	11.7569	1.3	2.4550	3.1	0.2094	2.8	0.91	1225.8	8.0	20.0	20.6	1258.9	6.3	14.7	15.3	1316.8	9.5	19.0	20.4	1316.8	19.0	93.1
Trial3_280	105	NA()	2.1	11.5001	1.1	2.7607	2.9	0.2304	2.6	0.92	1336.4	8.1	21.4	22.1	1345.0	6.0	15.0	15.6	1359.5	8.5	18.5	19.9	1359.5	18.5	98.3
Trial3_281	62	NA()	1.0	5.5569	0.8	12.2147	2.5	0.4925	2.4	0.94	2581.5	10.4	36.5	37.6	2621.0	6.4	18.5	19.3	2652.5	8.0	16.2	17.4	2652.5	16.2	97.3
Trial3_282	236	NA()	2.4	9.0529	0.8	4.8222	1.6	0.3168	1.3	0.85	1773.9	6.3	26.2	27.1	1788.8	4.1	16.1	16.8	1807.0	4.8	16.2	17.6	1807.0	16.2	98.2
Trial3_283	87	NA()	1.3	11.7050	1.3	2.6164	3.1	0.2222	2.8	0.91	1293.6	8.5	21.0	21.7	1305.3	6.5	15.1	15.7	1325.4	10.0	19.3	20.6	1325.4	19.3	97.6
Trial3_284	130	NA()	1.2	19.9343	3.2	0.2419	4.7	0.0350	3.4	0.73	221.7	1.9	4.0	4.1	220.0	3.2	4.8	5.0	202.6	31.6	37.2	38.2	221.7	4.0	NA
Trial3_285	78	NA()	2.0	13.2925	1.5	1.9054	3.4	0.1838	3.0	0.89	1087.5	7.8	18.2	18.7	1083.0	6.4	13.9	14.4	1074.7	11.5	20.6	22.0	1074.7	20.6	101.2
Trial3_286	35	NA()	2.0	12.8498	2.2	2.0432	4.0	0.1905	3.3	0.83	1124.1	8.9	19.2	19.7	1130.0	8.5	15.2	15.7	1142.3	17.9	24.6	25.7	1142.3	24.6	98.4
Trial3_287	211	NA()	1.6	9.6741	0.9	3.7769	2.0	0.2651	1.8	0.90	1516.0	6.6	23.1	23.9	1587.9	4.5	15.5	16.2	1685.4	5.4	16.6	18.0	1685.4	16.6	89.9
Trial3_288	155	NA()	0.8	17.7476	2.0	0.5489	3.7	0.0707	3.1	0.84	440.3	3.2	7.7	7.9	444.3	4.0	7.8	8.0	466.0	18.1	26.2	27.5	440.3	7.7	94.5
Trial3_289	47	NA()	1.1	12.7235	1.9	1.9448	3.8	0.1795	3.3	0.86	1064.5	8.4	18.2	18.7	1096.7	7.7	14.6	15.1	1161.9	15.1	22.6	23.8	1161.9	22.6	91.6
Trial3_290	39	NA()	2.0	13.5387	2.3	1.6841	4.1	0.1654	3.3	0.82	986.9	8.0	17.0	17.5	1002.5	8.4	14.5	15.0	1037.7	19.9	26.3	27.4	1037.7	26.3	95.1
Trial3_291	48	NA()	1.8	13.0551	1.9	1.9173	3.8	0.1816	3.2	0.86	1075.8	8.3	18.2	18.8	1087.1	7.6	14.5	15.0	1110.7	15.6	23.1	24.3	1110.7	23.1	96.9
Trial3_292	730	NA()	2.7	13.3164	0.8	1.7624	1.4	0.1703	1.1	0.79	1013.7	3.6	15.8	16.4	1031.7	2.8	12.3	12.9	1071.1	4.0	17.5	19.1	1071.1	17.5	94.6
Trial3_293	129	NA()	0.7	19.8336	3.2	0.2520	4.7	0.0363	3.4	0.73	229.7	1.9	4.2	4.3	228.2	3.4	5.1	5.2	214.4	33.1	38.5	39.5	229.7	4.2	NA
Trial3_294	709	NA()	7.6	12.5240	0.8	2.0687	1.3	0.1880	1.1	0.79	1110.5	3.4	17.1	17.7	1138.5	2.6	12.9	13.5	1193.2	3.7	17.2	18.7	1193.2	17.2	93.1
Trial3_295	184	NA()	1.6	19.6312	2.7	0.2394	4.3	0.0341	3.3	0.77	216.2	1.5	3.8	3.9	217.9	2.8	4.6	4.7	238.1	27.8	34.0	35.1	216.2	3.8	NA
Trial3_296	62	NA()	2.7	12.2724	1.6	2.3778	3.4	0.2117	3.0	0.89	1238.1	8.8	20.5	21.1	1236.0	7.2	15.0	15.6	1233.2	12.4	20.8	22.1	1233.2	20.8	100.4
Trial3_297	107	NA()	1.9	9.0254	0.9	4.8894	2.5	0.3202	2.3	0.93	1790.7	9.3	27.3	28.2	1800.4	6.0	16.7	17.4	1812.5	7.2	17.0	18.4	1812.5	17.0	98.8
Trial3_298	142	NA()	1.3	12.6926	1.1	1.9927	2.8	0.1835	2.6	0.91	1086.2	6.6	17.7	18.2	1113.0	5.3	13.6	14.1	1166.7	8.4	18.8	20.3	1166.7	18.8	93.1
Trial3_299	26	NA()	1.4	13.2735	2.7	1.7170	4.4	0.1654	3.4	0.78	986.5	8.5	17.3	17.7	1014.9	9.8	15.4	15.8	1077.5	24.2	29.6	30.6	1077.5	29.6	91.6
Trial3_300	131	NA()	1.9	11.1085	1.0	2.8973	2.6	0.2335	2.4	0.92	1353.0	7.6	21.4	22.1	1381.2	5.6	15.0	15.7	1426.0	7.5	17.9	19.3	1426.0	17.9	94.9
Trial3_301	152	NA()	1.6	11.1267	0.9	3.0765	2.4	0.2484	2.2	0.92	1430.1	7.4	22.3	23.0	1426.9	5.2	15.1	15.7	1422.8	6.9	17.7	19.1	1422.8	17.7	100.5
Trial3_302	107	NA()	1.3	11.0326	1.1	3.2088	2.7	0.2569	2.5	0.92	1473.8	8.5	23.2	23.9	1459.3	5.9	15.5	16.1	1439.0	8.0	18.1	19.5	1439.0	18.1	102.4
Trial3_303	96	NA()	2.3	19.5790	3.1	0.3307	4.6	0.0470	3.4	0.74	295.9	2.6	5.4	5.5	290.1	4.1	6.2	6.4	244.2	31.1	36.8	37.8	295.9	5.4	NA
Trial3_304	59	NA()	1.7	12.9539	1.8	1.8492	3.6	0.1738	3.2	0.87	1033.1	7.7	17.5	18.0	1063.1	7.1	14.1	14.6	1126.3	14.2	22.1	23.4	1126.3	22.1	91.7
Trial3_305	681	NA()	1.1	18.4406	1.0	0.4929	2.2	0.0660	1.9	0.88	411.7	1.4	6.7	6.9	406.9	1.7	6.4	6.7	380.5	8.1	20.8	22.5	411.7	6.7	108.2
Trial3_306	107	NA()	0.5	5.3702	0.9	12.8998	2.0	0.5026	1.7	0.89	2625.2	8.8	36.5	37.7	2672.4	5.6	18.3	19.1	2709.0	7.0	15.7	16.9	2709.0	15.7	96.9
Trial3_307	46	NA()	1.3	11.2585	1.6	3.0791	3.5	0.2515	3.1	0.89	1446.4	9.8	23.4	24.1	1427.5	7.5	16.1	16.7	1400.3	12.1	20.3	21.6	1400.3	20.3	103.3
Trial3_308	68	NA()	1.5	13.7340	1.7	1.7453	3.5	0.1739	3.1	0.88	1033.7	7.5	17.4	17.9	1025.4	6.7	13.7	14.2	1008.7	13.5	21.9	23.2	1008.7	21.9	102.5
Trial3_309	48	NA()	0.6	10.7742	1.5	3.1568	3.4	0.2468	3.1	0.90	1421.9	9.6	23.1	23.7	1446.7	7.5	16.1	16.7	1484.1	11.6	19.8	21.1	1484.1	19.8	95.8
Trial3_310	52	NA()	0.4	18.9309	4.6	0.1939	5.6	0.0266	3.2	0.57	169.5	1.8	3.3	3.4	180.0	6.0	6.7	6.8	321.3	78.4	80.8	81.3	169.5	3.3	NA
Trial3_311	49	NA()	0.8	11.1667	1.6	2.8005	3.5	0.2269	3.1	0.89	1318.3	9.4	21.7	22.4	1355.7	7.8	15.9	16.5	1416.0	13.0	20.8	22.0	1416.0	20.8	93.1
Trial3_312	92	NA()	0.3	5.2306	0.9	13.3697	2.1	0.5074	1.9	0.90	2645.6	9.3	36.8	38.1	2706.1	5.9	18.4	19.2	2752.3	7.4	15.8	17.0	2752.3	15.8	96.1
Trial3_313	22	NA()	0.7	13.5032	3.1	1.6944	4.6	0.1660	3.5	0.75	990.1	8.7	17.4	17.9	1006.4	12.0	16.8	17.2	1043.0	32.7	36.9	37.8	1043.0	36.9	94.9
Trial3_314	37	NA()	1.3	10.9404	1.8	3.0846	3.6	0.2449	3.2	0.88	1411.9	10.1	23.2	23.8	1428.9	8.2	16.4	17.0	1455.0	13.7	21.2	22.4	1455.0	21.2	97.0
Trial3_315	201	NA()	1.5	13.8138	1.1	1.5890	2.6	0.1593	2.3	0.91	952.7	5.1	15.4	15.9	965.9	4.3	12.3	12.8	997.0	8.0	19.0	20.5	997.0	19.0	95.6
Trial3_316	78	NA()	1.4	11.3166	1.3	3.0902	3.1	0.2537	2.8	0.91	1457.8	9.2	23.3	24.0	1430.3	6.6	15.7	16.3	1390.4	9.6	18.9	20.3	1390.4	18.9	104.8
Trial3_317	29	NA()	1.5	12.5876	2.5	1.8802	4.2	0.1717	3.4	0.81	1021.6	8.8	17.8	18.3	1074.1	9.6	15.6	16.0	1183.2	21.8	27.5	28.5	1183.2	27.5	86.3
Trial3_318	43	NA()	1.2	13.1897	2.1	1.7605	3.9	0.1685	3.3	0.84	1003.8	8.3	17.4	17.9	1031.0	8.1	14.5	14.9	1090.2	17.5	24.4	25.6	1090.2	24.4	92.1
Trial3_319	105	NA()	1.2	13.5333	1.4	1.8116	3.1	0.1779	2.8	0.90	1055.5	7.3	17.6	18.1	1049.7	6.0	13.5	14.0	1038.5	10.8	20.3	21.7	1038.5	20.3	101.6
Trial3_320	80	NA()	2.1	11.9502	1.3	2.5169	3.2	0.2182	2.9	0.91	1272.6	8.7	20.9	21.5	1276.9	6.7	15.0	15.6	1285.1	10.3	19.5	20.8	1285.1	19.5	99.0
Trial3_321	92	NA()	0.5	10.7374	1.1	3.3320	2.8	0.2596	2.6	0.92	1487.8	9.1	23.7	24.4	1488.6	6.4	15.8	16.5	1490.6	8.5	18.2	19.6	1490.6	18.2	99.8
Trial3_322	139	NA()	2.4	13.0757	1.2	1.9561	2.8	0.1856	2.6	0.91	1097.4	6.8	17.9	18.5	1100.5	5.4	13.6	14.1	1107.6	8.7	19.1	20.6	1107.6	19.1	99.1
Trial3_323	102	NA()	11.4	11.0031	1.1	2.9135	2.8	0.2326	2.6	0.92	1348.1	8.4	21.7	22.3	1385.4	6.2	15.3	15.9	1444.1	8.4	18.2	19.6	1444.1	18.2	93.4
Trial3_324	316	NA()	2.2	17.1273	1.3	0.6650	2.8	0.0826	2.5	0.89	511.8	2.0	8.3	8.6	517.6	2.5	7.9	8.3	544.3	9.9	21.1	22.7	511.8	8.3	94.0
Trial3_325	245	NA()	1.9	11.7488	0.8	2.6239	1.9	0.2237	1.7	0.91	1301.3	5.3	20.1	20.7	1307.4	4.0	14.2	14.8	1318.1	5.6	17.4	18.9	1318.1	17.4	98.7
Trial3_326	32	NA()	1.1	13.2672	2.4	1.8762	4.1	0.1806	3.4	0.82	1070.3	8.8	18.4	18.9	1072.7	9.2	15.3	15.8	1078.5	21.4	27.4	28.4	1078.5	27.4	99.2
Trial3_327	66	NA()	0.7	9.0787	1.1	4.8233	2.9	0.3177	2.7	0.93	1778.7	10.3	27.5	28.4	1789.0	6.9	17.0	17.7	1801.8	8.9	17.8	19.1	1801.8	17.8	98.7
Trial3_328	54	NA()	1.5	11.8404	1.6	2.5563	3.5	0.2196	3.1	0.89	1279.9	9.3	21.2	21.8	1288.2	7.4	15.4	16.0	1303.1	12.2	20.5	21.8	1303.1	20.5	98.2
Trial3_329	104	NA()	1.6	13.2589	1.4	1.8935	3.1	0.1822	2.8	0.90	1078.8	7.4	17.9	18.4	1078.8	6.0	13.7	14.2	1079.7	10.4	20.0	21.4	1079.7	20.0	99.9
Trial3_330	50	NA()	1.4	12.8169	1.9	1.8337	3.8	0.1705	3.2	0.86	1015.0	8.2	17.4	17.9	1057.6	7.7	14.4	14.9	1147.4	15.5	22.9	24.1	1147.4	22.9	88.5
Trial3_331	87	NA()	2.1	13.0994	1.5	1.8614	3.3	0.1769	3.0	0.89	1050.1	7.6	17.6	18.1	1067.5	6.3	13.8	14.3	1104.0	11.2	20.4	21.7	1104.0	20.4	95.1
Trial3_332	96	NA()	2.8	9.0601	0.9	4.7638	2.6	0.3132	2.4	0.93	1756.3	9.7	27.0	27.9	1778.5	6.4	16.8	17.5	1805.6	7.7	17.3	18.6	1805.6	17.3	97.3
Trial3_333	312	NA()	2.8	18.0039	1.5	0.4635	3.1	0.0605	2.7	0.87	378.9	1.7	6.3	6.5	386.7	2.4	6.4	6.7	434.1	13.1	23.0	24.6	378.9	6.3	NA
Trial3_334	38	NA()	2.0	11.3929	1.8	2.9175	3.7	0.2412	3.2	0.88	1392.8	10.2	23.0	23.6	1386.5	8.1	16.2	16.8	1377.5	13.4	21.1	22.4	1377.5	21.1	101.1

Trial3_335	36	NA()	2.2	12.8174	2.2	1.9102	4.0	0.1777	3.3	0.83	1054.1	8.7	18.2	18.7	1084.7	8.5	15.0	15.5	1147.4	18.2	24.8	25.9	1147.4	24.8	91.9
Trial3_336	212	NA()	2.4	13.4587	1.0	1.8928	2.3	0.1848	2.1	0.91	1093.4	5.5	17.4	18.0	1078.6	4.3	13.0	13.6	1049.7	7.2	18.6	20.1	1049.7	18.6	104.2
Trial3_337	175	NA()	1.5	8.2907	0.9	5.2068	1.9	0.3132	1.7	0.90	1756.6	7.8	26.4	27.3	1853.7	5.1	16.6	17.3	1965.3	5.6	16.2	17.5	1965.3	16.2	89.4
Trial3_338	31	NA()	3.4	13.3739	2.5	1.8266	4.2	0.1772	3.4	0.81	1051.9	8.9	18.2	18.7	1055.0	9.2	15.3	15.7	1062.4	21.6	27.6	28.6	1062.4	27.6	99.0
Trial3_339	83	NA()	2.2	11.4849	1.3	2.7856	3.1	0.2321	2.8	0.91	1345.7	8.8	21.8	22.5	1351.7	6.6	15.3	15.9	1362.1	9.5	18.9	20.3	1362.1	18.9	98.8
Trial3_340	85	NA()	2.2	16.0318	2.1	0.8327	3.9	0.0969	3.3	0.84	596.0	4.9	10.6	10.9	615.1	5.5	10.1	10.5	687.0	17.3	25.1	26.4	596.0	10.6	86.7
Trial3_341	102	NA()	2.3	12.8150	1.3	2.0679	3.1	0.1923	2.8	0.90	1133.7	7.6	18.7	19.2	1138.2	6.1	14.0	14.6	1147.7	10.0	19.6	21.0	1147.7	19.6	98.8
Trial3_342	47	NA()	1.4	13.0829	2.0	1.8343	3.8	0.1741	3.3	0.85	1034.8	8.3	17.7	18.2	1057.8	8.0	14.5	15.0	1106.5	17.1	24.1	25.3	1106.5	24.1	93.5
Trial3_343	95	NA()	1.5	11.0087	1.1	3.0430	2.9	0.2431	2.7	0.92	1402.6	8.7	22.5	23.1	1418.5	6.3	15.5	16.1	1443.2	8.5	18.3	19.7	1443.2	18.3	97.2
Trial3_344	110	NA()	2.6	13.5797	1.4	1.7645	3.1	0.1739	2.8	0.90	1033.3	7.0	17.2	17.7	1032.5	5.8	13.3	13.8	1031.6	10.4	20.1	21.5	1031.6	20.1	100.2
Trial3_345	56	NA()	1.4	13.4668	1.8	1.8730	3.6	0.1830	3.2	0.87	1083.4	8.3	18.4	18.9	1071.6	7.2	14.2	14.7	1048.5	14.1	22.2	23.5	1048.5	22.2	103.3
Trial3_346	71	NA()	1.4	11.4405	1.4	2.7711	3.2	0.2300	2.9	0.91	1334.7	9.1	21.8	22.4	1347.8	6.9	15.4	16.0	1369.5	10.4	19.4	20.7	1369.5	19.4	97.5
Trial3_347	591	NA()	0.8	18.0652	1.1	0.5196	2.3	0.0681	2.0	0.88	424.7	0.7	6.8	7.0	424.9	1.4	6.6	6.9	426.6	8.4	20.8	22.4	424.7	6.8	99.6
Trial3_348	19	NA()	1.6	12.7265	2.9	2.0936	4.3	0.1933	3.1	0.74	1139.4	10.0	19.8	20.4	1146.7	10.8	16.7	17.2	1161.5	24.7	29.9	30.9	1161.5	29.9	98.1
Trial3_349	87	NA()	1.0	10.7220	1.2	2.9307	3.0	0.2280	2.8	0.92	1324.0	8.7	21.5	22.1	1389.9	6.5	15.5	16.1	1493.3	9.0	18.4	19.8	1493.3	18.4	88.7
Trial3_350	27	NA()	1.5	7.7045	1.5	6.9013	3.5	0.3858	3.1	0.91	2103.3	12.5	32.0	33.0	2098.8	8.4	18.5	19.2	2095.2	11.4	18.8	20.0	2095.2	18.8	100.4
Trial3_351	589	NA()	3.5	13.4703	0.8	1.6737	1.3	0.1636	1.0	0.76	976.7	1.2	14.9	15.5	998.6	1.6	11.9	12.4	1048.0	4.2	17.7	19.3	1048.0	17.7	93.2
Trial3_352	188	NA()	3.5	11.4602	0.9	2.8949	2.2	0.2407	2.0	0.92	1390.5	6.8	21.7	22.3	1380.6	4.8	14.8	15.4	1366.2	6.4	17.6	19.0	1366.2	17.6	101.8
Trial3_353	108	NA()	1.2	9.0590	0.9	4.7946	2.4	0.3152	2.3	0.93	1766.0	8.8	26.8	27.7	1784.0	5.8	16.6	17.3	1805.8	7.0	17.0	18.3	1805.8	17.0	97.8
Trial3_354	118	NA()	2.7	13.6292	1.3	1.6914	3.1	0.1673	2.8	0.90	997.0	6.3	16.4	16.9	1005.3	5.4	13.0	13.5	1024.2	10.2	20.0	21.5	1024.2	20.0	97.3
Trial3_355	86	NA()	2.6	13.5237	1.6	1.6629	3.4	0.1632	3.0	0.89	974.4	6.7	16.3	16.8	994.5	6.0	13.2	13.7	1039.9	12.0	21.0	22.3	1039.9	21.0	93.7
Trial3_356	46	NA()	1.2	13.6805	2.1	1.6928	3.9	0.1680	3.3	0.84	1001.3	7.5	17.0	17.5	1005.8	7.4	14.0	14.4	1016.6	17.0	24.2	25.4	1016.6	24.2	98.5
Trial3_357	70	NA()	0.9	5.4956	0.9	12.4689	2.4	0.4972	2.2	0.93	2601.8	9.6	36.4	37.6	2640.4	6.0	18.4	19.2	2670.8	7.4	15.9	17.1	2670.8	15.9	97.4
Trial3_358	32	NA()	1.4	7.5288	1.3	6.8949	3.3	0.3767	3.0	0.92	2060.7	11.3	31.1	32.0	2098.0	7.6	18.1	18.8	2135.6	10.1	18.0	19.2	2135.6	18.0	96.5
Trial3_359	500	NA()	1.3	19.9532	1.8	0.1956	3.4	0.0283	2.9	0.84	180.0	0.3	2.9	3.0	181.4	1.2	3.3	3.4	200.5	16.7	25.8	27.3	180.0	2.9	NA
Trial3_360	284	NA()	3.0	12.0997	0.8	2.3540	1.8	0.2067	1.6	0.90	1211.1	3.9	18.5	19.2	1228.8	3.2	13.6	14.2	1260.9	5.3	17.4	19.0	1260.9	17.4	96.0
Trial3_361	53	NA()	2.2	13.0366	1.8	1.9831	3.7	0.1876	3.2	0.87	1108.3	7.9	18.5	19.0	1109.8	7.3	14.5	15.0	1113.6	15.1	22.8	24.0	1113.6	22.8	99.5
Trial3_362	115	NA()	1.7	19.1210	3.4	0.2483	4.8	0.0344	3.4	0.71	218.3	1.8	3.9	4.0	225.2	3.5	5.1	5.3	298.5	35.1	40.1	41.1	218.3	3.9	NA
Trial3_363	93	NA()	2.3	14.5471	1.6	1.4007	3.4	0.1478	3.0	0.88	888.9	6.0	14.9	15.3	889.2	5.6	12.3	12.8	891.0	12.9	21.8	23.2	888.9	14.9	99.8
Trial3_364	36	NA()	0.7	10.7112	1.7	3.3642	3.6	0.2615	3.2	0.88	1497.3	10.0	24.1	24.8	1496.1	7.9	16.5	17.1	1495.2	12.9	20.6	21.9	1495.2	20.6	100.1
Trial3_365	594	NA()	0.9	19.8163	1.7	0.1924	3.3	0.0277	2.8	0.85	175.9	0.5	2.9	3.0	178.7	1.1	3.2	3.4	216.4	15.0	24.7	26.3	175.9	2.9	NA
Trial3_366	51	NA()	1.8	7.6884	1.1	7.0583	3.0	0.3938	2.7	0.93	2140.2	10.6	31.7	32.7	2118.8	6.8	17.8	18.5	2098.8	8.7	17.3	18.5	2098.8	17.3	102.0
Trial3_367	135	NA()	2.3	17.3829	2.2	0.5025	3.9	0.0634	3.2	0.83	396.1	2.8	6.9	7.1	413.4	3.8	7.4	7.6	511.8	18.9	26.6	27.9	396.1	6.9	NA
Trial3_368	13	NA()	0.9	11.5973	3.1	2.6423	4.4	0.2224	3.1	0.72	1294.3	10.6	22.0	22.6	1312.5	12.1	18.2	18.7	1343.3	26.6	31.2	32.1	1343.3	31.2	96.4
Trial3_369	16	NA()	0.8	5.6169	1.5	11.3453	3.6	0.4624	3.3	0.91	2450.1	13.0	35.9	37.0	2551.9	8.7	19.3	20.1	2634.6	11.2	18.0	19.1	2634.6	18.0	93.0
Trial3_370	316	NA()	2.4	13.4210	0.8	1.6834	2.0	0.1639	1.8	0.90	978.6	2.9	15.2	15.7	1002.3	2.8	12.1	12.7	1055.3	5.9	18.1	19.7	1055.3	18.1	92.7
Trial3_371	27	NA()	1.3	4.6762	1.0	16.6605	3.1	0.5653	2.9	0.94	2888.5	11.1	39.8	41.1	2915.5	7.2	19.1	19.9	2934.9	9.3	16.6	17.7	2934.9	16.6	98.4
Trial3_372	568	NA()	2.5	13.6757	0.8	1.3706	1.5	0.1360	1.3	0.83	822.0	1.3	12.7	13.2	876.4	1.7	11.0	11.5	1017.3	4.8	17.9	19.5	1017.3	17.9	80.8
Trial3_373	163	NA()	5.6	13.6872	1.1	1.6771	2.7	0.1666	2.5	0.91	993.1	5.4	16.0	16.6	999.9	4.6	12.6	13.2	1015.6	8.5	19.2	20.7	1015.6	19.2	97.8
Trial3_374	103	NA()	2.3	11.8590	1.2	2.6531	2.9	0.2283	2.6	0.91	1325.6	7.5	21.1	21.7	1315.5	5.7	14.8	15.4	1300.0	8.6	18.6	20.1	1300.0	18.6	102.0
Trial3_375	63	NA()	0.6	11.6204	1.5	2.5557	3.4	0.2155	3.0	0.90	1258.0	8.4	20.6	21.2	1288.1	6.8	15.1	15.7	1339.4	11.1	19.8	21.1	1339.4	19.8	93.9
Trial3_376	73	NA()	0.6	5.3246	0.9	13.6766	2.3	0.5284	2.1	0.92	2734.7	9.4	37.8	39.0	2727.6	5.9	18.5	19.3	2723.0	7.7	16.0	17.2	2723.0	16.0	100.4
Trial3_377	629	NA()	6.0	13.5761	0.8	1.7927	1.3	0.1766	1.0	0.78	1048.3	2.4	16.1	16.6	1042.8	2.1	12.2	12.8	1032.1	4.1	17.7	19.3	1032.1	17.7	101.6
Trial3_378	157	NA()	1.9	19.6820	2.8	0.2654	4.3	0.0379	3.3	0.76	239.8	1.7	4.2	4.4	239.0	3.0	4.9	5.1	232.1	27.6	33.9	35.0	239.8	4.2	NA
Trial3_379	66	NA()	1.5	12.8611	1.6	2.0730	3.4	0.1935	3.0	0.89	1140.0	7.7	18.8	19.4	1139.9	6.5	14.2	14.8	1140.6	11.8	20.6	22.0	1140.6	20.6	100.0
Trial3_380	35	NA()	0.7	12.4529	2.0	2.3250	3.9	0.2101	3.3	0.85	1229.3	9.1	20.5	21.1	1220.0	8.2	15.5	16.0	1204.4	16.0	23.2	24.4	1204.4	23.2	102.1
Trial3_381	507	NA()	0.6	18.9788	1.7	0.2163	3.4	0.0298	2.9	0.85	189.2	0.3	3.1	3.2	198.8	1.2	3.6	3.7	315.5	14.7	24.3	25.8	189.2	3.1	NA
Trial3_382	36	NA()	0.9	9.5537	1.5	4.3798	3.4	0.3036	3.1	0.90	1709.2	10.7	26.9	27.7	1708.5	7.8	17.2	17.8	1708.5	11.5	19.4	20.6	1708.5	19.4	100.0
Trial3_383	34	NA()	0.8	12.8740	2.2	2.0669	4.0	0.1931	3.3	0.84	1138.0	8.8	19.2	19.8	1137.9	8.4	15.2	15.7	1138.6	17.8	24.6	25.7	1138.6	24.6	99.9
Trial3_384	82	NA()	0.6	5.3919	0.9	13.6404	2.2	0.5337	2.0	0.91	2756.8	9.2	38.0	39.2	2725.1	5.8	18.4	19.2	2702.3	7.6	15.9	17.1	2702.3	15.9	102.0
Trial3_385	240	NA()	2.6	12.6846	0.9	2.1637	2.1	0.1991	1.9	0.91	1170.7	5.3	18.4	19.0	1169.5	4.1	13.5	14.1	1168.0	6.3	18.0	19.5	1168.0	18.0	100.2
Trial3_386	114	NA()	1.3	17.9902	2.3	0.5366	4.0	0.0700	3.3	0.81	436.4	3.6	7.8	8.0	436.2	4.6	8.0	8.3	435.8	21.6	28.7	29.9	436.4	7.8	100.1
Trial3_387	89	NA()	3.0	13.0706	1.4	1.9158	3.3	0.1817	2.9	0.90	1076.2	7.8	18.0	18.6	1086.6	6.4	13.9	14.4	1108.4	10.9	20.2	21.6	1108.4	20.2	97.1
Trial3_388	87	NA()	1.5	13.5331	1.5	1.7736	3.3	0.1742	3.0	0.89	1035.0	7.6	17.5	18.0	1035.9	6.3	13.6	14.1	1038.5	11.5	20.6	22.0	1038.5	20.6	99.7
Trial3_389	43	NA()	1.8	4.8713	0.9	15.2175	2.8	0.5379	2.6	0.94	2774.5	11.2	38.7	39.9	2828.9	7.0	19.0	19.8	2868.7	8.9	16.4	17.6	2868.7	16.4	96.7
Trial3_390	113	NA()	0.9	8.7513	0.8	5.5502	2.3	0.3524	2.1	0.93	1946.2	9.8	29.3	30.2	1908.4	6.1	17.0	17.8	1868.4	7.3	17.0	18.3	1868.4	17.0	104.2
Trial3_391	500	NA()	1.7	14.1347	0.7	1.5272	1.4	0.1566	1.2	0.87	938.0	1.0	14.4	14.9	941.4	1.6	11.5	12.0	950.1	4.9	18.0	19.7	950.1	18.0	98.7
Trial3_392	115	NA()	1.0	10.8968	1.0	3.1353	2.7	0.2479	2.5	0.92	1427.6	8.7	22.8	23.4	1441.4	6.1	15.5	16.1	1462.6	8.0	18.0	19.4	1462.6	18.0	97.6
Trial3_393	53	NA()	1.4	11.5452	1.5	2.7934	3.4	0.2340	3.1	0.89	1355.4	9.9	22.4	23.0	1353.8	7.6	15.8	16.4	1352.0	11.8	20.2	21.5	1352.0	20.2	100.3
Trial3_394	117	NA()	2.7	14.0175	1.4	1.6194	3.1	0.1647	2.8	0.90	982.9	6.9	16.5	17.0	977.8	5.7	12.9	13.4	967.2	10.3	20.2	21.6	967.2	20.2	101.6
Trial3_395	161	NA()	1.1	9.0405	0.9	5.1193	2.0	0.3358	1.8	0.90	1866.5	8.6	28.0	28.9	1839.3	5.4	16.6	17.3	1809.5	6.3	16.7	18.1	1809.5	16.7	103.1
Trial3_396	140	NA()	4.7	13.4794	1.2	1.7335	2.9	0.1695	2.6	0.91	1009.6	6.7	16.7	17.2	1021.0	5.4	13.1	13.6	1046.6	9.2	19.4	20.9	1046.6	19.4	96.5
Trial3_397	108	NA()	2.4	10.9550	1.1	3.1204	2.7	0.2480	2.5	0.92	1428.3	8.9	22.8	23.5	1437.7	6.3	15.5	16.2	1452.5	8.1	18.1	19.5	1452.5	18.1	98.3
Trial3_398	28	NA()	1.3	13.4986	2.7	1.6930	4.4	0.1658	3.4	0.78	989.1	8.9	17.5	18.0	1005.9	9.9	15.4	15.8	1043.7	24.3	29.7	30.7	1043.7	29.7	94.8
Trial3_399	163	NA()	2.2	8.5685	0.9	5.4267	2.0	0.3374	1.7	0.90	1874.1	8.6	28.1	29.0	1889.1	5.4	16.8	17.5	1906.4	6.1	16.4	17.8	1906.4	16.4	98.3
Trial3_400	34	NA()	1.6	9.3682	1.6	4.4697	3.5	0.3038	3.2	0.90	1710.2	11.6	27.2	28.0	1725.4	8.4	17.5	18.1	1744.5	12.0	19.6	20.8	1744.5	19.6	98.0
Trial3_401	91	NA()	2.3	13.6871	1.5	1.7242	3.3	0.1712	3.0	0.89	1018.9	7.5	17.2	17.7	1017.6	6.3	13.5	14.0	1015.6	11.9	20.9	22.3	1015.6	20.9	100.3
Trial3_402	126	NA()	4.1	4.9754	0.9	15.8372	1.7	0.5717	1.4	0.84	2915.0	8.6	39.5	40.8	2867.0	5.5	18.5	19.3	2834.2	7.3	15.7	16.9	2834.2	15.7	102.8
Trial3_403	148	NA()	3.0	11.0355	0.9	3.2591	2.4	0.2610	2.2	0.92	1494.8	8.1	23.4	24.1	1471.3	5.5	15.4	16.0	1438.5	7.0	17.7	19.1	1438.5	17.7	103.9
Trial3_404	432	NA()	3.8	13.5004	0.7	1.7317	1.5	0.1696	1.3	0.88	1010.1	1.7	15.5	16.0	1020.4	2.0	12.1	12.6	1043.4	5.0	17.9	19.5	1043.4	17.9	96.8
Trial3_405	81	NA()	1.8	8.7988	1.0	5.0945	2.8	0.3253	2.6	0.93	1815.3	10.2	27.9	28.8	1835.2	6.6	17.0	17.7	1858.6	8.0	17.3	18.6	1858.6	17.3	97.7
Trial3_406	379	NA()	0.9	19.8983	2.2	0.1866	3.8	0.0269	3.1	0.82	171.4	0.6	2.8	2.9	173.7	1.5	3.3	3.4	206.8	20.3	28.3	29.7	171.4	2.8	NA
Trial3_407	72	NA()	1.2	10.8864	1.3	3.1375	3.1	0.2478	2.9	0.91	1427.3	9.3	23.0	23.7	1441.9	6.9	15.8	16.4	1464.4	9.9	18.9	20.3	1464.4	18.9	97.5
Trial3_408	152	NA()	1.1	11.5214	1.0	2.6930	2.5	0.2251	2.3	0.92	1308.9	7.2	20.7	21.4	1326.5	5.3	14.7	15.3	1355.9	7.2	17.9	19.4	1355.9	17.9	96.5
Trial3_409	128	NA()	1.6	11.6405	1.0	2.7807	2.6	0.2349	2.4	0.92	1359.9	7.7	21.6	22.2	1350.4	5.6	14.9	15.5	1336.1	7.8	18.2	19.6	1336.1	18.2	101.8
Trial3_410	196	NA()	3.2	5.1453	0.9	13.8711	1.3	0.5179	1.0	0.75	2690.1	7.0	36.8	38.1	2740.9	4.3	18.0	18.9	2779.3	5.3	14.9	16.2	2779.3	14.9	96.8
Trial3_411	89	NA()	0.5	16.5524	2.2	0.7793	3.9	0.0936	3.3	0.83	576.8	4.7	10.2	10.5	585.1	5.4	9.8	10.2	618.4	18.7	26.2	27.5	576.8	10.2	93.3
Trial3_412	261	NA()	1.2	17.1188	1.6	0.5359	3.2	0.0666	2.8	0.87	415.4	2.1	6.9	7.2	435.7	2.8	7.1	7.4	545.4	12.8	22.6	24.1	415.4	6.9	76.2
Trial3_413	53	NA()	1.0	5.3089	0.9	13.7451	2.6	0.5295	2.5	0.94	2739.3	10.6	38.2	39.4	2732.3	6.6	18.7	19.5	2727.9	8.5	16.4	17.6	2727.9	16.4	100.4
Trial3_414	550	NA()	2.5	18.3197	1.2	0.4313	2.6	0.0573	2.3	0.88	359.4	0.5	5.8	6.0	364.1	1.4	5.8	6.1	395.3	9.7	21.4	23.0	359.4	5.8	NA
Trial3_415	311	NA()	1.3	12.9446	0.8	2.0728	1.8	0.1947	1.6	0.90	1146.7	3.6	17.6	18.2	1139.8	3.0	13.0	13.6	1127.7	5.4	17.8	19.4	1127.7	17.8	101.7
Trial3_416	89	NA()	0.9	10.7501	1.1	3.2713	2.9	0.2552	2.7	0.92	1465.1	8.8	23.3	24.0	1474.2	6.3	15.7	16.4	1488.3	8.8	18.3	19.7	1488.3	18.3	98.4
Trial3_417	43	NA()	3.3	12.3682	1.9	2.3318	3.7	0.2093	3.2	0.86	1224.9	9.1	20.5	21.0	1222.0	8.1	15.4	16.0	1217.9	15.6	22.8	24.1	1217.9	22.8	100.6
Trial3_418	29	NA()	0.6	10.9503	2.0	3.1668	3.8	0.2516	3.3	0.86	1446.8	10.5	23.7	24.4	1449.1	8.8	16.8	17.4	1453.3	15.2	22.2	23.4	1453.3	22.2	99.6
Trial3_419	169	NA()	1.9	12.8873	1.0	2.1540	2.5	0.2014	2.3	0.91	1182.9	6.2	18.8	19.4	1166.3	4.8	13.7	14.3	1136.6	7.5	18.5	20.0	1136.6	18.5	104.1
Trial3_420	132	NA()	2.3	9.3482	0.8	4.7282	2.2	0.3207	2.1	0.93	1793.2	8.3	27.0	27.9	1772.3	5.4	16.4	17.1	1748.4	6.6	16.9	18.3	1748.4	16.9	102.6
Trial3_421	195	NA()	3.2	9.3225	0.9	4.3880	1.9	0.2968	1.7	0.89	1675.5	6.7	25.1	26.0	1710.1	4.5	15.9	16.6	1753.5	5.3	16.4	17.9	1753.5	16.4	95.6
Trial3_422	84	NA()	2.2	12.2510	1.4	2.2517	3.2	0.2002	2.9	0.90	1176.2	7.7	19.3	19.8	1197.3	6.3	14.4	15.0	1236.6	10.4	19.7	21.0	1236.6	19.7	95.1
Trial3_423	515	NA()	2.4	18.9487	1.5	0.2949	3.1	0.0405	2.7	0.87	256.2	0.4	4.1	4.3	262.4	1.3	4.5	4.7	319.1	12.7	23.1	24.7	256.2	4.1	NA
Trial3_424	301	NA()	0.4	11.0946	0.8	3.0148	1.6	0.2427	1.3	0.85	1400.7	4.1	21.1	21.8	1411.4	3.1	14.4	15.1	1428.3	4.6	16.9	18.4	1428.3	16.9	98.1
Trial3_425	101	NA()	0.7	5.7458	0.9	12.5156	2.0	0.5218	1.8	0.89	2706.8	8.6	37.3	38.6	2643.9	5.3	18.2	19.0	2596.9	6.9	15.8	17.0	2596.9	15.8	104.2
Trial3_426	126	NA()	1.4	9.7586	0.9	4.1153	2.4	0.2914	2.2	0.93	1648.5	7.8	25.1	25.9	1657.3	5.3	16.0	16.7	1669.4	6.7	17.1	18.5	1669.4	17.1	98.7
Trial3_427	197	NA()	1.2	18.3747	2.0	0.4100	3.7	0.0547	3.1	0.83	343.1	2.0	5.8	6.0	348.8	2.9	6.2	6.4	388.6	17.8	26.1	27.5	343.1	5.8	NA
Trial3_428	159	NA()	0.7	19.5562	3.4	0.1833	4.8	0.0260	3.4	0.71	165.5	1.2	2.9	3.0	170.9	2.8	4.0	4.1	246.9	37.3	42.1	43.0	165.5	2.9	NA
Trial3_429	98	NA()	2.1	10.8290	1.1	3.1566	2.8	0.2480	2.6	0.92	1428.3	7.9	22.5	23.2	1446.6	5.8	15.4	16.0	1474.5	8.1	18.0	19.4	1474.5	18.0	96.9
Trial3_430	66	NA()	1.6	10.0083	1.3	3.4195	3.2	0.2483	2.9	0.91	1429.8	8.8	22.8	23.5	1508.9	6.7	16.0	16.6	1622.5	9.4	18.4	19.7	1622.5	18.4	88.1
Trial3_431	56	NA()	0.6	14.8601	2.2	1.2434	3.9	0.1341	3.3	0.83	811.1	6.3	14.0	14.4	820.4	6.7	12.4	12.8	846.9	17.7	25.0	26.3	811.1	14.0	95.8
Trial3_432	231	NA()	1.1	13.4497	0.9	1.8281	2.3	0.1784	2.1	0.91	1058.2	4.4	16.6	17.2	1055.6	3.7	12.7	13.2	1051.0	6.7	18.4	20.0	1051.0	18.4	100.7
Trial3_433	116	NA()	2.0	16.6799	1.9	0.8025	3.6	0.0971	3.1	0.85	597.5	4.2	10.3	10.6	598.2	4.7	9.6	10.0	601.8	16.1	24.5	25.8	597.5	10.3	99.3
Trial3_434	170	NA()	2.5	19.6323	2.8	0.2377	4.4	0.0339	3.3	0.76	214.6	1.5	3.8	3.9	216.5	2.9	4.6	4.7	238.0	29.7	35.6	36.7	214.6	3.8	NA
Trial3_435	131	NA()	2.2	13.7362	1.3	1.7473	2.9	0.1742	2.7	0.90	1034.9	6.2	16.9	17.4	1026.2	5.1	13.0	13.5	1008.4	9.5	19.7	21.1	1008.4	19.7	102.6
Trial3_436	280	NA()	1.8	19.9023	2.2	0.2508	3.8	0.0362	3.1	0.82	229.4	1.1	3.9	4.0	227.2	2.0	4.2	4.4	206.4	19.4	27.6	29.0	229.4	3.9	NA
Trial3_437	319	NA()	1.5	19.3043	1.9	0.2835	3.5	0.0397	3.0	0.84	251.0	1.0	4.2	4.3	253.4	1.9	4.6	4.8	276.7	17.3	26.0	27.5	251.0	4.2	NA
Trial3_438	91	NA()	1.6	13.4239	1.5	1.7450	3.3	0.1700	2.9	0.89	1011.9	6.8	16.8	17.3	1025.3	5.9	13.3	13.8	1054.9	11.0	20.4	21.8	1054.9	20.4	95.9
Trial3_439	398	NA()	1.2	13.1131	0.8	1.8926	1.6	0.1801	1.3	0.85	1067.4	1.9	16.3	16.8	1078.5	2.0	12.5	13.0	1101.9	4.8	17.7	19.3	1101.9	17.7	96.9
Trial3_440	74	NA()	2.5	13.3461	1.6	1.7355	3.5	0.1681	3.0	0.88	1001.4	7.1	16.8	17.3	1021.8	6.4	13.5	14.0	1066.6	12.8	21.4	22.7	1066.6	21.4	93.9
Trial3_441	391	NA()	0.6	9.8807	0.8	3.9644	1.3	0.2842	1.0	0.78	1612.6	2.5	23.6	24.4	1626.9	2.0	15.1	15.9	1646.4	3.3	16.1	17.6	1646.4	16.1	97.9
Trial3_442	73	NA()	1.1	13.2772	1.6	1.8000	3.4	0.1734	3.0	0.88	1030.9	7.2	17.2	17.7	1045.5	6.4	13.7	14.2	1077.0	12.4	21.1	22.4	1077.0	21.1	95.7
Trial3_443	48	NA()	1.3	12.4368	1.9	2.0739	3.7	0.1872	3.2	0.86	1105.9	8.1	18.5	19.1	1140.2	7.5	14.7	15.3	1207.0	14.9	22.4	23.6	1207.0	22.4	91.6
Trial3_444	192	NA()	1.9	19.4330	2.6	0.2552	4.2	0.0360	3.3	0.78	227.9	1.5	4.0	4.1	230.8	2.7	4.7	4.8	261.4	26.2	32.7	33.8	227.9	4.0	NA
Trial3_445	121	NA()	0.7	11.3370	1.1	2.5391	2.8	0.2089	2.5	0.92	1222.8	7.0	19.6	20.2	1283.3	5.5	14.5	15.1	1387.0	7.8	18.1	19.5	1387.0	18.1	88.2
Trial3_446	16	NA()	0.8	17.6905	4.9	0.5638	5.9	0.0724	3.2	0.55	450.4	5.0	8.7	8.9	454.0	14.9	16.3	16.5	473.1	86.2	88.2	88.6	450.4	8.7	95.2
Trial3_447	131	NA()	2.4	12.3567	1.1	2.3087	2.7	0.2070	2.5	0.91	1212.8	6.8	19.4	20.0	1215.0	5.3	14.1	14.7	1219.7	8.2	18.6	20.1	1219.7	18.6	99.4
Trial3_448	261	NA()	0.9	8.1564	0.8	5.4355	1.4	0.3217	1.2	0.81	1798.0	5.3	26.3	27.2	1890.5	3.5	16.2	17.0	1994.4	4.1	15.7	17.1	1994.4	15.7	90.1
Trial3_449	27	NA()	1.4	13.0715	3.4	1.1378	4.6	0.1079	3.2	0.69	660.6	6.3	12.1	12.4	771.4	9.7	13.9	14.3	1108.2	29.9	34.4	35.2	660.6	12.1	59.6
Trial3_450	93	NA()	1.4	10.5219	1.1	3.1448	2.9	0.2401	2.7	0.92	1387.2	8.3	22.1	22.8	1443.7	6.2	15.5	16.2	1528.8	8.4	18.1	19.5	1528.8	18.1	90.7
Trial3_451	94	NA()	0.8	19.5246	3.6	0.2594	4.7	0.0367	3.1	0.66	232.7	2.0	4.3	4.4	234.2	4.4	5.9	6.0	250.6	43.9	48.1	48.9	232.7	4.3	NA
Trial3_452	24	NA()	3.7	9.3982	1.8	4.5810	3.7	0.3124	3.3	0.88	1752.5	11.4	27.7	28.5	1745.8	8.8	17.7	18.4	1738.7	13.7	20.7	21.9	1738.7	20.7	100.8
Trial3_453	103	NA()	6.3	4.3053	0.9	18.9602	1.8	0.5923	1.6	0.86	2998.7	8.1	40.2	41.6	3039.8	5.5	18.7	19.5	3067.8	7.2	15.4	16.6	3067.8	15.4	97.8
Trial3_454	441	NA()	1.1	19.4028	1.7	0.2784	3.2	0.0392	2.8	0.86	247.9	0.5	4.0	4.2	249.4	1.4	4.3	4.5	265.0	14.3	24.2	25.7	247.9	4.0	NA
Trial3_455	106	NA()	1.8	9.0373	0.9	4.4732	2.5	0.2933	2.4	0.93	1658.1	8.8	25.6	26.4	1726.0	6.0	16.5	17.2	1810.1	7.1	17.0	18.4	1810.1	17.0	91.6
Trial3_456	52	NA()	2.0	11.8834	1.7	2.2462	3.6	0.1937	3.2	0.88	1141.3	8.3	19.1	19.6	1195.6	7.3	14.9	15.5	1296.0	13.0	21.0	22.3	1296.0	21.0	88.1
Trial3_457	125	NA()	0.9	10.1053	0.9	3.8405	2.4	0.2816	2.2	0.93	1599.4	8.1	24.7	25.4	1601.3	5.5	15.9	16.6	1604.5	7.0	17.3	18.7	1604.5	17.3	99.7
Trial3_458	204	NA()	12.3	17.4242	1.7	0.5613	3.4	0.0710	2.9	0.86	442.0	2.6	7.5	7.7	452.4	3.2	7.5	7.8	506.6	13.9	23.3	24.8	442.0	7.5	87.2
Trial3_459	366	NA()	0.7	15.6052	1.3	0.5946	2.8	0.0673	2.5	0.89	420.0	1.2	6.8	7.0	473.8	2.1	7.3	7.6	744.3	9.6	20.4	21.9	420.0	6.8	56.4
Trial3_460	40	NA()	0.7	12.8277	2.1	1.8502	3.9	0.1722	3.3	0.84	1024.3	8.1	17.5	18.0	1063.5	8.0	14.6	15.1	1145.8	17.0	23.9	25.1	1145.8	23.9	89.4
Trial3_461	88	NA()	1.3	11.1872	1.2	2.9929	2.9	0.2429	2.7	0.92	1402.0	8.5	22.4	23.0	1405.8	6.2	15.4	16.0	1412.5	8.9	18.6	19.9	1412.5	18.6	99.3
Trial3_462	55	NA()	2.0	13.1347	1.8	1.8542	3.7	0.1767	3.2	0.86	1049.0	7.8	17.7	18.2	1064.9	7.3	14.2	14.7	1098.6	14.9	22.6	23.9	1098.6	22.6	95.5
Trial3_463	128	NA()	1.8	9.3220	0.8	4.7838	2.2	0.3236	2.0	0.93	1807.2	8.5	27.2	28.1	1782.1	5.5	16.5	17.2	1753.6	6.7	16.9	18.3	1753.6	16.9	103.1
Trial3_464	66	NA()	1.5	13.7633	1.8	1.5720	3.6	0.1570	3.1	0.87	940.0	7.0	16.0	16.4	959.2	6.6	13.2	13.7	1004.4	13.9	22.2	23.5	1004.4	22.2	93.6
Trial3_465	79	NA()	2.4	13.2141	1.6	1.7327	3.4	0.1661	3.0	0.89	990.8	7.0	16.6	17.1	1020.8	6.2	13.4	14.0	1086.5	12.0	20.8	22.2	1086.5	20.8	91.2
Trial3_466	171	NA()	3.0	13.3636	1.1	1.6320	2.7	0.1582	2.5	0.91	947.0	5.3	15.4	15.9	982.6	4.6	12.5	13.0	1064.0	8.4	19.0	20.5	1064.0	19.0	89.0
Trial3_467	309	NA()	2.3	12.4951	0.8	1.9999	1.9	0.1813	1.7	0.90	1074.2	3.4	16.6	17.2	1115.5	3.0	12.9	13.5	1197.8	5.5	17.6	19.2	1197.8	17.6	89.7
Trial3_468	205	NA()	0.8	19.1336	2.9	0.1810	4.4	0.0251	3.3	0.75	160.0	1.1	2.8	2.9	168.9	2.2	3.6	3.8	297.0	28.9	34.8	35.9	160.0	2.8	NA
Trial3_469	76	NA()	0.9	20.1236	3.7	0.2363	4.9	0.0345	3.2	0.65	218.6	2.1	4.1	4.2	215.4	4.7	5.9	6.0	180.6	51.7	55.4	56.1	218.6	4.1	NA
Trial3_470	180	NA()	2.1	12.4720	1.0	2.2372	2.4	0.2025	2.2	0.91	1188.5	6.0	18.8	19.4	1192.8	4.6	13.8	14.4	1201.4	7.1	18.2	19.7	1201.4	18.2	98.9
Trial3_471	223	NA()	1.9	19.1549	2.3	0.2864	3.9	0.0398	3.2	0.81	251.6	1.5	4.3	4.5	255.7	2.5	4.9	5.0	294.4	21.1	28.7	30.0	251.6	4.3	NA
Trial3_472	66	NA()	1.2	10.9541	1.3	3.0272	3.2	0.2406	2.9	0.91	1389.9	8.9	22.4	23.0	1414.5	6.7	15.6	16.3	1452.6	10.0	19.0	20.3	1452.6	19.0	95.7

Trial3_473	24	NA()	2.5	12.9682	2.7	1.8890	4.4	0.1777	3.4	0.78	1054.7	8.8	18.2	18.7	1077.2	9.8	15.7	16.2	1124.1	23.3	28.8	29.7	1124.1	28.8	93.8
Trial3_474	37	NA()	1.0	11.0860	1.8	2.8890	3.7	0.2324	3.2	0.87	1347.0	9.6	22.1	22.8	1379.0	8.1	16.1	16.7	1429.8	13.9	21.4	22.6	1429.8	21.4	94.2
Trial3_475	271	NA()	10.7	12.8488	0.9	1.8502	2.1	0.1725	1.9	0.91	1025.9	3.8	16.0	16.6	1063.5	3.3	12.6	13.2	1142.5	6.1	17.9	19.5	1142.5	17.9	89.8
Trial3_476	156	NA()	0.8	19.1441	2.8	0.2680	4.3	0.0372	3.3	0.76	235.6	1.7	4.2	4.3	241.1	3.1	5.0	5.2	295.7	28.3	34.3	35.4	235.6	4.2	NA
Trial3_477	335	NA()	7.1	4.5503	0.8	16.2797	1.3	0.5375	1.0	0.78	2773.0	3.9	35.4	36.8	2893.4	2.5	16.7	17.6	2979.0	3.1	12.0	13.5	2979.0	12.0	93.1
Trial3_478	30	NA()	1.4	13.1297	2.2	1.8553	3.9	0.1768	3.2	0.83	1049.2	8.8	17.5	18.0	1065.3	9.6	14.9	15.4	1099.4	22.8	27.0	28.0	1099.4	27.0	95.4
Trial3_479	36	NA()	4.3	9.3552	1.3	4.2888	3.2	0.2911	3.0	0.91	1647.2	10.9	25.2	26.0	1691.2	8.1	16.3	17.0	1747.1	11.7	17.6	19.0	1747.1	17.6	94.3
Trial3_480	75	NA()	2.7	12.6535	1.3	2.0494	3.1	0.1882	2.8	0.91	1111.4	7.9	17.8	18.4	1132.1	6.7	13.5	14.1	1172.9	11.9	18.6	20.1	1172.9	18.6	94.8
Trial3_481	110	NA()	1.8	13.2098	1.1	1.8373	2.8	0.1761	2.6	0.92	1045.7	7.0	16.6	17.2	1058.9	5.7	12.7	13.2	1087.2	9.7	17.4	19.0	1087.2	17.4	96.2
Trial3_482	48	NA()	2.2	12.6597	1.7	1.9114	3.5	0.1756	3.1	0.88	1042.8	8.2	17.2	17.7	1085.1	8.1	14.1	14.6	1171.9	17.2	22.3	23.6	1171.9	22.3	89.0
Trial3_483	52	NA()	1.9	11.7006	1.4	2.6265	3.2	0.2230	2.9	0.91	1297.6	9.3	20.6	21.2	1308.1	7.4	14.7	15.3	1326.1	12.1	18.5	19.9	1326.1	18.5	97.9
Trial3_484	39	NA()	1.9	13.0938	1.9	1.9084	3.7	0.1813	3.1	0.86	1074.2	8.6	17.7	18.2	1084.0	8.4	14.2	14.8	1104.8	18.3	23.3	24.5	1104.8	23.3	97.2
Trial3_485	60	NA()	2.5	11.1343	1.3	2.7781	3.1	0.2244	2.9	0.92	1305.3	9.1	20.6	21.2	1349.7	7.2	14.7	15.3	1421.5	10.9	17.6	19.0	1421.5	17.6	91.8
Trial3_486	85	NA()	1.6	12.4803	1.2	2.1203	3.0	0.1920	2.7	0.92	1132.2	7.8	18.0	18.6	1155.4	6.4	13.5	14.1	1200.1	10.6	17.7	19.3	1200.1	17.7	94.3
Trial3_487	39	NA()	1.5	13.3594	1.9	1.8181	3.7	0.1762	3.2	0.85	1046.4	8.5	17.3	17.9	1052.0	8.5	14.1	14.6	1064.6	19.0	23.9	25.1	1064.6	23.9	98.3
Trial3_488	71	NA()	2.5	13.5862	1.4	1.7280	3.3	0.1703	2.9	0.90	1014.0	7.6	16.5	17.0	1019.0	6.6	12.9	13.4	1030.6	12.8	19.4	20.9	1030.6	19.4	98.4
Trial3_489	88	NA()	1.2	5.3966	0.9	12.8324	2.1	0.5025	1.9	0.89	2624.5	9.6	35.0	36.3	2667.4	6.0	17.3	18.2	2700.9	7.6	14.1	15.5	2700.9	14.1	97.2
Trial3_490	169	NA()	2.7	13.8176	1.0	1.5459	2.6	0.1550	2.4	0.92	928.9	5.4	14.6	15.1	948.9	4.7	11.6	12.2	996.4	8.8	17.1	18.8	996.4	17.1	93.2
Trial3_491	124	NA()	1.4	4.4403	0.9	17.5375	1.6	0.5650	1.3	0.81	2887.4	8.1	37.3	38.7	2964.7	5.3	17.4	18.3	3018.3	6.9	13.5	14.8	3018.3	13.5	95.7
Trial3_492	29	NA()	0.8	10.6791	1.7	3.0223	3.6	0.2342	3.2	0.88	1356.4	10.4	21.8	22.4	1413.3	8.9	15.9	16.5	1500.9	15.2	20.4	21.7	1500.9	20.4	90.4
Trial3_493	19	NA()	1.6	9.3883	1.8	4.2227	3.7	0.2877	3.2	0.87	1629.8	12.0	25.5	26.3	1678.4	9.9	17.2	17.9	1740.6	15.9	20.7	21.8	1740.6	20.7	93.6
Trial3_494	44	NA()	1.3	10.8595	1.4	3.1507	3.3	0.2483	3.0	0.91	1429.5	10.2	22.5	23.2	1445.2	7.9	15.4	16.1	1469.1	12.3	18.4	19.8	1469.1	18.4	97.3
Trial3_495	273	NA()	2.5	13.4237	0.8	1.7977	1.9	0.1751	1.7	0.92	1040.1	4.3	15.6	16.2	1044.6	3.5	11.8	12.4	1054.9	6.2	15.8	17.6	1054.9	15.8	98.6
Trial3_496	210	NA()	1.3	16.3048	1.2	0.8634	2.8	0.1021	2.5	0.91	627.0	3.7	10.0	10.4	632.0	3.7	8.9	9.3	650.9	10.6	18.8	20.5	627.0	10.0	96.3
Trial3_497	443	NA()	2.5	8.5222	0.8	5.3952	1.3	0.3336	1.0	0.78	1855.9	2.1	25.3	26.3	1884.1	1.6	14.8	15.7	1916.1	2.6	13.2	14.9	1916.1	13.2	96.9
Trial3_498	37	NA()	2.9	12.6758	1.8	2.1116	3.6	0.1942	3.1	0.87	1144.2	9.3	18.9	19.4	1152.6	8.4	14.5	15.1	1169.4	16.4	21.7	23.0	1169.4	21.7	97.8
Trial3_499	268	NA()	0.9	17.4051	1.2	0.6350	2.8	0.0802	2.5	0.90	497.3	2.5	7.9	8.2	499.2	2.9	7.4	7.7	509.0	11.4	19.5	21.3	497.3	7.9	97.7
Trial3_500	101	NA()	0.7	12.5867	1.1	2.1546	2.8	0.1968	2.6	0.92	1158.0	8.0	18.4	19.0	1166.5	6.2	13.5	14.1	1183.4	9.6	17.2	18.8	1183.4	17.2	97.9
Trial3_501	42	NA()	1.9	13.5767	1.9	1.6790	3.7	0.1654	3.2	0.85	986.7	8.3	16.5	17.0	1000.6	8.4	13.8	14.3	1032.0	19.5	24.4	25.5	1032.0	24.4	95.6
Trial3_502	38	NA()	0.6	8.5095	1.2	5.3417	3.1	0.3298	2.9	0.93	1837.5	11.6	27.6	28.4	1875.6	8.0	16.8	17.5	1918.8	10.6	16.7	18.1	1918.8	16.7	95.8
Trial3_503	126	NA()	1.1	10.0839	0.8	3.8663	2.3	0.2829	2.1	0.94	1605.9	8.7	23.9	24.7	1606.7	5.8	15.1	15.8	1608.5	7.1	15.2	16.8	1608.5	15.2	99.8
Trial3_504	213	NA()	3.4	18.9979	2.0	0.2808	3.7	0.0387	3.0	0.83	244.8	1.6	4.1	4.2	251.3	2.6	4.6	4.8	313.2	21.8	27.3	28.6	244.8	4.1	NA
Trial3_505	71	NA()	1.6	9.5450	1.0	4.4137	2.7	0.3057	2.6	0.94	1719.4	10.3	25.8	26.6	1714.9	6.9	15.8	16.6	1710.2	8.7	15.9	17.4	1710.2	15.9	100.5
Trial3_506	41	NA()	1.1	9.1756	1.2	4.8204	3.1	0.3209	2.9	0.92	1794.3	11.4	27.0	27.9	1788.5	8.0	16.5	17.2	1782.5	11.0	17.1	18.5	1782.5	17.1	100.7
Trial3_507	37	NA()	2.2	13.7866	2.1	1.5926	3.8	0.1593	3.2	0.84	952.9	8.2	16.1	16.6	967.3	8.8	13.9	14.4	1000.9	21.6	26.0	27.2	1000.9	26.0	95.2
Trial3_508	48	NA()	1.0	12.9526	1.6	1.9673	3.5	0.1849	3.1	0.88	1093.7	8.7	18.0	18.5	1104.4	7.7	13.9	14.5	1126.5	14.7	20.6	21.9	1126.5	20.6	97.1
Trial3_509	87	NA()	2.9	13.6397	1.3	1.6805	3.1	0.1663	2.8	0.91	991.8	7.4	16.2	16.7	1001.2	6.3	12.6	13.2	1022.7	11.6	18.7	20.2	1022.7	18.7	97.0
Trial3_510	40	NA()	0.5	18.9011	5.9	0.1924	6.7	0.0264	3.3	0.48	167.9	2.1	3.3	3.4	178.7	8.6	9.0	9.1	324.8	115.5	116.7	117.0	167.9	3.3	NA
Trial3_511	94	NA()	1.1	18.0402	2.3	0.5220	3.9	0.0683	3.2	0.81	426.1	3.8	7.5	7.7	426.5	5.2	7.9	8.2	429.7	26.4	30.9	32.0	426.1	7.5	99.2
Trial3_512	118	NA()	1.0	16.8147	1.7	0.7436	3.4	0.0907	3.0	0.87	559.8	4.5	9.5	9.8	564.5	4.9	8.9	9.3	584.4	16.6	22.8	24.3	559.8	9.5	95.8
Trial3_513	283	NA()	6.1	18.4838	1.4	0.4549	3.0	0.0610	2.7	0.89	381.8	1.9	6.1	6.3	380.7	2.5	6.0	6.3	375.3	13.7	21.2	22.9	381.8	6.1	NA
Trial3_514	150	NA()	1.7	19.0154	2.3	0.3044	3.9	0.0420	3.2	0.81	265.2	2.1	4.6	4.7	269.8	3.3	5.2	5.4	311.1	25.6	30.4	31.6	265.2	4.6	NA
Trial3_515	50	NA()	0.7	10.6579	1.3	3.2392	3.2	0.2505	2.9	0.91	1441.0	10.0	22.5	23.2	1466.6	7.6	15.4	16.0	1504.6	11.4	17.8	19.2	1504.6	17.8	95.8
Trial3_516	51	NA()	1.8	16.6633	2.6	0.6833	4.2	0.0826	3.3	0.78	511.7	4.8	9.1	9.3	528.7	6.9	9.9	10.2	604.0	29.4	33.3	34.3	511.7	9.1	84.7
Trial3_517	35	NA()	1.0	8.8108	1.3	4.9914	3.2	0.3191	3.0	0.92	1785.4	11.5	27.0	27.8	1817.9	8.2	16.7	17.4	1856.1	11.3	17.3	18.6	1856.1	17.3	96.2
Trial3_518	213	NA()	1.7	9.1267	0.8	4.8358	1.6	0.3202	1.3	0.85	1790.9	6.6	25.3	26.2	1791.1	4.2	15.1	15.9	1792.2	5.1	14.1	15.7	1792.2	14.1	99.9
Trial3_519	155	NA()	2.1	13.1552	1.0	1.9203	2.5	0.1833	2.3	0.93	1085.0	6.2	16.8	17.4	1088.2	4.9	12.5	13.1	1095.5	8.1	16.6	18.2	1095.5	16.6	99.0
Trial3_520	48	NA()	1.7	13.7478	1.8	1.7079	3.6	0.1704	3.1	0.87	1014.1	7.9	16.7	17.2	1011.5	7.7	13.5	14.0	1006.7	17.4	22.7	24.0	1006.7	22.7	100.7
Trial3_521	105	NA()	2.6	13.4231	1.2	1.6778	3.0	0.1634	2.7	0.91	975.7	6.6	15.6	16.1	1000.2	5.8	12.4	12.9	1055.0	10.8	18.1	19.7	1055.0	18.1	92.5
Trial3_522	463	NA()	0.1	18.4452	1.2	0.3892	2.7	0.0521	2.4	0.89	327.3	0.4	5.0	5.2	333.8	1.5	5.1	5.4	380.0	11.5	19.9	21.6	327.3	5.0	NA
Trial3_523	326	NA()	2.0	12.1930	0.9	1.1167	2.3	0.0988	2.1	0.93	607.3	1.9	9.3	9.6	761.4	2.6	9.6	10.1	1245.9	7.0	15.7	17.4	1245.9	15.7	48.7
Trial3_524	50	NA()	1.3	10.9568	1.3	3.1401	3.2	0.2496	2.9	0.91	1436.6	9.4	22.2	23.0	1442.6	7.2	15.1	15.8	1452.2	11.2	17.7	19.1	1452.2	17.7	98.9
Trial3_525	50	NA()	1.2	19.6143	4.6	0.2588	5.6	0.0368	3.2	0.57	233.2	2.4	4.3	4.4	233.7	7.6	8.4	8.5	240.1	80.8	82.5	82.9	233.2	4.3	NA
Trial3_526	100	NA()	1.6	13.4893	1.3	1.4725	3.1	0.1441	2.8	0.91	868.0	6.0	14.0	14.5	919.2	5.6	11.8	12.4	1045.1	11.6	18.6	20.1	1045.1	18.6	83.1
Trial3_527	167	NA()	2.2	12.1427	0.9	2.1807	2.3	0.1921	2.2	0.93	1132.9	5.9	17.3	17.9	1174.9	4.7	12.9	13.5	1253.9	7.2	15.8	17.5	1253.9	15.8	90.4
Trial3_528	193	NA()	1.4	16.1709	1.2	0.8425	2.9	0.0989	2.6	0.90	607.7	3.4	9.7	10.0	620.5	3.7	8.8	9.2	668.6	11.3	19.1	20.8	607.7	9.7	90.9
Trial3_529	90	NA()	2.0	8.9331	0.9	4.1089	2.6	0.2663	2.5	0.94	1522.1	8.9	23.0	23.8	1656.1	6.4	15.4	16.2	1831.2	7.7	15.2	16.7	1831.2	15.2	83.1
Trial3_530	327	NA()	2.1	12.5602	0.6	2.1363	1.5	0.1947	1.4	0.91	1146.8	3.2	16.7	17.4	1160.6	2.8	12.2	12.9	1187.5	5.0	15.1	16.9	1187.5	15.1	96.6
Trial3_531	58	NA()	1.1	10.6596	1.2	3.2279	3.1	0.2497	2.8	0.92	1436.8	9.4	22.2	23.0	1463.9	7.1	15.1	15.8	1504.3	10.2	17.0	18.5	1504.3	17.0	95.5
Trial3_532	36	NA()	1.0	5.2741	0.9	13.1456	2.8	0.5031	2.6	0.95	2627.0	11.1	35.5	36.8	2690.2	7.1	17.7	18.6	2738.7	8.9	14.8	16.1	2738.7	14.8	95.9
Trial3_533	164	NA()	0.9	9.7743	0.7	3.9541	1.9	0.2804	1.8	0.94	1593.5	7.1	23.2	24.0	1624.8	4.8	14.8	15.5	1666.4	5.9	14.6	16.2	1666.4	14.6	95.6
Trial3_534	152	NA()	1.6	9.1049	0.9	4.8711	1.9	0.3218	1.7	0.89	1798.5	7.7	25.7	26.6	1797.3	5.0	15.3	16.1	1796.6	6.0	14.4	16.0	1796.6	14.4	100.1
Trial3_535	16	NA()	1.3	12.3834	2.9	2.0507	4.3	0.1843	3.1	0.74	1090.2	9.6	18.4	18.9	1132.5	12.5	17.1	17.6	1215.5	30.7	33.8	34.6	1215.5	33.8	89.7
Trial3_536	135	NA()	2.1	12.4147	1.0	2.2547	2.5	0.2031	2.4	0.93	1192.0	6.9	18.3	19.0	1198.3	5.3	13.2	13.8	1210.5	8.2	16.4	18.0	1210.5	16.4	98.5
Trial3_537	76	NA()	0.9	5.2425	0.9	13.5428	2.2	0.5152	2.0	0.91	2678.6	9.7	35.6	36.9	2718.3	6.0	17.4	18.2	2748.6	7.6	14.1	15.4	2748.6	14.1	97.5
Trial3_538	29	NA()	1.5	13.6167	2.3	1.6929	4.0	0.1673	3.2	0.82	997.0	8.4	16.7	17.2	1005.9	9.6	14.6	15.1	1026.1	24.2	28.3	29.3	1026.1	28.3	97.2
Trial3_539	230	NA()	0.4	19.3492	2.1	0.2532	3.7	0.0355	3.0	0.83	225.1	1.3	3.7	3.8	229.1	2.4	4.3	4.5	271.3	23.5	28.8	30.0	225.1	3.7	NA
Trial3_540	173	NA()	2.0	9.2514	0.9	4.7747	1.8	0.3205	1.6	0.88	1792.2	7.5	25.6	26.5	1780.5	4.8	15.2	16.0	1767.5	5.8	14.4	16.0	1767.5	14.4	101.4
Trial3_541	104	NA()	0.5	5.4516	0.9	12.9770	1.9	0.5133	1.6	0.87	2670.8	9.1	35.4	36.7	2678.0	5.6	17.2	18.0	2684.2	7.1	13.9	15.2	2684.2	13.9	99.5
Trial3_542	50	NA()	2.5	9.3220	1.1	4.5495	3.0	0.3077	2.8	0.93	1729.5	10.6	26.0	26.8	1740.1	7.3	16.1	16.8	1753.6	9.8	16.5	17.9	1753.6	16.5	98.6
Trial3_543	74	NA()	2.0	13.6337	1.5	1.6358	3.3	0.1618	2.9	0.90	966.9	7.5	15.9	16.4	984.1	6.6	12.7	13.2	1023.6	13.0	19.6	21.0	1023.6	19.6	94.5
Trial3_544	58	NA()	1.6	13.2680	1.6	1.7138	3.4	0.1650	3.0	0.89	984.4	7.9	16.3	16.8	1013.7	7.1	13.2	13.7	1078.4	14.1	20.2	21.6	1078.4	20.2	91.3
Trial3_545	127	NA()	1.4	8.7714	0.7	5.4315	2.0	0.3457	1.9	0.94	1914.0	9.2	27.5	28.4	1889.8	5.8	15.9	16.6	1864.2	6.8	14.7	16.2	1864.2	14.7	102.7
Trial3_546	90	NA()	1.1	17.9156	2.3	0.5676	3.9	0.0738	3.2	0.81	458.9	4.0	8.0	8.2	456.5	5.5	8.4	8.7	445.1	26.6	31.1	32.2	458.9	8.0	103.1
Trial3_547	112	NA()	1.9	17.5101	1.9	0.5851	3.7	0.0743	3.1	0.85	462.3	3.9	8.0	8.2	467.8	4.8	8.0	8.3	495.8	20.5	26.0	27.3	462.3	8.0	93.2
Trial3_548	49	NA()	1.4	11.0102	1.3	3.1312	3.2	0.2502	2.9	0.91	1439.3	10.3	22.7	23.4	1440.4	7.8	15.4	16.0	1442.9	12.0	18.2	19.6	1442.9	18.2	99.7
Trial3_549	72	NA()	1.1	19.2512	3.5	0.2698	4.8	0.0377	3.2	0.67	238.5	2.4	4.4	4.5	242.5	4.6	5.9	6.1	283.0	43.2	46.2	47.0	238.5	4.4	NA
Trial3_550	114	NA()	1.1	18.4119	2.3	0.4085	3.9	0.0546	3.2	0.81	342.6	3.0	6.0	6.2	347.8	4.2	6.6	6.8	384.0	24.5	29.4	30.6	342.6	6.0	NA
Trial3_551	62	NA()	4.1	13.0704	1.4	1.9899	3.3	0.1887	3.0	0.90	1114.4	8.7	18.2	18.8	1112.1	7.2	13.7	14.2	1108.4	12.9	19.3	20.7	1108.4	19.3	100.5
Trial3_552	22	NA()	1.9	13.2250	2.4	1.9060	4.1	0.1829	3.3	0.81	1082.8	9.7	18.4	18.9	1083.2	10.4	15.5	16.0	1084.9	24.4	28.4	29.4	1084.9	28.4	99.8
Trial3_553	100	NA()	1.7	16.6983	1.8	0.7390	3.6	0.0895	3.1	0.86	552.8	4.7	9.5	9.8	561.8	5.2	9.1	9.4	599.4	17.8	23.7	25.1	552.8	9.5	92.2
Trial3_554	93	NA()	0.9	11.6663	1.0	2.6556	2.8	0.2248	2.6	0.93	1307.2	8.8	20.5	21.2	1316.2	6.5	14.3	14.9	1331.8	9.2	16.7	18.3	1331.8	16.7	98.2
Trial3_555	37	NA()	1.9	13.5250	1.9	1.8372	3.7	0.1803	3.2	0.86	1068.6	9.1	17.9	18.4	1058.9	8.3	14.1	14.6	1039.7	17.7	22.9	24.1	1039.7	22.9	102.8
Trial3_556	22	NA()	1.6	6.0611	1.2	10.5944	3.2	0.4659	3.0	0.93	2465.7	13.1	34.6	35.8	2488.2	8.5	18.1	18.9	2507.4	10.9	16.3	17.5	2507.4	16.3	98.3
Trial3_557	24	NA()	1.8	13.1380	2.4	1.8723	4.1	0.1785	3.3	0.81	1058.7	9.5	18.0	18.5	1071.4	10.7	15.6	16.1	1098.1	25.9	29.6	30.6	1098.1	29.6	96.4
Trial3_558	63	NA()	1.2	17.4731	2.4	0.6876	4.0	0.0872	3.2	0.81	538.8	5.0	9.5	9.8	531.4	6.3	9.5	9.8	500.5	25.8	30.3	31.4	538.8	9.5	107.7
Trial3_559	206	NA()	0.7	13.0819	0.8	1.9064	2.2	0.1810	2.0	0.93	1072.2	6.0	16.6	17.1	1083.3	4.7	12.4	13.0	1106.6	7.1	16.1	17.8	1106.6	16.1	96.9
Trial3_560	103	NA()	0.8	13.1223	1.2	1.8481	2.9	0.1760	2.7	0.92	1044.9	7.7	16.9	17.5	1062.8	6.2	12.9	13.5	1100.5	10.2	17.7	19.3	1100.5	17.7	95.0
Trial3_561	49	NA()	1.9	15.9195	2.3	0.9793	4.0	0.1131	3.2	0.82	690.8	6.5	12.1	12.5	693.2	7.5	11.5	11.8	702.0	23.9	28.4	29.5	690.8	12.1	98.4
Trial3_562	364	NA()	2.3	13.3435	0.6	1.8654	1.5	0.1806	1.3	0.90	1070.3	3.2	15.7	16.3	1068.9	2.7	11.7	12.3	1067.0	5.1	15.4	17.2	1067.0	15.4	100.3
Trial3_563	159	NA()	1.3	13.2431	1.0	1.8091	2.6	0.1738	2.4	0.93	1033.2	6.6	16.3	16.9	1048.8	5.3	12.4	13.0	1082.1	8.3	16.7	18.3	1082.1	16.7	95.5
Trial3_564	346	NA()	1.6	13.5993	0.7	1.6923	1.7	0.1670	1.5	0.91	995.5	3.3	14.8	15.4	1005.6	2.9	11.4	12.0	1028.7	5.5	15.6	17.4	1028.7	15.6	96.8
Trial3_565	343	NA()	2.2	13.8291	0.7	1.6544	1.7	0.1660	1.5	0.91	990.1	3.4	14.7	15.3	991.2	2.9	11.3	11.9	994.7	5.7	15.7	17.5	994.7	15.7	99.5
Trial3_566	114	NA()	1.0	13.2377	1.1	1.8575	2.8	0.1784	2.6	0.92	1058.3	7.5	17.0	17.6	1066.1	6.0	12.9	13.4	1082.9	9.7	17.4	19.1	1082.9	17.4	97.7
Trial3_567	27	NA()	1.7	13.7423	2.4	1.6900	4.0	0.1685	3.3	0.81	1003.9	9.2	17.2	17.7	1004.8	10.2	15.0	15.4	1007.5	25.3	29.3	30.3	1007.5	29.3	99.6
Trial3_568	100	NA()	1.5	12.7234	1.1	2.1046	2.9	0.1943	2.6	0.92	1144.6	8.2	18.3	18.9	1150.3	6.3	13.4	14.0	1161.9	9.6	17.2	18.8	1161.9	17.2	98.5
Trial3_569	87	NA()	1.4	11.1367	1.0	2.9558	2.8	0.2389	2.6	0.93	1380.7	9.4	21.6	22.3	1396.3	6.8	14.7	15.4	1421.1	9.2	16.6	18.1	1421.1	16.6	97.2
Trial3_570	94	NA()	2.6	20.0543	3.6	0.1815	4.8	0.0264	3.2	0.66	168.0	1.7	3.1	3.2	169.3	3.5	4.4	4.5	188.7	46.1	49.0	49.8	168.0	3.1	NA
Trial3_571	13	NA()	1.4	13.6656	3.5	1.8044	4.8	0.1789	3.2	0.67	1061.1	10.5	18.6	19.1	1047.0	14.1	18.1	18.5	1018.8	38.0	40.7	41.4	1018.8	40.7	104.1
Trial3_572	93	NA()	1.7	7.7331	0.8	6.4217	2.3	0.3603	2.1	0.94	1983.7	10.5	28.7	29.7	2035.2	6.6	16.5	17.3	2088.7	7.5	14.7	16.2	2088.7	14.7	95.0
Trial3_573	77	NA()	1.2	16.9593	2.1	0.7373	3.8	0.0907	3.2	0.83	559.9	5.1	9.8	10.1	560.8	6.1	9.6	10.0	565.8	23.2	28.0	29.2	559.9	9.8	99.0
Trial3_574	198	NA()	0.6	16.9847	1.3	0.7207	3.0	0.0888	2.7	0.90	548.6	3.6	9.0	9.3	551.1	3.8	8.2	8.6	562.5	12.0	19.8	21.4	548.6	9.0	97.5
Trial3_575	80	NA()	2.5	18.8202	3.5	0.3292	4.9	0.0450	3.3	0.69	283.5	2.8	5.2	5.3	289.0	5.4	6.9	7.0	334.6	42.4	45.4	46.2	283.5	5.2	NA
Trial3_576	151	NA()	1.6	11.2187	0.8	2.9338	2.2	0.2388	2.1	0.93	1380.6	8.0	21.0	21.7	1390.7	5.6	14.2	14.9	1407.1	7.0	15.5	17.1	1407.1	15.5	98.1
Trial3_577	136	NA()	3.0	13.7264	1.1	1.6706	2.8	0.1664	2.5	0.92	992.2	6.8	15.9	16.4	997.4	5.6	12.3	12.8	1009.8	9.6	17.5	19.1	1009.8	17.5	98.3
Trial3_578	80	NA()	1.0	7.3857	0.8	7.4530	2.3	0.3994	2.2	0.94	2166.3	10.8	30.7	31.8	2167.4	6.7	16.8	17.6	2169.1	8.0	14.9	16.3	2169.1	14.9	99.9
Trial3_579	29	NA()	0.6	7.8394	1.2	6.4412	3.2	0.3664	3.0	0.92	2012.4	12.5	29.8	30.8	2037.9	8.4	17.3	18.1	2064.6	11.0	16.8	18.1	2064.6	16.8	97.5
Trial3_580	81	NA()	1.9	13.5091	1.3	1.7978	3.1	0.1762	2.8	0.91	1046.3	7.9	17.1	17.6	1044.7	6.5	13.0	13.5	1042.1	11.6	18.6	20.1	1042.1	18.6	100.4
Trial3_581	152	NA()	1.4	18.5834	1.8	0.4853	3.5	0.0654	3.0	0.86	408.6	3.2	7.0	7.2	401.7	3.8	6.9	7.1	363.2	18.6	24.7	26.2	408.6	7.0	112.5
Trial3_582	27	NA()	1.6	7.8212	1.3	6.5382	3.3	0.3710	3.0	0.92	2034.3	12.7	30.1	31.1	2051.1	8.6	17.4	18.2	2068.7	11.5	17.1	18.4	2068.7	17.1	98.3
Trial3_583	142	NA()	1.2	13.6271	1.0	1.8027	2.6	0.1782	2.4	0.92	1057.4	6.9	16.7	17.3	1046.4	5.5	12.5	13.1	1024.5	8.9	17.1	18.8	1024.5	17.1	103.2
Trial3_584	63	NA()	1.1	5.2698	1.0	13.3494	2.4	0.5104	2.2	0.92	2658.5	10.9	35.8	37.0	2704.7	6.8	17.6	18.5	2740.1	8.5	14.6	15.9	2740.1	14.6	97.0
Trial3_585	99	NA()	1.9	13.7943	1.2	1.7020	3.0	0.1704	2.7	0.91	1014.1	7.5	16.5	17.0	1009.3	6.1	12.6	13.2	999.8	10.8	18.2	19.8	999.8	18.2	101.4
Trial3_586	114	NA()	2.4	13.7740	1.2	1.5377	3.0	0.1537	2.7	0.91	921.6	6.8	15.0	15.5	945.6	5.8	12.1	12.6	1002.8	10.7	18.1	19.7	1002.8	18.1	91.9
Trial3_587	263	NA()	1.7	19.6367	1.9	0.2486	3.5	0.0354	3.0	0.85	224.4	1.3	3.7	3.8	225.4	2.1	4.1	4.2	237.5	19.3	25.5	27.0	224.4	3.7	NA
Trial3_588	97	NA()	2.1	9.1570	0.8	4.4723	2.5	0.2972	2.3	0.94	1677.2	9.9	25.1	25.9	1725.8	6.6	15.7	16.5	1786.2	7.7	15.2	16.7	1786.2	15.2	93.9
Trial3_589	40	NA()	0.9	13.7081	1.9	1.6617	3.7	0.1653	3.2	0.86	986.1	8.5	16.7	17.1	994.0	8.1	13.6	14.1	1012.5	17.6	22.9	24.2	1012.5	22.9	97.4
Trial3_590	137	NA()	1.8	9.1660	0.7	4.8220	2.0	0.3207	1.9	0.94	1793.1	9.1	26.1	27.0	1788.7	5.8	15.6	16.3	1784.4	6.7	14.8	16.3	1784.4	14.8	100.5
Trial3_591	117	NA()	0.7	11.1091	0.9	2.8310	2.6	0.2282	2.4	0.93	1325.0	8.4	20.5	21.2	1363.8	6.1	14.3	14.9	1425.9	7.9	15.9	17.5	1425.9	15.9	92.9
Trial3_592	28	NA()	0.8	10.7038	1.7	3.2666	3.6	0.2537	3.1	0.88	1457.5	11.3	23.3	24.0	1473.1	9.1	16.2	16.8	1496.5	15.0	20.3	21.5	1496.5	20.3	97.4
Trial3_593	52	NA()	2.1	13.9297	1.7	1.6034	3.5	0.1621	3.1	0.88	968.2	8.1	16.2	16.7	971.5	7.3	13.0	13.5	980.0	15.4	21.3	22.7	980.0	21.3	98.8
Trial3_594	43	NA()	1.6	11.1428	1.4	2.9344	3.3	0.2373	3.0	0.90	1372.4	10.2	21.9	22.5	1390.8	8.1	15.4	16.0	1420.1	13.0	18.9	20.3	1420.1	18.9	96.6
Trial3_595	94	NA()	1.6	8.9986	0.8	4.6642	2.5	0.3045	2.3	0.94	1713.8	9.9	25.5	26.4	1760.8	6.5	15.8	16.5	1817.9	7.6	15.2	16.7	1817.9	15.2	94.3
Trial3_596	56	NA()	0.5	5.5296	0.8	12.3056	2.4	0.4937	2.3	0.95	2586.8	11.0	35.1	36.3	2628.0	6.9	17.6	18.4	2660.6	8.6	14.7	16.0	2660.6	14.7	97.2
Trial3_597	150	NA()	2.0	7.9213	0.9	5.8768	1.9	0.3378	1.7	0.89	1876.0	8.7	26.9	27.8	1957.8	5.5	15.9	16.7	2046.3	6.1	14.1	15.7	2046.3	14.1	91.7
Trial3_598	49	NA()	1.4	5.3687	0.8	12.7498	2.5	0.4967	2.4	0.95	2599.4	11.2	35.3	36.5	2661.3	7.0	17.7	18.5	2709.5	8.8	14.8	16.1	2709.5	14.8	95.9
Trial3_599	31	NA()	2.9	13.8689	2.2	1.6436	3.9	0.1654	3.2	0.83	986.7	8.7	16.8	17.2	987.1	8.9	14.0	14.5	988.9	21.1	25.7	26.8	988.9	25.7	99.8
Trial3_600	100	NA()	1.0	5.5184	0.9	11.9377	2.0	0.4780	1.8	0.89	2518.6	9.9	34.1	35.3	2599.5	6.1	17.2	18.1	2664.0	7.4	14.0	15.4	2664.0	14.0	94.5
Trial3_601	117	NA()	13.1	16.5825	1.7	0.7336	3.4	0.0883	3.0	0.87	545.3	4.4	9.3	9.6	558.6	4.8	8.8	9.2	614.5	16.2	22.4	23.9	545.3	9.3	88.7
Trial3_602	825	NA()	9.0	17.3102	1.0	0.3501	2.3	0.0440	2.1	0.91	277.4	1.7	4.6	4.7	304.8	1.9	4.9	5.2	521.0	8.5	18.0	19.8	277.4	4.6	NA
Trial3_603	184	NA()	0.9	5.4289	0.9	12.9042	1.3	0.5083	1.0	0.74	2649.4	7.8	34.8	36.1	2672.7	4.8	16.9	17.8	2691.0	6.0	13.4	14.8	2691.0	13.4	98.5
Trial3_604	49	NA()	0.3	5.7388	0.8	11.4373	2.6	0.4763	2.5	0.95	2510.9	11.4	34.4	35.6	2559.5	7.1	17.6	18.4	2598.9	8.8	14.9	16.2	2598.9	14.9	96.6
Trial3_605	230	NA()	0.8	13.2933	0.8	1.7095	2.2	0.1649	2.0	0.92	983.9	4.9	15.1	15.6	1012.1	4.1	11.8	12.3	1074.6	6.9	16.0	17.8	1074.6	16.0	91.6
Trial3_606	74	NA()	1.7	13.0883	1.3	1.9925	3.2	0.1892	2.9	0.91	1117.2	8.3	18.1	18.6	1113.0	6.8	13.5	14.0	1105.7	11.8	18.7	20.1	1105.7	18.7	101.0
Trial3_607	198	NA()	0.8	9.2830	0.9	4.4082	1.8	0.2969	1.5	0.87	1676.0	7.3	24.2	25.1	1713.9	4.8	15.0	15.8	1761.2	5.5	14.3	15.9	1761.2	14.3	95.2
Trial3_608	17	NA()	2.1	8.8328	1.8	5.0339	3.7	0.3226	3.2	0.87	1802.5	12.7	27.7	28.5	1825.0	9.9	17.6	18.3	1851.6	15.1	20.0	21.1	1851.6	20.0	97.3
Trial3_609	133	NA()	3.1	16.7647	1.6	0.7353	3.3	0.0894	2.9	0.88	552.3	4.0	9.2	9.5	559.7	4.4	8.6	9.0	590.9	14.7	21.4	23.0	552.3	9.2	93.5
Trial3_610	27	NA()	0.8	6.9268	1.2	7.4015	3.2	0.3720	3.0	0.93	2038.8	12.2	29.9	30.9	2161.2	8.3	17.5	18.3	2280.2	10.6	16.3	17.6	2280.2	16.3	89.4

Trial3_611	203	NA()	2.5	10.7538	0.7	2.9483	2.0	0.2301	1.8	0.93	1334.8	6.2	19.8	20.5	1394.4	4.5	13.8	14.5	1487.7	5.8	14.9	16.5	1487.7	14.9	89.7
Trial3_612	206	NA()	2.4	13.3918	0.9	1.7255	2.3	0.1677	2.1	0.92	999.3	5.1	15.4	15.9	1018.1	4.3	11.9	12.4	1059.7	7.4	16.3	18.0	1059.7	16.3	94.3
Trial3_613	391	NA()	2.1	18.3984	1.3	0.4053	2.8	0.0541	2.5	0.89	339.7	0.9	5.3	5.5	345.5	1.8	5.3	5.6	385.7	12.0	20.2	21.9	339.7	5.3	NA
Trial3_614	100	NA()	1.4	9.7962	0.9	3.5988	2.6	0.2558	2.4	0.94	1468.3	8.8	22.4	23.1	1549.3	6.3	15.1	15.7	1662.3	7.6	15.4	16.9	1662.3	15.4	88.3
Trial3_615	68	NA()	1.4	16.1706	2.1	0.8055	3.8	0.0945	3.2	0.83	582.1	5.1	10.1	10.4	599.9	6.1	9.9	10.2	668.6	21.0	26.1	27.3	582.1	10.1	87.1
Trial3_616	33	NA()	1.1	6.1302	1.0	10.9773	2.9	0.4883	2.7	0.94	2563.2	12.0	35.2	36.4	2521.2	7.6	17.7	18.5	2488.3	9.9	15.7	16.9	2488.3	15.7	103.0
Trial3_617	69	NA()	1.3	12.0475	1.3	2.4151	3.1	0.2111	2.8	0.91	1234.8	8.7	19.6	20.2	1247.1	6.9	14.2	14.8	1269.3	11.1	17.9	19.4	1269.3	17.9	97.3
Trial3_618	53	NA()	1.6	9.5620	1.1	4.4395	3.0	0.3080	2.7	0.93	1730.9	10.6	26.0	26.9	1719.7	7.3	16.0	16.7	1706.9	9.8	16.5	17.9	1706.9	16.5	101.4
Trial3_619	55	NA()	1.5	17.5856	2.8	0.5685	4.3	0.0725	3.3	0.77	451.5	4.2	8.0	8.3	457.1	6.2	8.8	9.1	486.3	30.0	34.0	35.0	451.5	8.0	92.8
Trial3_620	75	NA()	1.3	13.2774	1.4	1.8227	3.2	0.1756	2.9	0.90	1042.9	7.7	16.9	17.4	1053.7	6.7	13.1	13.7	1077.0	12.6	19.2	20.7	1077.0	19.2	96.8
Trial3_621	72	NA()	0.8	5.5353	0.9	11.4120	2.4	0.4584	2.2	0.92	2432.3	10.5	33.4	34.5	2557.4	6.4	17.3	18.1	2658.9	7.5	14.1	15.5	2658.9	14.1	91.5
Trial3_622	337	NA()	3.7	13.7909	0.7	1.6020	1.7	0.1603	1.6	0.91	958.5	2.9	14.2	14.7	971.0	2.7	11.1	11.7	1000.3	5.8	15.8	17.6	1000.3	15.8	95.8
Trial3_623	61	NA()	2.5	13.3712	1.6	1.7793	3.4	0.1726	3.0	0.89	1026.6	7.9	16.8	17.3	1037.9	7.2	13.3	13.8	1062.8	14.5	20.5	21.9	1062.8	20.5	96.6
Trial3_624	243	NA()	1.0	19.4791	2.3	0.1856	3.9	0.0262	3.1	0.80	166.9	1.0	2.8	2.9	172.8	1.9	3.4	3.5	256.0	24.4	29.5	30.8	166.9	2.8	NA
Trial3_625	118	NA()	2.2	13.1748	1.1	1.8787	2.8	0.1796	2.6	0.92	1064.8	7.1	16.9	17.4	1073.6	5.7	12.8	13.3	1092.5	9.5	17.3	18.9	1092.5	17.3	97.5
Trial3_626	150	NA()	1.7	12.5780	0.9	2.1005	2.5	0.1917	2.3	0.93	1130.6	6.8	17.6	18.1	1149.0	5.2	13.0	13.6	1184.7	7.8	16.2	17.9	1184.7	16.2	95.4
Trial3_627	117	NA()	1.2	8.9740	0.7	4.9895	2.2	0.3249	2.1	0.94	1813.6	9.3	26.4	27.3	1817.5	5.9	15.7	16.5	1822.9	7.0	14.8	16.4	1822.9	14.8	99.5
Trial3_628	38	NA()	1.2	5.3924	0.9	12.5043	2.8	0.4893	2.6	0.95	2567.4	11.7	35.1	36.3	2643.0	7.3	17.8	18.6	2702.2	9.0	14.9	16.2	2702.2	14.9	95.0
Trial3_629	82	NA()	1.3	16.8968	1.9	0.8077	3.7	0.0990	3.1	0.85	608.7	5.1	10.4	10.7	601.2	5.8	9.7	10.1	573.8	20.4	25.7	27.0	608.7	10.4	106.1
Trial3_630	17	NA()	0.6	5.1834	1.2	13.3786	3.3	0.5032	3.0	0.93	2627.4	13.4	36.3	37.5	2706.7	8.6	18.4	19.2	2767.2	11.0	16.2	17.3	2767.2	16.2	94.9
Trial3_631	152	NA()	1.2	10.9955	0.8	3.0793	2.2	0.2457	2.1	0.93	1416.1	7.5	21.2	22.0	1427.6	5.3	14.2	14.9	1445.5	6.8	15.3	16.9	1445.5	15.3	98.0
Trial3_632	113	NA()	3.9	9.1048	0.8	4.6875	2.3	0.3097	2.1	0.94	1739.1	9.1	25.5	26.4	1765.0	5.9	15.6	16.3	1796.6	7.0	14.9	16.4	1796.6	14.9	96.8
Trial3_633	217	NA()	1.7	10.1731	0.8	3.6578	1.8	0.2700	1.6	0.88	1540.8	6.4	22.3	23.2	1562.2	4.3	14.4	15.1	1592.1	5.3	14.5	16.1	1592.1	14.5	96.8
Trial3_634	74	NA()	1.2	19.6729	3.9	0.2675	5.0	0.0382	3.2	0.63	241.6	2.3	4.4	4.5	240.7	5.7	6.8	6.9	233.2	57.3	59.7	60.4	241.6	4.4	NA
Trial3_635	78	NA()	2.3	13.6028	1.4	1.7255	3.2	0.1703	2.9	0.90	1013.8	7.5	16.5	17.0	1018.1	6.4	12.8	13.3	1028.1	12.1	18.9	20.5	1028.1	18.9	98.6
Trial3_636	98	NA()	1.3	13.7162	1.2	1.7531	3.0	0.1745	2.7	0.91	1036.7	7.2	16.6	17.2	1028.3	6.0	12.7	13.2	1011.3	11.0	18.3	19.8	1011.3	18.3	102.5
Trial3_637	78	NA()	1.3	11.4098	1.1	2.6621	2.9	0.2204	2.7	0.92	1283.9	8.7	20.2	20.8	1318.0	6.7	14.4	15.0	1374.7	10.0	17.1	18.6	1374.7	17.1	93.4
Trial3_638	115	NA()	2.3	13.9258	1.2	1.6962	2.9	0.1714	2.6	0.91	1019.8	7.0	16.3	16.8	1007.1	5.7	12.4	13.0	980.5	10.5	18.0	19.6	980.5	18.0	104.0
Trial3_639	651	NA()	2.3	13.7402	0.8	1.7265	1.3	0.1721	1.0	0.78	1023.8	2.4	15.0	15.6	1018.4	2.1	11.3	11.9	1007.8	4.0	15.2	17.0	1007.8	15.2	101.6
Trial3_640	153	NA()	2.7	9.2629	0.7	4.4949	2.0	0.3021	1.8	0.94	1701.7	8.6	24.9	25.8	1730.0	5.6	15.4	16.1	1765.2	6.4	14.6	16.2	1765.2	14.6	96.4
Trial3_641	54	NA()	1.0	5.4745	0.8	12.5462	2.5	0.4984	2.4	0.95	2606.8	11.3	35.4	36.6	2646.2	7.0	17.7	18.5	2677.2	8.8	14.8	16.1	2677.2	14.8	97.4
Trial3_642	111	NA()	1.5	11.3568	0.9	2.8286	2.6	0.2331	2.4	0.93	1350.7	8.6	20.9	21.6	1363.2	6.2	14.3	15.0	1383.6	8.3	16.1	17.7	1383.6	16.1	97.6
Trial3_643	85	NA()	2.5	12.8235	1.3	1.9486	3.1	0.1813	2.8	0.91	1074.1	8.1	17.4	18.0	1097.9	6.6	13.3	13.9	1146.4	11.0	18.0	19.6	1146.4	18.0	93.7
Trial3_644	146	NA()	1.7	16.3023	1.4	0.8616	3.2	0.1019	2.8	0.89	625.6	4.6	10.4	10.7	631.0	4.6	9.3	9.7	651.2	12.9	20.1	21.8	625.6	10.4	96.1
Trial3_645	35	NA()	3.1	12.9919	1.9	1.9251	3.7	0.1815	3.2	0.86	1075.0	9.3	18.0	18.6	1089.8	8.7	14.4	14.9	1120.5	18.1	23.1	24.3	1120.5	23.1	95.9
Trial3_646	231	NA()	13.1	17.8339	1.4	0.5351	3.1	0.0692	2.7	0.89	431.6	2.7	7.0	7.3	435.2	3.1	6.8	7.2	455.3	13.6	21.0	22.6	431.6	7.0	94.8
Trial3_647	27	NA()	2.5	12.7142	2.2	1.9528	3.9	0.1802	3.3	0.83	1067.8	9.5	18.1	18.6	1099.4	9.7	15.1	15.6	1163.4	21.3	25.6	26.7	1163.4	25.6	91.8
Trial3_648	222	NA()	1.5	19.3702	1.9	0.2930	3.6	0.0412	3.0	0.84	260.1	1.7	4.4	4.5	260.9	2.6	4.7	4.9	268.8	20.8	26.6	28.0	260.1	4.4	NA
Trial3_649	16	NA()	0.6	9.7862	2.0	4.0132	3.8	0.2850	3.3	0.85	1616.4	12.7	25.7	26.5	1636.9	10.8	17.7	18.3	1664.2	18.3	22.6	23.7	1664.2	22.6	97.1
Trial3_650	256	NA()	2.0	12.6689	0.7	2.0328	1.9	0.1869	1.8	0.92	1104.4	5.1	16.7	17.3	1126.6	4.0	12.4	13.0	1170.4	6.0	15.5	17.3	1170.4	15.5	94.4
Trial3_651	246	NA()	4.1	13.4604	0.8	1.7494	2.1	0.1709	1.9	0.92	1016.9	5.1	15.6	16.1	1027.0	4.1	11.9	12.4	1049.4	6.7	16.0	17.8	1049.4	16.0	96.9
Trial3_652	55	NA()	2.4	12.5720	1.5	2.1278	3.4	0.1941	3.0	0.90	1143.6	9.2	18.8	19.4	1157.9	7.7	14.2	14.7	1185.7	13.5	19.7	21.1	1185.7	19.7	96.4
Trial3_653	65	NA()	1.7	12.9291	1.4	1.9746	3.3	0.1852	2.9	0.90	1095.6	8.7	18.0	18.5	1106.9	7.3	13.7	14.2	1130.1	12.8	19.2	20.7	1130.1	19.2	96.9
Trial3_654	84	NA()	2.3	13.8751	1.4	1.6107	3.2	0.1622	2.9	0.90	968.8	7.7	16.0	16.5	974.4	6.6	12.6	13.2	988.0	12.6	19.3	20.8	988.0	19.3	98.1
Trial3_655	110	NA()	2.7	13.1443	1.1	2.0371	2.8	0.1943	2.6	0.92	1144.6	7.9	18.2	18.8	1128.0	6.1	13.2	13.8	1097.1	9.5	17.3	18.9	1097.1	17.3	104.3
Trial3_656	56	NA()	1.6	13.0518	1.6	1.9075	3.4	0.1807	3.0	0.89	1070.5	8.6	17.6	18.2	1083.7	7.6	13.8	14.3	1111.2	14.8	20.6	22.0	1111.2	20.6	96.3
Trial3_657	68	NA()	2.1	12.4565	1.3	2.1961	3.2	0.1985	2.9	0.91	1167.2	8.7	18.8	19.4	1179.8	7.1	13.9	14.5	1203.8	11.9	18.5	20.0	1203.8	18.5	97.0
Trial3_658	160	NA()	1.3	10.9283	0.8	3.0111	2.2	0.2388	2.0	0.93	1380.3	7.5	20.8	21.5	1410.4	5.3	14.2	14.9	1457.1	6.7	15.3	16.9	1457.1	15.3	94.7
Trial3_659	51	NA()	2.1	8.7498	1.1	4.9794	2.9	0.3161	2.7	0.93	1770.8	11.1	26.6	27.5	1815.8	7.5	16.4	17.1	1868.7	9.4	16.1	17.5	1868.7	16.1	94.8
Trial3_660	29	NA()	0.6	5.6414	1.0	11.6546	3.0	0.4771	2.8	0.94	2514.5	12.2	34.8	36.0	2577.1	7.7	17.9	18.7	2627.4	9.7	15.4	16.7	2627.4	15.4	95.7
Trial3_661	35	NA()	1.7	13.1994	2.0	1.8713	3.7	0.1792	3.2	0.85	1062.7	8.8	17.7	18.2	1071.0	8.7	14.3	14.8	1088.8	19.0	23.9	25.1	1088.8	23.9	97.6
Trial3_662	45	NA()	1.1	12.3646	1.6	2.3015	3.4	0.2065	3.0	0.89	1210.0	9.3	19.6	20.2	1212.7	8.0	14.6	15.1	1218.4	14.8	20.5	21.8	1218.4	20.5	99.3
Trial3_663	22	NA()	2.2	10.3023	1.9	3.3560	3.7	0.2509	3.2	0.87	1443.0	11.2	23.1	23.8	1494.2	9.8	16.6	17.3	1568.5	16.9	21.6	22.8	1568.5	21.6	92.0
Trial3_664	199	NA()	3.7	12.7396	0.8	2.1801	2.1	0.2015	2.0	0.93	1183.5	6.0	17.9	18.6	1174.7	4.5	12.8	13.4	1159.4	6.9	15.9	17.6	1159.4	15.9	102.1
Trial3_665	86	NA()	0.8	5.5011	0.9	12.8712	2.1	0.5138	1.9	0.89	2672.7	9.8	35.6	36.9	2670.3	6.1	17.3	18.2	2669.2	7.7	14.2	15.6	2669.2	14.2	100.1
Trial3_666	121	NA()	1.3	8.0356	0.9	6.5931	2.0	0.3844	1.8	0.90	2096.9	9.5	29.5	30.6	2058.4	5.8	16.2	17.0	2020.9	6.9	14.5	16.0	2020.9	14.5	103.8
Trial3_667	32	NA()	1.0	11.9335	1.8	2.5360	3.6	0.2196	3.2	0.87	1279.7	10.1	20.8	21.4	1282.4	9.1	15.5	16.1	1287.8	17.5	22.4	23.6	1287.8	22.4	99.4
Trial3_668	63	NA()	0.9	5.1571	0.9	14.0885	2.3	0.5272	2.1	0.91	2729.6	10.5	36.3	37.6	2755.7	6.6	17.6	18.4	2775.6	8.3	14.5	15.8	2775.6	14.5	98.3
Trial3_669	54	NA()	0.9	13.6818	1.6	1.7203	3.5	0.1708	3.1	0.88	1016.4	8.1	16.8	17.3	1016.1	7.3	13.2	13.8	1016.4	15.0	20.9	22.3	1016.4	20.9	100.0
Trial3_670	21	NA()	1.2	13.0174	2.5	1.8879	4.2	0.1783	3.3	0.80	1057.8	9.5	18.0	18.5	1076.9	10.3	15.4	15.9	1116.5	24.3	28.2	29.3	1116.5	28.2	94.7
Trial3_671	103	NA()	1.7	10.9900	0.9	3.1654	2.6	0.2524	2.4	0.93	1450.9	8.7	22.1	22.8	1448.8	6.1	14.6	15.3	1446.4	8.1	16.0	17.5	1446.4	16.0	100.3
Trial3_672	398	NA()	3.0	19.7144	1.6	0.2557	3.1	0.0366	2.7	0.87	231.6	0.6	3.6	3.8	231.2	1.6	3.9	4.1	228.3	16.3	23.3	24.9	231.6	3.6	NA
Trial3_673	199	NA()	1.0	16.6817	1.3	0.7584	3.0	0.0918	2.7	0.90	566.1	3.4	9.1	9.4	573.1	3.6	8.4	8.8	601.6	11.7	19.5	21.2	566.1	9.1	94.1
Trial3_674	57	NA()	0.7	10.8059	1.2	3.2972	3.0	0.2585	2.8	0.92	1482.3	9.8	22.9	23.7	1480.4	7.2	15.2	15.9	1478.5	10.4	17.2	18.6	1478.5	17.2	100.3
Trial3_675	236	NA()	1.4	16.4810	1.2	0.8005	2.7	0.0957	2.5	0.91	589.3	3.1	9.3	9.7	597.1	3.4	8.5	8.9	627.7	10.7	18.9	20.6	589.3	9.3	93.9
Trial3_676	444	NA()	1.9	19.7824	1.5	0.2444	3.1	0.0351	2.7	0.87	222.3	0.5	3.4	3.6	222.0	1.5	3.8	3.9	220.4	16.9	23.7	25.3	222.3	3.4	NA
Trial3_677	57	NA()	1.0	13.2944	1.6	1.7510	3.4	0.1689	3.0	0.89	1006.1	8.0	16.6	17.1	1027.5	7.1	13.2	13.8	1074.4	14.0	20.2	21.6	1074.4	20.2	93.6
Trial3_678	20	NA()	2.6	12.1984	2.3	2.1996	4.0	0.1947	3.3	0.82	1146.7	10.0	19.2	19.8	1180.9	10.2	15.8	16.3	1245.0	21.7	25.8	26.9	1245.0	25.8	92.1
Trial3_679	62	NA()	1.4	13.4065	1.6	1.7248	3.4	0.1678	3.0	0.89	999.9	7.9	16.5	17.0	1017.8	7.2	13.2	13.7	1057.5	15.0	20.9	22.2	1057.5	20.9	94.5
Trial3_680	64	NA()	1.4	13.3022	1.5	1.8552	3.3	0.1791	3.0	0.90	1061.9	8.3	17.4	18.0	1065.3	7.1	13.4	13.9	1073.2	13.1	19.5	21.0	1073.2	19.5	98.9
Trial3_681	294	NA()	2.0	13.7046	0.7	1.6803	1.9	0.1671	1.7	0.92	996.1	4.0	15.0	15.5	1001.1	3.4	11.5	12.1	1013.1	6.1	15.8	17.6	1013.1	15.8	98.3
Trial3_682	86	NA()	1.3	5.4468	0.9	12.4742	2.1	0.4930	1.9	0.89	2583.6	10.1	34.8	36.0	2640.8	6.3	17.4	18.2	2685.6	7.9	14.3	15.7	2685.6	14.3	96.2
Trial3_683	61	NA()	1.4	19.0773	4.2	0.2676	5.2	0.0370	3.2	0.61	234.5	2.5	4.4	4.5	240.8	6.6	7.6	7.7	303.7	65.8	67.9	68.4	234.5	4.4	NA
Trial3_684	143	NA()	1.7	13.4620	1.0	1.8076	2.6	0.1766	2.4	0.92	1048.2	6.6	16.5	17.0	1048.2	5.3	12.4	13.0	1049.2	8.8	17.0	18.7	1049.2	17.0	99.9
Trial3_685	54	NA()	1.1	16.4219	2.5	0.7637	4.1	0.0910	3.3	0.80	561.4	5.2	9.9	10.2	576.1	7.0	10.3	10.6	635.5	27.0	31.1	32.2	561.4	9.9	88.3
Trial3_686	47	NA()	1.2	13.4191	1.8	1.7115	3.6	0.1667	3.1	0.87	993.6	8.2	16.6	17.1	1012.9	7.9	13.6	14.1	1055.6	17.1	22.4	23.7	1055.6	22.4	94.1
Trial3_687	97	NA()	2.4	9.0917	0.8	4.3466	2.5	0.2867	2.3	0.94	1625.2	9.5	24.4	25.2	1702.2	6.4	15.6	16.3	1799.2	7.6	15.2	16.7	1799.2	15.2	90.3
Trial3_688	92	NA()	1.0	12.4438	1.1	2.2161	2.9	0.2001	2.7	0.92	1175.8	8.1	18.7	19.3	1186.1	6.4	13.7	14.3	1205.9	10.3	17.5	19.0	1205.9	17.5	97.5
Trial3_689	260	NA()	1.2	16.3919	1.1	0.7888	2.7	0.0938	2.4	0.91	578.1	2.9	9.1	9.4	590.5	3.1	8.3	8.7	639.5	9.8	18.4	20.1	578.1	9.1	90.4
Trial3_690	93	NA()	0.8	13.1258	1.2	1.8390	3.0	0.1751	2.8	0.91	1040.4	7.6	16.8	17.4	1059.5	6.2	12.9	13.5	1099.9	10.6	17.9	19.4	1099.9	17.9	94.6
Trial3_691	37	NA()	1.7	5.2273	0.9	13.4631	2.8	0.5106	2.6	0.95	2659.4	11.7	36.0	37.3	2712.7	7.4	17.9	18.7	2753.4	9.4	15.1	16.4	2753.4	15.1	96.6
Trial3_692	612	NA()	3.6	13.3976	0.8	1.6110	1.3	0.1566	1.0	0.78	937.9	1.5	13.7	14.3	974.5	1.7	10.9	11.5	1058.8	4.2	15.1	17.0	1058.8	15.1	88.6
Trial3_693	110	NA()	2.9	13.3306	1.2	1.7092	2.9	0.1653	2.7	0.92	986.3	7.0	15.9	16.4	1012.0	5.9	12.5	13.0	1068.9	10.1	17.6	19.2	1068.9	17.6	92.3
Trial3_694	40	NA()	2.9	13.5713	1.9	1.7079	3.7	0.1682	3.2	0.85	1002.1	8.5	16.8	17.3	1011.5	8.5	13.9	14.4	1032.8	19.2	24.1	25.3	1032.8	24.1	97.0
Trial3_695	104	NA()	1.6	9.1206	0.8	4.7537	2.3	0.3146	2.2	0.94	1763.3	9.8	26.0	26.9	1776.8	6.3	15.8	16.5	1793.4	7.4	15.1	16.6	1793.4	15.1	98.3
Trial3_696	31	NA()	0.8	13.6171	2.1	1.7211	3.9	0.1701	3.2	0.84	1012.4	9.0	17.2	17.7	1016.4	8.9	14.2	14.7	1026.0	20.2	24.9	26.1	1026.0	24.9	98.7
Trial3_697	30	NA()	1.4	7.7130	1.2	6.7835	3.2	0.3796	2.9	0.93	2074.6	12.7	30.5	31.5	2083.6	8.4	17.4	18.1	2093.2	10.9	16.7	18.0	2093.2	16.7	99.1
Trial3_698	65	NA()	0.9	5.3940	0.9	12.5941	2.4	0.4929	2.2	0.92	2583.3	11.1	35.1	36.3	2649.8	6.8	17.6	18.4	2701.7	8.3	14.5	15.8	2701.7	14.5	95.6
Trial3_699	228	NA()	2.8	12.2697	0.7	2.4011	1.9	0.2138	1.8	0.93	1248.9	6.0	18.7	19.4	1242.9	4.4	13.1	13.8	1233.6	6.2	15.4	17.2	1233.6	15.4	101.2
Trial3_700	63	NA()	2.4	9.4774	1.0	4.2845	2.9	0.2946	2.7	0.93	1664.7	10.8	25.4	26.2	1690.4	7.4	16.0	16.7	1723.3	9.3	16.2	17.6	1723.3	16.2	96.6
Trial3_701	48	NA()	2.7	10.2728	1.3	3.2675	3.2	0.2436	3.0	0.91	1405.2	10.5	22.3	23.0	1473.3	8.0	15.6	16.2	1573.8	11.4	17.7	19.1	1573.8	17.7	89.3
Trial3_702	20	NA()	1.0	11.9835	2.2	2.6361	3.9	0.2292	3.3	0.83	1330.4	11.2	21.9	22.5	1310.8	10.7	16.6	17.1	1279.7	21.5	25.7	26.8	1279.7	25.7	104.0
Trial3_703	49	NA()	1.1	9.2207	1.1	4.8312	3.0	0.3232	2.8	0.93	1805.5	11.6	27.2	28.1	1790.4	7.8	16.4	17.2	1773.6	10.2	16.6	18.0	1773.6	16.6	101.8
Trial3_704	48	NA()	1.5	7.7058	1.0	6.8470	2.8	0.3828	2.7	0.94	2089.5	12.0	30.4	31.4	2091.8	7.6	17.0	17.8	2094.9	9.3	15.7	17.1	2094.9	15.7	99.7
Trial3_705	174	NA()	3.6	10.7058	0.7	3.2938	2.0	0.2559	1.9	0.93	1468.7	7.7	21.9	22.7	1479.6	5.2	14.4	15.1	1496.1	6.3	15.0	16.7	1496.1	15.0	98.2
Trial3_706	300	NA()	5.6	13.5710	0.7	1.7188	1.8	0.1693	1.7	0.92	1008.0	4.0	15.1	15.7	1015.6	3.3	11.5	12.1	1032.9	6.0	15.8	17.6	1032.9	15.8	97.6
Trial3_707	240	NA()	2.8	13.3577	0.8	1.9103	2.0	0.1852	1.9	0.92	1095.1	5.0	16.5	17.1	1084.7	4.0	12.1	12.7	1064.9	6.5	15.9	17.6	1064.9	15.9	102.8
Trial3_708	281	NA()	1.8	10.8039	0.8	3.0281	1.6	0.2374	1.4	0.86	1373.1	4.9	19.9	20.7	1414.7	3.6	13.6	14.3	1478.9	4.8	14.5	16.2	1478.9	14.5	92.8
Trial3_709	98	NA()	1.6	8.9970	0.8	5.0828	2.4	0.3318	2.2	0.94	1847.2	9.9	27.0	27.9	1833.2	6.3	15.9	16.7	1818.3	7.5	15.1	16.6	1818.3	15.1	101.6
Trial3_710	100	NA()	1.3	13.6119	1.2	1.7276	3.0	0.1706	2.7	0.91	1015.6	7.2	16.4	16.9	1018.9	6.0	12.6	13.1	1026.8	10.8	18.2	19.7	1026.8	18.2	98.9
Trial3_711	135	NA()	2.9	13.7390	1.1	1.7678	2.7	0.1762	2.5	0.92	1046.3	6.7	16.5	17.1	1033.7	5.4	12.4	13.0	1008.0	9.2	17.3	18.9	1008.0	17.3	103.8
Trial3_712	160	NA()	1.4	12.5666	0.9	2.1014	2.4	0.1916	2.3	0.93	1130.1	6.6	17.5	18.1	1149.2	5.1	12.9	13.5	1186.5	7.7	16.2	17.9	1186.5	16.2	95.2
Trial3_713	160	NA()	1.5	10.9530	0.8	3.1308	2.1	0.2488	2.0	0.93	1432.4	7.5	21.5	22.2	1440.3	5.3	14.3	15.0	1452.8	6.8	15.3	16.9	1452.8	15.3	98.6
Trial3_714	62	NA()	1.6	10.9456	1.2	3.2019	3.0	0.2543	2.8	0.92	1460.6	9.9	22.7	23.4	1457.6	7.2	15.1	15.8	1454.1	10.2	17.1	18.6	1454.1	17.1	100.4
Trial3_715	419	NA()	0.6	17.5354	1.1	0.5369	2.5	0.0683	2.2	0.90	426.0	1.0	6.5	6.8	436.4	1.8	6.4	6.7	492.6	9.6	18.6	20.4	426.0	6.5	86.5
Trial3_716	1002	NA()	14.4	13.7670	0.8	1.5126	1.6	0.1511	1.3	0.85	907.1	7.5	15.2	15.6	935.5	5.6	11.9	12.4	1003.8	4.6	15.3	17.2	1003.8	15.3	90.4
Trial3_717	31	NA()	1.1	5.4484	1.0	13.1112	2.9	0.5183	2.7	0.94	2692.1	12.0	36.4	37.7	2687.7	7.6	17.9	18.8	2685.1	9.9	15.5	16.7	2685.1	15.5	100.3
Trial3_718	23	NA()	1.5	13.1112	2.5	1.7679	4.1	0.1682	3.3	0.80	1002.1	9.1	17.1	17.6	1033.8	10.4	15.3	15.7	1102.2	25.4	29.2	30.2	1102.2	29.2	90.9
Trial3_719	35	NA()	1.6	13.0016	2.0	1.8554	3.8	0.1750	3.2	0.85	1039.8	8.8	17.4	17.9	1065.3	8.7	14.3	14.8	1119.0	18.8	23.7	24.9	1119.0	23.7	92.9
Trial3_720	62	NA()	1.0	5.2280	0.9	13.2807	2.4	0.5038	2.2	0.92	2630.1	10.5	35.4	36.6	2699.8	6.6	17.5	18.4	2753.2	8.2	14.4	15.7	2753.2	14.4	95.5
Trial3_721	89	NA()	2.3	11.8882	1.1	2.2940	2.9	0.1979	2.7	0.92	1163.9	8.0	18.5	19.1	1210.4	6.4	13.8	14.4	1295.2	9.8	17.1	18.6	1295.2	17.1	89.9
Trial3_722	96	NA()	2.3	13.4873	1.3	1.6124	3.1	0.1578	2.8	0.91	944.5	6.9	15.4	15.9	975.1	6.1	12.4	12.9	1045.4	11.6	18.6	20.1	1045.4	18.6	90.4
Trial3_723	34	NA()	1.5	12.9202	2.1	1.6224	3.9	0.1521	3.2	0.84	912.7	8.0	15.5	16.0	978.9	8.8	14.0	14.4	1131.5	20.8	25.3	26.4	1131.5	25.3	80.7
Trial3_724	56	NA()	2.3	11.9750	1.4	2.4740	3.2	0.2150	2.9	0.91	1255.2	9.2	20.1	20.7	1264.5	7.3	14.4	15.0	1281.1	11.7	18.3	19.7	1281.1	18.3	98.0
Trial3_725	64	NA()	1.2	18.9009	3.8	0.2641	5.0	0.0362	3.2	0.64	229.4	2.3	4.2	4.3	238.0	5.5	6.6	6.8	324.9	54.5	56.9	57.5	229.4	4.2	NA
Trial3_726	215	NA()	3.2	13.2867	0.9	1.6593	2.3	0.1600	2.1	0.93	956.6	5.0	14.8	15.3	993.1	4.2	11.7	12.3	1075.6	7.2	16.1	17.9	1075.6	16.1	88.9
Trial3_727	69	NA()	2.7	13.3781	1.5	1.6141	3.3	0.1567	3.0	0.89	938.3	7.4	15.5	16.0	975.7	6.8	12.7	13.2	1061.8	13.5	19.8	21.3	1061.8	19.8	88.4
Trial3_728	110	NA()	2.0	13.0859	1.2	1.8161	2.9	0.1724	2.7	0.92	1025.5	6.9	16.4	16.9	1051.3	5.8	12.7	13.2	1106.0	10.0	17.5	19.1	1106.0	17.5	92.7
Trial3_729	547	NA()	4.2	19.2037	1.3	0.2678	2.8	0.0373	2.5	0.88	236.1	0.4	3.6	3.8	240.9	1.3	3.9	4.1	288.6	13.0	21.0	22.7	236.1	3.6	NA
Trial3_730	64	NA()	2.3	16.7688	2.3	0.6711	4.0	0.0817	3.2	0.81	506.0	4.6	8.9	9.1	521.4	5.9	9.2	9.5	590.3	23.9	28.6	29.8	506.0	8.9	85.7
Trial3_731	168	NA()	1.4	8.3694	0.9	5.3394	1.8	0.3242	1.6	0.88	1810.4	7.9	25.9	26.8	1875.2	5.1	15.6	16.4	1948.5	5.7	14.1	15.7	1948.5	14.1	92.9
Trial3_732	162	NA()	2.3	10.7849	0.7	3.2826	2.1	0.2569	1.9	0.93	1473.8	7.5	21.9	22.7	1476.9	5.2	14.4	15.1	1482.2	6.5	15.1	16.8	1482.2	15.1	99.4
Trial3_733	124	NA()	1.5	10.7580	0.8	3.2205	2.4	0.2514	2.2	0.93	1445.7	8.2	21.8	22.6	1462.1	5.8	14.5	15.2	1486.9	7.5	15.6	17.2	1486.9	15.6	97.2
Trial3_734	69	NA()	1.8	13.0941	1.4	1.9290	3.2	0.1833	2.9	0.90	1084.8	8.0	17.5	18.1	1091.2	6.8	13.3	13.9	1104.8	12.3	18.9	20.4	1104.8	18.9	98.2
Trial3_735	32	NA()	2.1	11.2449	1.7	2.8235	3.5	0.2304	3.1	0.88	1336.5	10.1	21.4	22.1	1361.8	8.6	15.5	16.1	1402.6	14.8	20.3	21.5	1402.6	20.3	95.3
Trial3_736	81	NA()	1.4	11.4600	1.1	2.7067	2.9	0.2251	2.7	0.92	1308.6	8.8	20.5	21.2	1330.3	6.6	14.4	15.0	1366.2	9.5	16.8	18.4	1366.2	16.8	95.8
Trial3_737	161	NA()	1.3	8.8535	0.9	4.9419	1.9	0.3175	1.7	0.89	1777.4	8.2	25.6	26.5	1809.4	5.3	15.5	16.2	1847.4	6.0	14.4	15.9	1847.4	14.4	96.2
Trial3_738	23	NA()	1.6	12.6849	2.3	2.0638	4.0	0.1900	3.3	0.82	1121.1	9.9	18.9	19.4	1136.9	10.1	15.5	16.0	1167.9	22.1	26.3	27.4	1167.9	26.3	96.0
Trial3_739	36	NA()	2.2	13.1812	2.0	1.8365	3.7	0.1756	3.2	0.85	1043.1	9.0	17.5	18.0	1058.6	8.9	14.4	14.9	1091.5	19.7	24.4	25.6	1091.5	24.4	95.6
Trial3_740	35	NA()	1.6	17.4618	3.5	0.6543	4.8	0.0829	3.2	0.67	513.4	5.3	9.4	9.6	511.1	8.7	11.1	11.4	501.9	41.6	44.5	45.3	513.4	9.4	102.3
Trial3_741	124	NA()	1.6	14.4880	1.2	1.4584	2.9	0.1533	2.7	0.91	919.5	6.5	14.9	15.4	913.4	5.6	11.8	12.3	899.4	10.7	18.4	20.0	899.4	18.4	102.2
Trial3_742	203	NA()	9.3	13.7918	0.9	1.6398	2.3	0.1641	2.2	0.92	979.5	5.5	15.2	15.8	985.6	4.5	11.8	12.3	1000.2	7.6	16.5	18.2	1000.2	16.5	97.9
Trial3_743	25	NA()	1.4	13.1059	2.4	1.7962	4.1	0.1708	3.3	0.81	1016.6	9.2	17.4	17.9	1044.1	10.4	15.3	15.8	1103.0	25.1	28.9	29.9	1103.0	28.9	92.2
Trial3_744	88	NA()	1.8	13.2892	1.3	1.6287	3.1	0.1571	2.9	0.91	940.4	7.2	15.5	15.9	981.4	6.3	12.5	13.1	1075.2	11.3	18.4	19.9	1075.2	18.4	87.5
Trial3_745	152	NA()	2.5	10.2085	0.8	3.4691	2.2	0.2570	2.0	0.94	1474.3	8.1	22.1	22.9	1520.2	5.6	14.7	15.4	1585.6	6.6	15.0	16.6	1585.6	15.0	93.0
Trial3_746	179	NA()	2.9	13.5171	0.9	1.7943	2.4	0.1760	2.2	0.93	1045.0	6.1	16.3	16.8	1043.4	4.8	12.2	12.8	1040.9	7.7	16.5	18.2	1040.9	16.5	100.4
Trial3_747	35	NA()	1.3	6.0333	1.0	10.4175	2.9	0.4560	2.7	0.94	2422.1	12.2	33.9	35.0	2472.6	7.8	17.7	18.5	2515.1	9.7	15.5	16.8	2515.1	15.5	96.3
Trial3_748	103	NA()	1.3	11.6653	1.0	2.3911	2.8	0.2024	2.6	0.93	1188.1	8.1	18.8	19.4	1239.9	6.3	13.8	14.4	1332.0	8.8	16.5	18.1	1332.0	16.5	89.2

Trial3_749	46	NA()	2.7	14.0923	1.9	1.4727	3.7	0.1506	3.2	0.86	904.2	7.9	15.4	15.8	919.2	7.9	13.1	13.6	956.3	18.7	23.8	25.0	956.3	23.8	94.6
Trial3_750	90	NA()	1.7	18.3588	2.4	0.4604	4.0	0.0613	3.2	0.80	383.7	3.5	6.8	7.0	384.6	5.0	7.5	7.7	390.5	27.9	32.3	33.4	383.7	6.8	NA
Trial3_751	159	NA()	1.7	5.4462	0.9	12.5655	1.5	0.4966	1.2	0.79	2599.0	8.5	34.5	35.8	2647.6	5.3	17.0	17.9	2685.8	6.5	13.6	15.0	2685.8	13.6	96.8
Trial3_752	77	NA()	0.8	11.0276	1.1	3.1613	2.8	0.2530	2.6	0.93	1453.7	9.8	22.6	23.3	1447.8	6.9	15.0	15.6	1439.9	9.3	16.6	18.1	1439.9	16.6	101.0
Trial3_753	154	NA()	15.3	13.3231	1.1	1.7811	2.4	0.1722	2.1	0.89	1024.1	7.6	16.2	16.8	1038.6	5.8	12.9	13.5	1070.1	7.8	20.1	21.5	1070.1	21.7	95.7
Trial3_754	91	NA()	3.7	13.5324	1.4	1.7895	3.0	0.1757	2.6	0.88	1043.5	9.1	17.2	17.7	1041.6	7.0	13.5	14.0	1038.6	10.3	21.3	22.6	1038.6	28.2	100.5
Trial3_755	170	NA()	2.0	10.9792	0.9	3.0005	1.9	0.2390	1.7	0.90	1381.7	8.7	20.8	21.5	1407.7	5.9	14.8	15.4	1448.3	6.6	18.7	20.1	1448.3	16.2	95.4
Trial3_756	497	NA()	1.4	19.6053	1.5	0.2719	2.8	0.0387	2.3	0.84	244.7	0.4	3.7	3.8	244.2	1.2	4.0	4.2	241.1	12.4	24.6	26.1	244.7	5.6	NA
Trial3_757	64	NA()	0.8	9.7032	1.2	3.8844	2.8	0.2735	2.5	0.91	1558.5	12.0	24.2	24.9	1610.4	8.1	16.5	17.1	1679.9	9.3	19.4	20.6	1679.9	21.8	92.8
Trial3_758	101	NA()	1.4	13.7125	1.4	1.7018	2.9	0.1693	2.6	0.88	1008.4	8.6	16.6	17.1	1009.2	6.7	13.2	13.7	1011.9	9.8	21.1	22.5	1011.9	27.5	99.7
Trial3_759	151	NA()	2.6	13.3782	1.1	1.8182	2.4	0.1765	2.2	0.89	1047.8	7.8	16.6	17.1	1052.0	5.9	13.0	13.6	1061.8	7.9	20.2	21.6	1061.8	22.2	98.7
Trial3_760	93	NA()	2.2	9.1621	1.0	4.5992	2.3	0.3058	2.1	0.91	1719.8	11.6	25.6	26.5	1749.1	7.4	16.6	17.3	1785.2	8.2	18.7	20.0	1785.2	17.4	96.3
Trial3_761	43	NA()	1.1	9.9248	1.4	3.7446	3.1	0.2697	2.8	0.89	1539.1	12.6	24.3	25.0	1580.9	8.8	16.7	17.3	1638.1	10.9	20.3	21.5	1638.1	26.7	94.0
Trial3_762	45	NA()	1.8	11.5589	1.6	2.7680	3.3	0.2322	2.9	0.87	1345.8	11.8	21.9	22.5	1346.9	8.6	15.8	16.4	1349.7	12.2	21.5	22.8	1349.7	30.9	99.7
Trial3_763	176	NA()	2.1	12.5933	1.0	2.1008	2.1	0.1920	1.9	0.89	1132.0	7.7	17.5	18.1	1149.1	5.6	13.4	14.0	1182.3	7.0	19.5	20.9	1182.3	19.0	95.7
Trial3_764	57	NA()	2.6	13.1614	1.8	1.7457	3.4	0.1667	2.9	0.86	994.0	9.5	16.9	17.4	1025.6	8.0	14.0	14.5	1094.6	13.6	22.9	24.2	1094.6	35.1	90.8
Trial3_765	44	NA()	0.9	5.5542	1.0	12.3438	2.5	0.4975	2.3	0.92	2602.9	12.9	34.9	36.2	2630.9	8.3	18.6	19.4	2653.3	10.6	18.6	19.7	2653.3	16.4	98.1
Trial3_766	270	NA()	5.5	13.2145	1.3	0.7241	2.7	0.0694	2.4	0.88	432.7	2.7	6.9	7.1	553.1	3.3	8.2	8.6	1086.5	8.5	20.3	21.7	432.7	10.1	39.8
Trial3_767	121	NA()	2.2	12.5602	1.1	2.2358	2.5	0.2038	2.3	0.90	1195.5	9.4	19.0	19.6	1192.4	6.7	14.2	14.7	1187.5	8.4	20.1	21.4	1187.5	22.2	100.7
Trial3_768	153	NA()	0.6	5.4669	0.9	12.5308	1.4	0.4971	1.0	0.73	2601.2	10.1	34.0	35.2	2645.0	6.3	17.8	18.7	2679.5	7.9	17.2	18.3	2679.5	15.4	97.1
Trial3_769	684	NA()	5.4	16.7397	0.9	0.6637	1.5	0.0806	1.3	0.83	499.8	1.7	7.5	7.8	516.8	1.8	7.4	7.8	594.1	5.9	20.8	22.4	499.8	6.2	84.1
Trial3_770	345	NA()	1.9	10.4732	0.8	3.3999	1.4	0.2584	1.1	0.79	1481.5	5.0	20.7	21.5	1504.4	3.4	14.3	15.0	1537.6	4.0	17.8	19.2	1537.6	15.6	96.4
Trial3_771	120	NA()	1.8	11.7419	1.0	2.6346	2.4	0.2245	2.2	0.90	1305.4	10.1	20.6	21.2	1310.4	7.0	14.8	15.4	1319.3	8.0	19.6	20.9	1319.3	20.1	99.0
Trial3_772	38	NA()	1.1	18.8905	4.4	0.2774	5.5	0.0380	3.3	0.60	240.5	3.3	4.8	4.9	248.6	7.7	8.6	8.7	326.1	72.7	75.6	76.1	240.5	7.8	NA
Trial3_773	270	NA()	1.3	18.2225	1.5	0.5100	2.9	0.0674	2.5	0.85	420.7	2.8	6.8	7.0	418.4	2.9	6.7	7.0	407.2	11.4	23.5	25.0	420.7	10.0	103.3
Trial3_774	51	NA()	1.3	12.2827	1.6	2.3941	3.3	0.2134	2.9	0.87	1246.7	11.7	20.8	21.4	1240.8	8.7	15.4	15.9	1231.5	12.5	21.9	23.2	1231.5	31.7	101.2
Trial3_775	71	NA()	1.5	13.0687	1.5	1.9205	3.2	0.1821	2.8	0.88	1078.5	10.2	18.2	18.7	1088.2	7.8	14.2	14.7	1108.7	11.2	21.6	22.9	1108.7	30.6	97.3
Trial3_776	45	NA()	2.1	13.2305	2.0	1.7845	3.7	0.1713	3.0	0.83	1019.3	10.3	17.6	18.1	1039.8	8.9	14.6	15.1	1084.0	16.6	24.9	26.0	1084.0	40.4	94.0
Trial3_777	127	NA()	1.1	10.7081	1.0	3.0049	2.3	0.2335	2.1	0.91	1352.7	10.0	21.0	21.7	1408.9	6.9	15.2	15.8	1495.7	7.4	18.9	20.3	1495.7	18.4	90.4
Trial3_778	31	NA()	1.5	7.6251	1.3	6.7658	3.1	0.3743	2.8	0.90	2049.8	14.6	30.3	31.3	2081.3	9.3	18.2	18.9	2113.3	11.2	19.6	20.7	2113.3	23.5	97.0
Trial3_779	22	NA()	0.5	10.7057	2.1	3.3171	3.7	0.2577	3.1	0.83	1477.9	13.6	24.2	24.9	1485.1	10.4	17.3	17.9	1496.2	16.1	23.7	24.8	1496.2	39.3	98.8
Trial3_780	178	NA()	1.3	5.5250	0.9	12.6908	1.4	0.5088	1.1	0.77	2651.3	9.8	34.4	35.7	2657.0	6.0	17.7	18.6	2662.0	7.5	17.0	18.1	2662.0	15.2	99.6
Trial3_781	67	NA()	0.7	12.9013	1.5	1.9972	3.2	0.1870	2.8	0.88	1104.9	10.2	18.4	19.0	1114.6	7.8	14.3	14.8	1134.4	11.2	21.5	22.8	1134.4	30.7	97.4
Trial3_782	116	NA()	2.1	13.1343	1.2	1.9991	2.7	0.1905	2.4	0.89	1124.2	8.9	18.0	18.5	1115.2	6.5	13.7	14.2	1098.7	8.8	20.4	21.8	1098.7	24.1	102.3
Trial3_783	159	NA()	1.6	17.0294	1.6	0.7946	3.1	0.0982	2.6	0.86	603.7	5.0	10.0	10.4	593.8	4.6	9.2	9.6	556.8	12.0	23.4	24.8	603.7	15.1	108.4
Trial3_784	135	NA()	1.6	19.1887	2.8	0.2797	4.2	0.0389	3.1	0.74	246.3	2.4	4.4	4.5	250.4	3.4	5.2	5.4	290.4	26.5	33.8	34.9	246.3	7.5	NA
Trial3_785	141	NA()	2.0	13.2853	1.1	1.8720	2.5	0.1805	2.3	0.89	1069.5	8.0	16.9	17.5	1071.2	6.0	13.2	13.8	1075.8	8.3	20.3	21.7	1075.8	22.8	99.4
Trial3_786	515	NA()	1.9	17.5596	1.1	0.5221	2.1	0.0665	1.8	0.85	415.2	0.4	6.1	6.4	426.5	1.3	6.3	6.6	489.6	8.2	21.9	23.5	415.2	7.3	84.8
Trial3_787	123	NA()	0.8	5.3081	0.9	13.6685	1.5	0.5264	1.2	0.78	2726.5	10.5	35.3	36.6	2727.0	6.5	18.0	18.8	2728.2	8.3	17.3	18.4	2728.2	15.6	99.9
Trial3_788	64	NA()	0.9	10.1627	1.2	3.8841	2.8	0.2864	2.5	0.90	1623.6	12.3	25.0	25.8	1610.4	8.1	16.4	17.1	1594.0	9.8	19.8	21.0	1594.0	22.6	101.9
Trial3_789	46	NA()	0.9	5.4374	1.0	12.9417	2.5	0.5106	2.2	0.91	2659.2	13.0	35.5	36.8	2675.4	8.1	18.6	19.4	2688.5	10.2	18.3	19.4	2688.5	16.7	98.9
Trial3_790	143	NA()	3.9	5.0425	1.0	15.7155	1.4	0.5750	1.1	0.75	2928.3	9.2	36.8	38.3	2859.7	6.2	18.0	18.9	2812.4	8.4	17.3	18.4	2812.4	15.6	104.1
Trial3_791	344	NA()	2.4	12.9362	0.8	1.9977	1.4	0.1875	1.1	0.80	1107.9	3.8	15.9	16.5	1114.7	3.0	12.4	13.0	1129.0	4.7	18.9	20.4	1129.0	18.9	98.1
Trial3_792	63	NA()	3.0	9.6127	1.2	4.4474	2.7	0.3102	2.5	0.91	1741.7	12.1	26.1	27.0	1721.2	7.8	16.6	17.3	1697.2	9.3	19.4	20.6	1697.2	19.4	102.6
Trial3_793	89	NA()	0.7	20.5222	4.8	0.1021	5.8	0.0152	3.2	0.55	97.2	1.2	1.9	1.9	98.7	3.8	4.1	4.1	134.7	89.5	92.0	92.5	97.2	1.9	NA
Trial3_794	123	NA()	0.6	8.8259	0.9	5.0900	2.0	0.3260	1.8	0.89	1818.8	10.5	26.2	27.1	1834.4	6.5	16.4	17.1	1853.0	7.2	18.1	19.4	1853.0	18.1	98.2
Trial3_795	156	NA()	2.5	13.9286	1.1	1.6250	2.5	0.1642	2.2	0.89	980.2	6.9	15.4	15.9	979.9	5.4	12.4	12.9	980.1	8.1	20.4	21.9	980.1	20.4	100.0
Trial3_796	164	NA()	2.2	13.6340	1.1	1.7256	2.4	0.1707	2.1	0.89	1016.0	6.9	15.8	16.4	1018.1	5.3	12.6	13.1	1023.5	7.7	20.2	21.6	1023.5	20.2	99.3
Trial3_797	214	NA()	3.3	5.1090	0.9	13.2839	1.4	0.4924	1.1	0.77	2581.2	7.7	33.1	34.4	2700.0	4.8	17.4	18.3	2790.9	5.8	16.1	17.3	2790.9	16.1	92.5
Trial3_798	103	NA()	1.4	13.7032	1.4	1.6224	2.9	0.1613	2.6	0.88	964.1	8.0	15.7	16.2	978.9	6.4	12.9	13.4	1013.3	10.2	21.3	22.6	1013.3	21.3	95.1
Trial3_799	339	NA()	1.5	13.0300	0.8	1.8431	1.5	0.1743	1.2	0.82	1035.5	3.5	14.9	15.5	1061.0	2.9	12.0	12.6	1114.6	4.9	19.0	20.5	1114.6	19.0	92.9
Trial3_800	201	NA()	0.6	16.4698	1.4	0.8340	2.8	0.0997	2.4	0.87	612.5	4.2	9.8	10.2	615.8	4.0	9.1	9.5	629.2	10.5	22.5	23.9	612.5	9.8	97.3
Trial3_801	170	NA()	1.7	19.9629	2.6	0.2585	4.0	0.0375	3.0	0.76	237.0	2.0	4.1	4.2	233.5	2.9	4.7	4.8	199.3	24.5	32.5	33.7	237.0	4.1	NA
Trial3_802	260	NA()	0.6	20.3695	2.5	0.1776	3.9	0.0262	2.9	0.76	167.0	1.1	2.7	2.8	166.0	1.8	3.3	3.4	152.3	22.6	31.2	32.5	167.0	2.7	NA
Trial3_803	158	NA()	3.0	13.2437	1.0	1.9849	2.3	0.1907	2.1	0.89	1125.4	7.6	17.4	18.0	1110.4	5.6	13.2	13.8	1082.0	7.6	20.0	21.4	1082.0	20.0	104.0
Trial3_804	43	NA()	2.7	12.3638	1.8	2.1558	3.5	0.1934	3.0	0.85	1139.8	10.5	19.0	19.5	1166.9	8.5	14.9	15.5	1218.5	13.8	22.8	24.0	1218.5	22.8	93.5
Trial3_805	58	NA()	1.4	13.7355	1.9	1.5473	3.5	0.1542	3.0	0.85	924.5	8.9	15.8	16.3	949.4	7.7	13.4	13.8	1008.5	14.0	23.3	24.6	1008.5	23.3	91.7
Trial3_806	191	NA()	1.0	17.5768	1.7	0.5530	3.2	0.0705	2.7	0.84	439.4	3.5	7.3	7.5	447.0	3.6	7.4	7.7	487.4	12.9	24.1	25.5	439.4	7.3	90.1
Trial3_807	61	NA()	1.4	13.4039	1.7	1.7025	3.4	0.1656	2.9	0.86	987.7	9.1	16.6	17.1	1009.5	7.6	13.7	14.2	1057.9	13.5	23.0	24.2	1057.9	23.0	93.4
Trial3_808	54	NA()	0.6	11.1795	1.4	2.9901	3.1	0.2425	2.7	0.89	1399.9	11.3	22.2	22.9	1405.1	8.0	15.7	16.3	1413.8	10.6	20.5	21.8	1413.8	20.5	99.0
Trial3_809	220	NA()	2.4	11.1201	0.8	2.6013	1.7	0.2099	1.6	0.89	1228.2	6.6	18.2	18.9	1301.0	4.8	13.9	14.5	1424.0	5.6	18.4	19.8	1424.0	18.4	86.3
Trial3_810	193	NA()	0.9	18.3352	1.8	0.4963	3.3	0.0660	2.7	0.83	412.2	3.3	6.9	7.1	409.2	3.5	6.9	7.2	393.4	14.6	25.3	26.7	412.2	6.9	104.8
Trial3_811	103	NA()	1.8	15.9013	1.8	0.9056	3.4	0.1045	2.8	0.85	640.7	6.1	11.1	11.4	654.7	5.7	10.3	10.7	704.4	13.7	23.9	25.3	640.7	11.1	90.9
Trial3_812	12	NA()	2.2	4.7904	1.5	15.6901	3.3	0.5454	3.0	0.90	2805.9	15.0	37.6	38.9	2858.1	9.9	19.6	20.4	2895.9	13.0	19.8	20.7	2895.9	19.8	96.9
Trial3_813	25	NA()	0.6	7.6442	1.5	6.5178	3.3	0.3615	2.9	0.89	1989.4	14.4	29.7	30.6	2048.3	9.5	18.3	19.0	2109.0	11.9	20.1	21.2	2109.0	20.1	94.3
Trial3_814	212	NA()	2.6	9.1891	0.9	4.7983	1.4	0.3199	1.1	0.80	1789.4	8.4	25.1	26.1	1784.6	5.3	15.8	16.5	1779.8	5.8	17.8	19.1	1779.8	17.8	100.5
Trial3_815	202	NA()	1.2	13.4912	1.0	1.7787	2.1	0.1741	1.9	0.89	1034.8	6.7	16.0	16.5	1037.7	5.1	12.6	13.1	1044.8	6.9	19.8	21.3	1044.8	19.8	99.0
Trial3_816	123	NA()	1.7	16.1691	1.8	0.8200	3.3	0.0962	2.8	0.85	592.1	5.5	10.2	10.5	608.1	5.4	9.7	10.1	668.8	14.1	24.2	25.5	592.1	10.2	88.5
Trial3_817	48	NA()	1.3	9.0215	1.3	4.8241	2.9	0.3158	2.6	0.90	1769.1	13.2	26.9	27.8	1789.1	8.6	17.2	17.9	1813.3	10.3	19.6	20.8	1813.3	19.6	97.6
Trial3_818	142	NA()	1.5	17.3771	1.9	0.5508	3.5	0.0694	2.9	0.83	432.8	4.0	7.5	7.7	445.5	4.2	7.7	7.9	512.5	15.0	25.2	26.6	432.8	7.5	84.4
Trial3_819	151	NA()	1.9	10.7809	0.9	3.0289	2.1	0.2369	1.9	0.90	1370.8	9.1	20.8	21.5	1414.9	6.3	14.9	15.6	1482.9	6.8	18.7	20.1	1482.9	18.7	92.4
Trial3_820	112	NA()	1.4	17.8706	2.2	0.5659	3.7	0.0734	3.0	0.81	456.5	4.5	8.1	8.3	455.4	4.8	8.1	8.4	450.7	18.1	27.3	28.6	456.5	8.1	101.3
Trial3_821	181	NA()	1.9	12.4611	1.0	2.1445	2.1	0.1939	1.9	0.89	1142.5	7.7	17.6	18.2	1163.3	5.6	13.5	14.1	1203.1	7.0	19.4	20.8	1203.1	19.4	95.0
Trial3_822	32	NA()	0.5	5.0867	1.1	13.4351	2.7	0.4959	2.5	0.92	2596.1	13.5	35.1	36.3	2710.7	8.5	18.8	19.6	2798.1	10.5	18.4	19.4	2798.1	18.4	92.8
Trial3_823	47	NA()	1.4	12.9042	1.8	1.9880	3.5	0.1861	3.0	0.85	1100.4	10.6	18.6	19.2	1111.4	8.5	14.7	15.2	1133.9	13.5	22.8	24.0	1133.9	22.8	97.0
Trial3_824	20	NA()	1.8	13.3771	2.9	1.8068	4.3	0.1754	3.3	0.75	1041.7	11.2	18.4	18.8	1047.9	10.7	15.8	16.2	1061.9	23.2	29.7	30.7	1061.9	29.7	98.1
Trial3_825	29	NA()	1.9	13.3625	2.4	1.7639	4.0	0.1710	3.2	0.80	1017.7	10.6	17.8	18.2	1032.3	9.5	14.9	15.4	1064.1	18.8	26.4	27.5	1064.1	26.4	95.6
Trial3_826	172	NA()	1.4	10.9259	0.8	2.9946	1.9	0.2374	1.7	0.90	1373.2	8.8	20.7	21.4	1406.2	6.0	14.8	15.4	1457.6	6.5	18.7	20.0	1457.6	18.7	94.2
Trial3_827	31	NA()	0.5	19.0304	5.3	0.1929	6.3	0.0266	3.3	0.53	169.5	2.4	3.5	3.6	179.1	7.6	8.2	8.2	309.3	100.9	103.0	103.4	169.5	3.5	NA
Trial3_828	78	NA()	1.4	10.6576	1.2	2.8935	2.8	0.2238	2.5	0.90	1301.7	10.7	20.8	21.4	1380.2	7.7	15.4	16.1	1504.7	9.0	19.6	20.9	1504.7	19.6	86.5
Trial3_829	181	NA()	3.2	9.2181	0.9	4.0699	1.7	0.2722	1.5	0.86	1552.1	8.6	22.6	23.4	1648.3	5.7	15.5	16.3	1774.1	5.9	17.8	19.1	1774.1	17.8	87.5
Trial3_830	200	NA()	0.9	18.1392	1.7	0.5290	3.1	0.0696	2.6	0.84	433.9	3.3	7.2	7.4	431.2	3.5	7.1	7.4	417.4	13.4	24.5	26.0	433.9	7.2	103.9
Trial3_831	82	NA()	2.5	13.1362	1.4	1.8778	3.0	0.1790	2.7	0.88	1061.4	9.1	17.4	17.9	1073.3	7.1	13.7	14.3	1098.4	10.6	21.3	22.6	1098.4	21.3	96.6
Trial3_832	53	NA()	0.2	19.0655	3.5	0.3438	4.7	0.0476	3.2	0.68	299.6	3.6	5.7	5.8	300.1	5.5	7.2	7.3	305.1	39.7	44.9	45.7	299.6	5.7	NA
Trial3_833	197	NA()	2.7	16.3960	1.4	0.8529	2.8	0.1015	2.4	0.87	623.0	4.5	10.1	10.4	626.2	4.2	9.3	9.7	638.9	10.6	22.5	23.9	623.0	10.1	97.5
Trial3_834	91	NA()	1.4	8.6404	0.9	5.3234	2.2	0.3337	2.0	0.91	1856.5	11.8	27.1	28.1	1872.6	7.4	16.9	17.6	1891.3	8.2	18.5	19.7	1891.3	18.5	98.2
Trial3_835	128	NA()	1.4	19.0192	2.9	0.2689	4.3	0.0371	3.1	0.73	234.9	2.4	4.2	4.3	241.8	3.5	5.2	5.3	310.7	28.5	35.4	36.5	234.9	4.2	NA
Trial3_836	122	NA()	1.2	13.4544	1.2	1.7742	2.7	0.1732	2.4	0.89	1029.8	8.2	16.6	17.1	1036.1	6.3	13.1	13.7	1050.3	8.9	20.6	22.0	1050.3	20.6	98.0
Trial3_837	58	NA()	1.9	13.6543	1.8	1.7100	3.4	0.1694	2.9	0.85	1008.9	9.2	16.9	17.4	1012.3	7.6	13.7	14.2	1020.5	13.4	22.9	24.2	1020.5	22.9	98.9
Trial3_838	35	NA()	2.3	13.0010	2.1	1.9041	3.8	0.1796	3.1	0.82	1064.9	10.2	18.1	18.6	1082.5	8.8	14.7	15.3	1119.1	16.4	24.6	25.8	1119.1	24.6	95.2
Trial3_839	79	NA()	1.5	13.7530	1.5	1.7657	3.1	0.1762	2.7	0.87	1046.2	9.0	17.2	17.7	1032.9	7.0	13.5	14.0	1005.9	11.3	21.8	23.1	1005.9	21.8	104.0
Trial3_840	42	NA()	1.8	9.3664	1.4	4.4764	3.0	0.3042	2.7	0.89	1712.2	12.8	26.1	26.9	1726.6	8.6	17.0	17.7	1744.9	10.8	20.0	21.2	1744.9	20.0	98.1
Trial3_841	95	NA()	34.4	13.5011	1.4	1.7056	3.0	0.1671	2.6	0.88	996.0	8.5	16.4	16.9	1010.6	6.7	13.2	13.7	1043.3	10.3	21.3	22.6	1043.3	21.3	95.5
Trial3_842	120	NA()	1.0	17.7365	2.1	0.5540	3.6	0.0713	2.9	0.81	444.0	4.2	7.7	8.0	447.6	4.6	7.9	8.1	467.4	17.4	26.8	28.1	444.0	7.7	95.0
Trial3_843	120	NA()	0.9	12.8607	1.2	2.0488	2.6	0.1912	2.3	0.89	1127.8	8.6	17.9	18.5	1131.9	6.4	13.7	14.3	1140.6	8.7	20.2	21.6	1140.6	20.2	98.9
Trial3_844	113	NA()	4.2	13.4419	1.3	1.7431	2.8	0.1700	2.5	0.89	1012.2	8.1	16.4	16.9	1024.6	6.4	13.1	13.6	1052.2	9.5	20.8	22.2	1052.2	20.8	96.2
Trial3_845	49	NA()	1.2	13.4209	1.9	1.8285	3.5	0.1781	3.0	0.85	1056.4	10.1	17.9	18.4	1055.7	8.2	14.2	14.7	1055.4	14.0	23.3	24.5	1055.4	23.3	100.1
Trial3_846	25	NA()	0.8	8.3501	1.6	5.2309	3.4	0.3169	3.0	0.88	1774.7	14.0	27.3	28.2	1857.7	9.7	17.9	18.6	1952.6	12.3	20.6	21.7	1952.6	20.6	90.9
Trial3_847	71	NA()	2.1	13.4308	1.6	1.8142	3.2	0.1768	2.8	0.87	1049.5	9.5	17.5	18.0	1050.6	7.4	13.8	14.3	1053.9	11.5	21.9	23.2	1053.9	21.9	99.6
Trial3_848	114	NA()	1.5	19.9946	3.4	0.2397	4.6	0.0348	3.2	0.69	220.3	2.3	4.0	4.1	218.1	3.8	5.1	5.3	195.6	37.3	43.0	43.9	220.3	4.0	NA
Trial3_849	72	NA()	1.9	13.7684	1.6	1.6673	3.2	0.1666	2.8	0.87	993.2	9.1	16.7	17.1	996.2	7.3	13.4	13.9	1003.6	12.1	22.3	23.6	1003.6	22.3	99.0
Trial3_850	21	NA()	1.0	7.5410	1.6	6.7670	3.4	0.3703	3.0	0.88	2030.7	14.7	30.2	31.2	2081.4	9.8	18.5	19.2	2132.8	12.5	20.4	21.5	2132.8	20.4	95.2
Trial3_851	113	NA()	2.5	10.8794	1.0	3.2034	2.4	0.2529	2.1	0.90	1453.3	10.3	22.2	23.0	1458.0	6.9	15.4	16.0	1465.6	7.9	19.2	20.5	1465.6	19.2	99.2
Trial3_852	150	NA()	1.8	8.3150	0.9	5.6541	1.7	0.3411	1.5	0.86	1892.1	10.2	26.9	27.8	1924.4	6.3	16.5	17.3	1960.1	6.8	17.8	19.0	1960.1	17.8	96.5
Trial3_853	121	NA()	4.5	12.7858	1.2	1.9299	2.7	0.1790	2.4	0.89	1061.7	8.4	17.1	17.6	1091.5	6.5	13.5	14.1	1152.2	8.7	20.3	21.6	1152.2	20.3	92.1
Trial3_854	86	NA()	1.3	16.6572	2.1	0.7870	3.7	0.0951	3.0	0.82	585.8	6.0	10.4	10.7	589.5	5.9	9.9	10.2	604.8	16.8	26.1	27.4	585.8	10.4	96.9
Trial3_855	103	NA()	1.9	12.4989	1.2	2.3042	2.7	0.2090	2.4	0.89	1223.3	9.9	19.6	20.2	1213.6	7.1	14.4	15.0	1197.2	8.9	20.2	21.6	1197.2	20.2	102.2
Trial3_856	25	NA()	0.8	5.4938	1.2	12.7492	3.0	0.5082	2.7	0.91	2649.0	14.3	35.9	37.1	2661.3	9.1	19.0	19.8	2671.4	11.7	19.2	20.2	2671.4	19.2	99.2
Trial3_857	58	NA()	0.6	11.1967	1.4	3.1031	3.0	0.2521	2.7	0.89	1449.3	12.1	23.1	23.8	1433.5	8.3	16.0	16.6	1410.9	10.6	20.6	21.8	1410.9	20.6	102.7
Trial3_858	75	NA()	1.0	9.5202	1.1	4.2550	2.6	0.2939	2.4	0.91	1661.1	12.3	25.4	26.2	1684.7	8.0	16.6	17.3	1715.0	8.9	19.1	20.4	1715.0	19.1	96.9
Trial3_859	53	NA()	1.6	13.5563	1.8	1.8497	3.4	0.1819	2.9	0.85	1077.6	10.5	18.3	18.8	1063.3	8.2	14.3	14.8	1035.1	13.4	22.9	24.2	1035.1	22.9	104.1
Trial3_860	126	NA()	2.0	11.7568	1.1	2.2787	2.6	0.1944	2.3	0.90	1145.1	9.2	18.3	18.9	1205.7	6.8	14.2	14.8	1316.8	7.9	19.5	20.9	1316.8	19.5	87.0
Trial3_861	158	NA()	1.1	13.7938	1.1	1.7473	2.4	0.1749	2.2	0.89	1039.0	7.8	16.5	17.1	1026.2	5.9	12.9	13.4	999.9	8.0	20.3	21.8	999.9	20.3	103.9
Trial3_862	92	NA()	1.0	8.7891	0.9	5.2229	2.3	0.3331	2.1	0.91	1853.3	12.3	27.3	28.2	1856.4	7.6	16.9	17.6	1860.6	8.4	18.6	19.9	1860.6	18.6	99.6
Trial3_863	52	NA()	0.8	8.8583	1.2	5.1389	2.8	0.3303	2.6	0.91	1839.9	13.6	27.8	28.7	1842.6	8.6	17.3	18.0	1846.4	10.0	19.4	20.6	1846.4	19.4	99.6
Trial3_864	20	NA()	1.1	13.2939	2.9	1.7642	4.4	0.1702	3.3	0.74	1013.1	11.1	18.0	18.5	1032.4	11.1	16.0	16.4	1074.5	24.8	31.0	31.9	1074.5	31.0	94.3
Trial3_865	19	NA()	0.6	5.5102	1.3	12.8238	3.1	0.5127	2.8	0.90	2668.2	14.9	36.3	37.5	2666.8	9.5	19.2	20.0	2666.4	12.4	19.6	20.6	2666.4	19.6	100.1
Trial3_866	124	NA()	1.7	13.2956	1.2	1.9570	2.6	0.1888	2.3	0.89	1114.8	8.9	17.9	18.5	1100.9	6.5	13.6	14.1	1074.2	8.6	20.4	21.8	1074.2	20.4	103.8
Trial3_867	29	NA()	1.0	5.0422	1.1	14.8424	2.8	0.5430	2.6	0.92	2796.1	13.7	37.0	38.3	2805.2	8.8	19.0	19.8	2812.5	11.4	18.9	19.9	2812.5	18.9	99.4
Trial3_868	110	NA()	1.4	16.6262	1.8	0.8489	3.4	0.1024	2.9	0.84	628.5	6.1	10.9	11.2	624.0	5.7	10.0	10.4	608.8	14.4	24.6	26.0	628.5	10.9	103.2
Trial3_869	388	NA()	2.2	20.2917	2.8	0.1032	4.0	0.0152	3.0	0.73	97.2	0.4	1.5	1.6	99.7	1.2	2.1	2.1	161.2	27.6	35.1	36.2	97.2	1.5	NA
Trial3_870	46	NA()	1.7	13.5633	2.0	1.6571	3.7	0.1631	3.1	0.83	973.9	9.9	16.9	17.4	992.3	8.5	14.1	14.6	1034.0	15.6	24.3	25.5	1034.0	24.3	94.2
Trial3_871	14	NA()	1.9	13.2841	3.4	1.8816	4.7	0.1814	3.2	0.69	1074.4	11.9	19.2	19.7	1074.6	13.1	17.6	18.0	1076.0	31.4	36.4	37.2	1076.0	36.4	99.9
Trial3_872	22	NA()	0.5	10.9172	2.1	3.2606	3.8	0.2583	3.1	0.82	1481.1	13.6	24.2	24.9	1471.7	10.5	17.3	17.9	1459.1	16.7	24.2	25.2	1459.1	24.2	101.5
Trial3_873	74	NA()	1.0	5.4875	1.0	13.0576	2.0	0.5199	1.8	0.87	2698.8	12.1	35.5	36.8	2683.8	7.6	18.4	19.2	2673.3	9.9	18.2	19.2	2673.3	18.2	101.0
Trial3_874	47	NA()	1.2	13.3841	1.9	1.8775	3.5	0.1823	3.0	0.85	1079.7	10.5	18.4	18.9	1073.2	8.4	14.4	14.9	1060.9	13.9	23.2	24.4	1060.9	23.2	101.8
Trial3_875	85	NA()	1.3	10.9825	1.2	3.0912	2.7	0.2463	2.4	0.90	1419.6	11.2	22.3	23.0	1430.5	7.7	15.6	16.3	1447.7	9.3	19.9	21.1	1447.7	19.9	98.1
Trial3_876	59	NA()	1.9	12.8047	1.6	1.9666	3.3	0.1827	2.9	0.87	1081.8	10.2	18.2	18.7	1104.1	8.1	14.4	14.9	1149.3	12.2	22.0	23.3	1149.3	22.0	94.1
Trial3_877	58	NA()	1.4	13.4998	1.7	1.8461	3.3	0.1808	2.9	0.86	1071.5	10.0	18.0	18.5	1062.0	7.9	14.1	14.6	1043.5	13.2	22.8	24.1	1043.5	22.8	102.7
Trial3_878	95	NA()	0.3	16.6171	2.0	0.8118	3.5	0.0979	2.9	0.83	602.0	6.0	10.6	10.9	603.5	5.8	9.9	10.3	610.0	15.8	25.4	26.7	602.0	10.6	98.7
Trial3_879	145	NA()	0.6	17.8877	2.0	0.5429	3.5	0.0705	2.9	0.83	439.0	4.0	7.6	7.8	440.4	4.2	7.6	7.9	448.5	15.4	25.6	27.0	439.0	7.6	97.9
Trial3_880	86	NA()	0.8	18.4012	2.8	0.4245	4.2	0.0567	3.2	0.75	355.4	3.9	6.5	6.7	359.3	4.6	7.1	7.3	385.3	23.6	31.3	32.5	355.4	6.5	NA
Trial3_881	60	NA()	1.2	5.5012	1.0	12.0389	2.3	0.4805	2.0	0.90	2529.7	12.7	34.1	35.4	2607.4	7.9	18.4	19.2	2669.2	9.6	18.0	19.1	2669.2	18.0	94.8
Trial3_882	82	NA()	0.8	5.6466	1.0	11.8844	2.0	0.4869	1.7	0.87	2557.3	12.0	34.2	35.4	2595.3	7.5	18.2	19.0	2625.9	9.3	17.9	19.0	2625.9	17.9	97.4
Trial3_883	32	NA()	2.1	13.2595	2.3	1.8541	3.9	0.1784	3.1	0.81	1058.2	10.7	18.3	18.8	1064.9	9.2	14.9	15.3	1079.7	17.0	25.1	26.3	1079.7	25.1	98.0
Trial3_884	42	NA()	1.3	4.6489	1.0	16.5306	2.4	0.5576	2.2	0.90	2856.8	12.4	37.1	38.5	2908.0	8.2	18.9	19.7	2944.4	10.8	18.4	19.4	2944.4	18.4	97.0
Trial3_885	268	NA()	2.3	18.6339	1.9	0.3034	3.4	0.0410	2.7	0.82	259.1	1.7	4.2	4.4	269.0	2.3	4.8	5.0	357.0	15.6	26.0	27.4	259.1	4.2	NA
Trial3_886	42	NA()	0.8	8.7484	1.3	5.0179	3.0	0.3185	2.7	0.90	1782.5	13.5	27.2	28.0	1822.3	8.8	17.4	18.1	1869.0	10.3	19.6	20.7	1869.0	19.6	95.4

Trial3_887	21	NA()	0.9	11.7986	2.4	2.5020	4.0	0.2142	3.2	0.80	1251.1	12.4	21.2	21.8	1272.6	10.5	16.6	17.1	1309.9	18.4	25.7	26.7	1309.9	25.7	95.5
Trial3_888	117	NA()	3.5	9.3377	0.9	4.5059	2.1	0.3053	1.9	0.91	1717.5	11.1	25.4	26.3	1732.1	7.0	16.3	17.0	1750.5	7.7	18.5	19.8	1750.5	18.5	98.1
Trial3_889	123	NA()	2.3	18.0947	2.3	0.5013	3.7	0.0658	3.0	0.80	410.9	4.1	7.3	7.5	412.6	4.8	7.7	7.9	422.9	21.2	29.5	30.7	410.9	7.3	97.2
Trial3_890	47	NA()	1.8	7.7389	1.1	6.6613	2.7	0.3741	2.5	0.91	2048.4	13.8	29.9	30.9	2067.5	8.6	17.9	18.6	2087.3	10.2	19.1	20.3	2087.3	19.1	98.1
Trial3_891	25	NA()	2.3	13.0438	2.5	1.9486	4.0	0.1844	3.2	0.79	1091.1	11.3	19.0	19.5	1098.0	9.8	15.4	15.9	1112.5	18.3	26.0	27.1	1112.5	26.0	98.1
Trial3_892	32	NA()	1.1	13.4294	2.4	1.6961	4.0	0.1653	3.2	0.80	986.0	10.4	17.3	17.8	1007.1	9.6	14.8	15.3	1054.1	19.5	26.9	28.0	1054.1	26.9	93.5
Trial3_893	54	NA()	1.5	11.3383	1.5	2.7776	3.2	0.2285	2.8	0.88	1326.7	11.8	21.7	22.3	1349.5	8.5	15.7	16.3	1386.8	11.1	20.9	22.1	1386.8	20.9	95.7
Trial3_894	18	NA()	2.9	12.9056	2.8	2.0257	4.3	0.1897	3.3	0.76	1119.7	11.8	19.6	20.1	1124.2	10.9	16.2	16.7	1133.7	22.2	28.8	29.8	1133.7	28.8	98.8
Trial3_895	102	NA()	2.2	10.7280	1.0	3.2393	2.5	0.2522	2.2	0.91	1449.6	10.9	22.5	23.2	1466.6	7.3	15.6	16.2	1492.2	8.2	19.3	20.6	1492.2	19.3	97.1
Trial3_896	25	NA()	1.6	12.9526	2.5	1.8842	4.1	0.1771	3.2	0.78	1051.0	11.1	18.4	18.9	1075.5	10.4	15.7	16.2	1126.5	21.2	28.1	29.1	1126.5	28.1	93.3
Trial3_897	54	NA()	1.9	13.2788	1.8	1.6784	3.5	0.1617	3.0	0.85	966.3	9.6	16.7	17.1	1000.4	8.2	13.9	14.4	1076.8	14.0	23.2	24.4	1076.8	23.2	89.7
Trial3_898	35	NA()	1.2	10.9824	1.7	3.0956	3.4	0.2467	3.0	0.86	1421.3	12.8	23.2	23.8	1431.6	9.3	16.5	17.1	1447.7	13.0	21.8	23.0	1447.7	21.8	98.2
Trial3_899	217	NA()	1.5	7.9820	0.9	5.4765	1.4	0.3172	1.1	0.79	1776.0	8.6	25.0	26.0	1896.9	5.5	16.2	16.9	2032.8	5.5	17.2	18.5	2032.8	17.2	87.4
Trial3_900	24	NA()	1.0	7.6470	1.5	6.8041	3.2	0.3775	2.9	0.89	2064.8	14.9	30.6	31.6	2086.3	9.7	18.4	19.1	2108.3	12.2	20.2	21.3	2108.3	20.2	97.9
Trial3_901	19	NA()	1.6	13.2652	3.1	1.7228	4.5	0.1658	3.3	0.73	989.0	10.9	17.7	18.1	1017.1	11.6	16.3	16.7	1078.8	27.4	33.0	33.9	1078.8	33.0	91.7
Trial3_902	78	NA()	1.9	11.1437	1.2	2.9207	2.8	0.2362	2.5	0.90	1366.7	11.2	21.8	22.4	1387.3	7.8	15.5	16.1	1419.9	9.3	19.9	21.2	1419.9	19.9	96.3
Trial3_903	38	NA()	1.4	11.2534	1.8	2.8238	3.4	0.2306	3.0	0.86	1337.5	12.3	22.0	22.7	1361.9	9.2	16.2	16.8	1401.2	13.2	22.1	23.2	1401.2	22.1	95.5
Trial3_904	26	NA()	0.7	13.1803	2.7	1.6296	4.2	0.1558	3.2	0.77	933.7	10.2	16.7	17.1	981.7	10.1	15.1	15.5	1091.7	21.8	28.6	29.6	1091.7	28.6	85.5
Trial3_905	238	NA()	0.8	17.8305	1.6	0.4991	3.1	0.0646	2.6	0.85	403.3	3.0	6.6	6.8	411.1	3.2	6.8	7.1	455.7	12.1	23.8	25.2	403.3	6.6	88.5
Trial3_906	349	NA()	1.6	13.8100	0.8	1.6070	1.5	0.1610	1.3	0.85	962.5	3.8	14.1	14.6	973.0	3.1	11.5	12.1	997.5	5.3	19.4	20.9	997.5	19.4	96.5
Trial3_907	136	NA()	2.9	13.3613	1.2	1.8099	2.6	0.1755	2.3	0.89	1042.2	8.4	16.8	17.3	1049.0	6.3	13.2	13.7	1064.3	8.5	20.4	21.8	1064.3	20.4	97.9
Trial3_908	28	NA()	1.3	11.4128	2.0	2.7627	3.7	0.2288	3.1	0.84	1328.1	12.7	22.2	22.8	1345.5	9.9	16.5	17.1	1374.2	15.3	23.4	24.5	1374.2	23.4	96.6
Trial3_909	303	NA()	0.7	17.8838	1.4	0.5515	2.7	0.0716	2.3	0.86	445.5	2.5	7.0	7.2	446.0	2.7	6.9	7.3	449.0	10.4	22.9	24.5	445.5	7.0	99.2
Trial3_910	128	NA()	1.0	19.6191	2.9	0.2596	4.3	0.0370	3.2	0.73	233.9	2.5	4.3	4.4	234.3	3.3	5.0	5.1	239.5	26.6	34.0	35.2	233.9	4.3	NA
Trial3_911	82	NA()	1.9	10.7837	1.2	3.1306	2.7	0.2450	2.4	0.90	1412.4	11.3	22.3	23.0	1440.2	7.8	15.7	16.3	1482.4	9.0	19.6	20.9	1482.4	19.6	95.3
Trial3_912	99	NA()	1.2	9.3264	0.9	4.5745	2.2	0.3096	2.0	0.91	1738.6	11.8	25.9	26.8	1744.6	7.4	16.5	17.2	1752.7	8.2	18.8	20.0	1752.7	18.8	99.2
Trial3_913	66	NA()	2.3	13.3811	1.7	1.7241	3.3	0.1674	2.9	0.86	997.7	9.5	16.9	17.4	1017.5	7.7	13.7	14.2	1061.3	12.1	22.2	23.5	1061.3	22.2	94.0
Trial3_914	145	NA()	2.8	11.0218	0.9	2.8910	2.2	0.2312	2.0	0.90	1340.8	9.4	20.6	21.3	1379.6	6.5	14.9	15.5	1440.9	7.1	18.9	20.3	1440.9	18.9	93.1
Trial3_915	216	NA()	3.8	14.0562	1.0	1.5176	2.2	0.1548	1.9	0.89	927.7	6.1	14.4	14.9	937.5	4.8	11.9	12.4	961.5	7.1	20.1	21.6	961.5	20.1	96.5
Trial3_916	217	NA()	0.8	16.3511	1.3	0.8257	2.7	0.0980	2.4	0.87	602.5	4.3	9.7	10.1	611.2	4.1	9.1	9.5	644.8	9.8	22.1	23.6	602.5	9.7	93.4
Trial3_917	129	NA()	1.7	8.8962	0.9	4.0449	2.1	0.2611	2.0	0.91	1495.5	10.3	22.7	23.4	1643.3	7.0	16.0	16.7	1838.7	6.9	18.0	19.3	1838.7	18.0	81.3
Trial3_918	65	NA()	1.3	10.8318	1.3	3.1048	2.9	0.2440	2.6	0.90	1407.6	11.7	22.5	23.1	1433.9	8.1	15.9	16.5	1474.0	9.8	20.0	21.3	1474.0	20.0	95.5
Trial3_919	140	NA()	2.5	12.7547	1.1	1.9355	2.5	0.1791	2.2	0.89	1062.2	8.0	16.8	17.4	1093.5	6.1	13.3	13.9	1157.1	7.9	19.9	21.3	1157.1	19.9	91.8
Trial3_920	93	NA()	1.6	9.6125	1.0	4.2531	2.4	0.2966	2.2	0.91	1674.6	11.4	25.1	25.9	1684.3	7.4	16.3	17.0	1697.2	8.3	18.9	20.2	1697.2	18.9	98.7
Trial3_921	34	NA()	1.6	20.0783	3.9	0.2779	5.1	0.0405	3.3	0.64	255.9	3.3	5.0	5.2	249.0	6.4	7.5	7.6	185.9	59.6	63.4	64.0	255.9	5.0	NA
Trial3_922	34	NA()	1.1	5.9611	1.1	10.9941	2.8	0.4755	2.5	0.92	2507.8	13.6	34.3	35.5	2522.6	8.6	18.6	19.4	2535.3	10.9	18.9	19.9	2535.3	18.9	98.9
Trial3_923	108	NA()	6.1	4.4425	1.0	18.3733	1.5	0.5922	1.1	0.76	2998.5	9.6	37.6	39.0	3009.5	6.7	18.3	19.2	3017.5	9.1	17.4	18.4	3017.5	17.4	99.4
Trial3_924	61	NA()	1.5	12.6783	1.6	2.0801	3.2	0.1914	2.8	0.87	1128.7	10.1	18.6	19.2	1142.3	7.8	14.4	15.0	1169.0	11.6	21.6	22.9	1169.0	21.6	96.6
Trial3_925	11	NA()	2.5	12.9601	3.4	1.8703	4.7	0.1759	3.2	0.69	1044.4	11.6	18.7	19.1	1070.7	13.1	17.6	18.0	1125.3	31.4	36.4	37.2	1125.3	36.4	92.8
Trial3_926	36	NA()	1.1	12.6970	2.1	2.0908	3.7	0.1926	3.1	0.83	1135.6	10.9	19.2	19.7	1145.8	9.1	15.2	15.7	1166.1	16.0	24.3	25.4	1166.1	24.3	97.4
Trial3_927	466	NA()	1.4	13.2432	0.8	1.7555	1.2	0.1687	0.9	0.72	1004.9	1.2	14.2	14.7	1029.2	1.6	11.6	12.2	1082.1	4.1	18.9	20.4	1082.1	18.9	92.9
Trial3_928	33	NA()	1.6	4.6442	1.0	16.9199	2.6	0.5702	2.4	0.92	2908.5	12.6	37.6	39.0	2930.3	8.3	18.9	19.7	2946.0	11.0	18.5	19.5	2946.0	18.5	98.7
Trial3_929	317	NA()	2.4	13.4207	0.8	1.7068	1.6	0.1662	1.4	0.86	991.2	4.0	14.5	15.1	1011.1	3.3	11.8	12.4	1055.4	5.5	19.4	20.8	1055.4	19.4	93.9
Trial3_930	45	NA()	1.6	11.2222	1.6	2.6355	3.3	0.2146	2.9	0.87	1253.3	11.2	20.6	21.2	1310.6	8.6	15.6	16.2	1406.5	11.9	21.3	22.5	1406.5	21.3	89.1
Trial3_931	63	NA()	1.4	13.3904	1.7	1.7598	3.3	0.1710	2.9	0.86	1017.5	9.4	17.1	17.6	1030.8	7.7	13.8	14.3	1059.9	12.9	22.6	23.8	1059.9	22.6	96.0
Trial3_932	38	NA()	2.0	13.2898	2.1	1.8772	3.7	0.1810	3.1	0.83	1072.5	10.4	18.2	18.8	1073.1	8.8	14.7	15.2	1075.1	16.0	24.5	25.6	1075.1	24.5	99.8
Trial3_933	307	NA()	0.9	18.2464	1.6	0.3836	3.0	0.0508	2.5	0.84	319.3	1.7	5.0	5.2	329.7	2.2	5.4	5.7	404.3	12.4	24.1	25.6	319.3	5.0	NA
Trial3_934	38	NA()	0.8	13.4072	2.2	1.6750	3.8	0.1630	3.1	0.82	973.2	9.6	16.7	17.2	999.1	8.6	14.2	14.7	1057.4	16.9	25.1	26.2	1057.4	25.1	92.0
Trial3_935	68	NA()	1.2	11.0592	1.3	3.0458	2.9	0.2444	2.6	0.90	1409.6	11.1	22.2	22.9	1419.2	7.7	15.6	16.2	1434.5	9.5	20.0	21.3	1434.5	20.0	98.3
Trial3_936	28	NA()	1.5	4.4649	1.0	18.8512	2.7	0.6107	2.5	0.92	3072.9	12.2	39.0	40.4	3034.2	8.3	19.0	19.8	3009.4	11.3	18.6	19.6	3009.4	18.6	102.1
Trial3_937	63	NA()	1.5	17.1083	2.7	0.6796	4.1	0.0844	3.2	0.76	522.1	5.5	9.4	9.6	526.5	6.3	9.6	9.9	546.7	23.4	30.8	31.9	522.1	9.4	95.5
Trial3_938	145	NA()	1.4	11.1568	0.9	3.0403	2.1	0.2461	1.9	0.90	1418.4	9.2	21.4	22.1	1417.8	6.2	14.9	15.6	1417.7	7.0	19.0	20.3	1417.7	19.0	100.1
Trial3_939	90	NA()	2.1	18.0829	2.7	0.4226	4.2	0.0554	3.2	0.75	347.9	3.7	6.3	6.5	357.9	4.5	7.0	7.2	424.4	22.7	30.6	31.8	347.9	6.3	NA
Trial3_940	49	NA()	2.0	13.5573	1.9	1.7270	3.6	0.1699	3.0	0.84	1011.5	9.7	17.2	17.7	1018.6	8.0	14.0	14.4	1034.9	14.3	23.5	24.7	1034.9	23.5	97.7
Trial3_941	45	NA()	0.9	19.1982	3.8	0.2730	5.0	0.0380	3.3	0.65	240.6	3.0	4.7	4.8	245.1	5.9	7.1	7.2	289.3	54.9	58.8	59.4	240.6	4.7	NA
Trial3_942	203	NA()	1.8	19.1891	2.4	0.2401	3.9	0.0334	3.0	0.77	212.0	1.7	3.6	3.7	218.5	2.4	4.2	4.4	290.3	20.8	29.6	30.9	212.0	3.6	NA
Trial3_943	78	NA()	2.6	13.3991	1.5	1.7891	3.1	0.1739	2.7	0.87	1033.8	9.1	17.1	17.6	1041.5	7.2	13.6	14.1	1058.6	11.2	21.7	23.0	1058.6	21.7	97.7
Trial3_944	159	NA()	5.4	13.6381	1.1	1.6877	2.4	0.1670	2.2	0.89	995.6	7.0	15.6	16.2	1003.9	5.5	12.6	13.1	1022.9	7.9	20.2	21.7	1022.9	20.2	97.3
Trial3_945	98	NA()	2.6	12.0192	1.2	2.6324	2.6	0.2296	2.3	0.90	1332.3	9.8	20.7	21.4	1309.7	6.8	14.7	15.3	1273.9	8.6	19.9	21.3	1273.9	19.9	104.6
Trial3_946	99	NA()	0.5	5.3200	1.0	13.6450	1.7	0.5267	1.4	0.83	2727.6	10.6	35.3	36.7	2725.4	6.7	18.1	18.9	2724.5	8.7	17.5	18.6	2724.5	17.5	100.1
Trial3_947	132	NA()	2.0	12.9589	1.2	1.9283	2.6	0.1813	2.3	0.89	1074.2	7.9	16.9	17.5	1091.0	6.0	13.3	13.9	1125.5	8.2	20.1	21.5	1125.5	20.1	95.4
Trial3_948	214	NA()	2.0	13.4017	0.9	1.8682	2.0	0.1817	1.8	0.88	1076.1	6.1	16.2	16.8	1069.9	4.6	12.6	13.2	1058.2	6.5	19.7	21.1	1058.2	19.7	101.7
Trial3_949	92	NA()	2.0	13.2524	1.4	1.9057	2.9	0.1832	2.6	0.88	1084.7	8.9	17.6	18.1	1083.1	6.8	13.6	14.2	1080.7	10.0	21.0	22.3	1080.7	21.0	100.4
Trial3_950	107	NA()	1.6	9.9922	1.0	3.9004	2.3	0.2828	2.1	0.91	1605.4	10.7	24.0	24.8	1613.8	6.9	15.9	16.6	1625.5	7.7	18.8	20.1	1625.5	18.8	98.8
Trial3_951	26	NA()	1.2	13.2034	2.5	1.8240	4.1	0.1747	3.2	0.78	1038.2	10.4	17.9	18.4	1054.1	9.8	15.2	15.7	1088.1	20.6	27.7	28.7	1088.1	27.7	95.4
Trial3_952	95	NA()	1.3	16.4561	2.0	0.8085	3.5	0.0965	2.9	0.83	594.1	5.7	10.3	10.6	601.6	5.6	9.8	10.2	631.0	15.8	25.4	26.7	594.1	10.3	94.1
Trial3_953	35	NA()	2.3	5.0247	1.0	14.2732	2.6	0.5204	2.4	0.92	2700.8	12.8	35.8	37.1	2768.0	8.2	18.7	19.5	2818.1	10.5	18.4	19.4	2818.1	18.4	95.8
Trial3_954	78	NA()	2.3	18.5607	3.3	0.3074	4.6	0.0414	3.3	0.70	261.5	2.9	4.8	5.0	272.2	4.3	6.0	6.2	365.9	31.3	37.6	38.6	261.5	4.8	NA
Trial3_955	63	NA()	1.8	12.7434	1.6	1.9851	3.2	0.1836	2.8	0.87	1086.3	9.5	17.9	18.4	1110.5	7.5	14.1	14.7	1158.8	11.5	21.6	22.9	1158.8	21.6	93.7
Trial3_956	203	NA()	1.3	9.8431	0.9	4.0345	1.5	0.2881	1.3	0.83	1632.3	7.9	23.2	24.1	1641.2	5.1	15.3	16.0	1653.4	5.6	18.0	19.3	1653.4	18.0	98.7
Trial3_957	82	NA()	9.8	11.1294	1.2	2.9898	2.7	0.2414	2.4	0.90	1394.2	10.4	21.7	22.4	1405.0	7.3	15.3	16.0	1422.4	9.0	19.8	21.1	1422.4	19.8	98.0
Trial3_958	12	NA()	0.9	13.4761	4.1	1.6554	5.2	0.1619	3.2	0.62	967.2	10.9	17.4	17.9	991.6	19.2	22.3	22.6	1047.1	56.1	59.1	59.6	1047.1	59.1	92.4
Trial3_959	117	NA()	1.9	8.7537	0.8	4.7808	2.0	0.3037	1.9	0.91	1709.4	10.4	25.0	25.9	1781.5	6.7	16.3	17.1	1867.9	7.1	18.1	19.4	1867.9	18.1	91.5
Trial3_960	178	NA()	2.9	8.9628	0.9	4.5769	1.7	0.2976	1.4	0.85	1679.6	8.6	24.0	24.9	1745.1	5.6	15.8	16.5	1825.2	5.9	17.7	19.0	1825.2	17.7	92.0
Trial3_961	440	NA()	6.0	3.9214	0.8	20.4066	1.3	0.5806	1.0	0.78	2951.4	2.0	35.9	37.4	3110.8	1.3	17.2	18.1	3216.1	1.6	14.6	15.8	3216.1	14.6	91.8
Trial3_962	246	NA()	1.8	10.7731	0.8	3.1148	1.5	0.2435	1.2	0.83	1404.8	6.3	20.2	20.9	1436.4	4.4	14.3	15.0	1484.3	5.0	18.2	19.6	1484.3	18.2	94.6
Trial3_963	222	NA()	1.9	11.9946	0.8	2.4381	1.7	0.2122	1.5	0.88	1240.5	6.5	18.3	19.0	1253.9	4.7	13.6	14.2	1277.9	5.8	18.8	20.3	1277.9	18.8	97.1
Trial3_964	28	NA()	3.9	9.4823	1.6	4.4442	3.4	0.3058	2.9	0.87	1719.9	13.3	26.4	27.3	1720.6	9.2	17.3	18.0	1722.3	12.3	20.9	22.1	1722.3	20.9	99.9
Trial3_965	111	NA()	2.2	13.1148	1.3	1.7794	2.8	0.1693	2.5	0.89	1008.4	7.9	16.2	16.7	1038.0	6.3	13.1	13.7	1101.7	9.1	20.6	22.0	1101.7	20.6	91.5
Trial3_966	115	NA()	2.1	13.1163	1.2	1.8548	2.7	0.1765	2.4	0.89	1048.0	8.0	16.7	17.2	1065.1	6.2	13.2	13.8	1101.4	8.9	20.4	21.8	1101.4	20.4	95.1
Trial3_967	174	NA()	7.2	18.8016	1.9	0.4572	3.4	0.0624	2.8	0.82	390.0	3.1	6.5	6.7	382.3	3.4	6.6	6.9	336.8	16.1	26.3	27.7	390.0	6.5	NA
Trial3_968	203	NA()	2.2	9.4537	0.9	4.5109	1.5	0.3094	1.2	0.81	1737.9	7.7	24.3	25.2	1733.0	4.9	15.5	16.3	1727.9	5.4	17.8	19.1	1727.9	17.8	100.6
Trial3_969	105	NA()	2.1	13.9711	1.4	1.6810	2.9	0.1704	2.5	0.88	1014.3	7.8	16.2	16.8	1001.3	6.1	12.8	13.4	973.9	9.7	21.1	22.5	973.9	21.1	104.2
Trial3_970	64	NA()	0.5	8.5271	1.0	5.3516	2.6	0.3311	2.3	0.91	1843.8	11.8	27.0	27.9	1877.1	7.5	16.9	17.6	1915.1	8.6	18.6	19.9	1915.1	18.6	96.3
Trial3_971	50	NA()	1.7	19.1882	3.9	0.2869	5.0	0.0399	3.2	0.64	252.5	2.9	4.8	4.9	256.1	6.2	7.3	7.5	290.5	55.9	59.7	60.4	252.5	4.8	NA
Trial3_972	25	NA()	1.5	13.7569	2.7	1.6356	4.2	0.1633	3.2	0.76	974.9	9.7	16.8	17.2	984.0	9.4	14.6	15.0	1005.3	21.0	28.1	29.2	1005.3	28.1	97.0
Trial3_973	41	NA()	0.9	5.2484	1.0	14.2042	2.5	0.5409	2.3	0.92	2787.3	12.5	36.5	37.8	2763.4	8.0	18.6	19.4	2746.8	10.4	18.4	19.4	2746.8	18.4	101.5
Trial3_974	57	NA()	0.8	5.5379	1.0	12.0521	2.3	0.4843	2.1	0.91	2545.9	12.4	34.2	35.4	2608.5	7.6	18.3	19.1	2658.1	9.2	17.8	18.9	2658.1	17.8	95.8
Trial3_975	115	NA()	3.0	13.5981	1.3	1.7046	2.8	0.1682	2.5	0.89	1002.1	7.9	16.1	16.6	1010.2	6.2	12.9	13.5	1028.8	9.3	20.8	22.2	1028.8	20.8	97.4
Trial3_976	69	NA()	2.4	4.5248	1.0	17.3153	2.0	0.5685	1.7	0.87	2901.6	11.0	37.1	38.5	2952.5	7.1	18.5	19.3	2988.0	9.3	17.5	18.5	2988.0	17.5	97.1
Trial3_977	242	NA()	9.8	13.1136	0.9	1.9846	1.8	0.1888	1.6	0.88	1115.1	5.7	16.5	17.1	1110.3	4.3	12.7	13.3	1101.8	6.0	19.4	20.8	1101.8	19.4	101.2
Trial3_978	477	NA()	1.5	12.4086	0.8	2.0424	1.3	0.1839	1.0	0.77	1088.2	1.0	15.2	15.8	1129.8	1.4	12.2	12.8	1211.4	3.7	18.5	20.0	1211.4	18.5	89.8
Trial3_979	64	NA()	1.0	13.2827	1.7	1.7747	3.3	0.1710	2.9	0.86	1017.9	9.2	17.0	17.5	1036.3	7.5	13.7	14.2	1076.2	12.1	22.1	23.4	1076.2	22.1	94.6
Trial3_980	103	NA()	2.2	11.6207	1.1	2.7466	2.6	0.2316	2.3	0.90	1342.8	10.0	20.9	21.6	1341.2	6.9	14.9	15.5	1339.4	8.5	19.7	21.0	1339.4	19.7	100.3
Trial3_981	24	NA()	0.8	7.5972	1.5	7.0108	3.2	0.3865	2.9	0.89	2106.4	14.4	30.8	31.8	2112.8	9.3	18.3	19.0	2119.8	11.7	19.9	21.0	2119.8	19.9	99.4
Trial3_982	106	NA()	1.0	8.6276	0.9	5.3354	2.1	0.3340	1.9	0.91	1857.8	11.4	27.0	27.9	1874.6	7.0	16.7	17.4	1894.0	7.7	18.2	19.5	1894.0	18.2	98.1
Trial3_983	156	NA()	1.9	12.6583	1.0	2.1005	2.3	0.1929	2.1	0.90	1137.2	8.1	17.8	18.3	1149.0	5.9	13.6	14.1	1172.1	7.4	19.7	21.1	1172.1	19.7	97.0
Trial3_984	21	NA()	1.9	13.3748	2.8	1.9604	4.3	0.1902	3.2	0.76	1122.7	11.5	19.4	19.9	1102.0	10.7	16.0	16.5	1062.3	22.6	29.2	30.2	1062.3	29.2	105.7
Trial3_985	229	NA()	3.6	12.5166	0.9	2.1155	1.9	0.1921	1.7	0.89	1132.9	6.5	17.0	17.6	1153.9	4.8	13.1	13.7	1194.3	6.0	19.1	20.5	1194.3	19.1	94.9
Trial3_986	40	NA()	0.8	10.8595	1.6	3.2679	3.3	0.2575	2.9	0.87	1477.0	12.6	23.6	24.3	1473.4	8.9	16.4	17.1	1469.1	12.3	21.4	22.5	1469.1	21.4	100.5
Trial3_987	49	NA()	1.6	13.8859	2.0	1.6413	3.6	0.1654	3.0	0.84	986.6	9.7	16.9	17.4	986.2	8.1	13.8	14.3	986.4	14.8	23.9	25.1	986.4	23.9	100.0
Trial3_988	61	NA()	0.9	16.5374	2.5	0.8077	4.0	0.0969	3.1	0.79	596.3	6.4	10.8	11.1	601.2	6.7	10.5	10.8	620.4	20.7	28.7	29.8	596.3	10.8	96.1
Trial3_989	300	NA()	1.9	5.3193	0.8	12.6917	1.3	0.4899	1.0	0.78	2570.0	6.1	32.7	34.0	2657.0	3.8	17.1	18.0	2724.7	4.5	15.8	17.0	2724.7	15.8	94.3
Trial3_990	31	NA()	1.5	18.8455	4.3	0.2903	5.4	0.0397	3.3	0.61	251.0	3.4	5.0	5.1	258.8	7.7	8.7	8.8	331.5	69.6	72.6	73.1	251.0	5.0	NA
Trial3_991	136	NA()	2.0	11.2880	0.9	3.0963	2.2	0.2536	1.9	0.90	1457.0	9.9	22.1	22.8	1431.8	6.5	15.1	15.8	1395.3	7.4	19.2	20.5	1395.3	19.2	104.4
Trial3_992	182	NA()	2.1	4.8430	0.9	14.5251	1.4	0.5104	1.1	0.77	2658.4	9.1	34.2	35.6	2784.6	5.6	17.8	18.6	2878.1	6.8	16.4	17.6	2878.1	16.4	92.4
Trial3_993	303	NA()	0.7	18.6036	1.6	0.3972	3.0	0.0536	2.5	0.84	336.7	1.9	5.3	5.5	339.6	2.3	5.6	5.9	360.7	12.4	24.2	25.7	336.7	5.3	NA
Trial3_994	86	NA()	0.6	10.7283	1.1	3.3907	2.6	0.2639	2.3	0.90	1510.0	11.4	23.3	24.1	1502.2	7.5	15.8	16.5	1492.2	8.7	19.5	20.8	1492.2	19.5	101.2
Trial3_995	132	NA()	2.1	12.2048	1.0	2.4692	2.4	0.2187	2.1	0.90	1274.8	9.5	20.0	20.6	1263.1	6.6	14.4	15.0	1244.0	7.8	19.7	21.1	1244.0	19.7	102.5
Trial3_996	174	NA()	3.1	13.5340	1.1	1.8004	2.3	0.1768	2.1	0.89	1049.5	7.4	16.4	17.0	1045.6	5.5	12.8	13.4	1038.4	7.7	20.1	21.5	1038.4	20.1	101.1
Trial3_997	84	NA()	1.4	5.3767	1.0	12.8081	1.9	0.4997	1.7	0.86	2612.4	11.8	34.6	35.9	2665.6	7.4	18.3	19.1	2707.0	9.2	17.8	18.8	2707.0	17.8	96.5
Trial3_998	36	NA()	0.8	8.8633	1.4	4.9705	3.1	0.3197	2.8	0.89	1788.1	13.7	27.3	28.2	1814.3	9.1	17.5	18.2	1845.4	11.2	20.1	21.2	1845.4	20.1	96.9
Trial3_999	57	NA()	1.7	13.8317	1.8	1.6779	3.5	0.1684	2.9	0.85	1003.3	9.8	17.2	17.6	1000.2	8.0	13.8	14.3	994.4	13.7	23.2	24.5	994.4	23.2	100.9
Trial3_1000	80	NA()	2.2	9.5710	1.1	4.2729	2.5	0.2967	2.3	0.91	1675.1	12.2	25.5	26.3	1688.2	7.9	16.6	17.3	1705.2	8.8	19.1	20.4	1705.2	19.1	98.2
Trial3_1001	99	NA()	1.5	13.3704	1.4	1.7364	3.0	0.1685	2.6	0.88	1003.6	8.9	16.6	17.1	1022.1	6.9	13.4	13.9	1062.9	10.0	21.0	22.4	1062.9	21.0	94.4
Trial3_1002	23	NA()	0.4	5.5714	1.3	12.7587	3.0	0.5158	2.7	0.91	2681.3	14.2	36.2	37.4	2662.0	9.2	19.1	19.8	2648.1	12.1	19.5	20.5	2648.1	19.5	101.3
Trial3_1003	75	NA()	0.6	10.8875	1.2	3.1328	2.8	0.2475	2.5	0.90	1425.5	11.4	22.5	23.2	1440.8	7.9	15.7	16.4	1464.2	9.4	19.8	21.1	1464.2	19.8	97.4
Trial3_1004	109	NA()	1.8	12.8488	1.2	2.0104	2.7	0.1874	2.4	0.89	1107.4	9.0	17.9	18.4	1119.0	6.7	13.8	14.3	1142.5	9.0	20.4	21.7	1142.5	20.4	96.9
Trial3_1005	52	NA()	2.0	9.2806	1.2	4.5946	2.9	0.3094	2.6	0.90	1737.7	13.0	26.5	27.3	1748.3	8.4	17.0	17.7	1761.7	9.9	19.5	20.7	1761.7	19.5	98.6
Trial3_1006	258	NA()	3.7	13.3737	0.9	1.7888	1.8	0.1736	1.6	0.88	1031.8	5.5	15.5	16.0	1041.4	4.3	12.3	12.9	1062.4	6.5	19.6	21.1	1062.4	19.6	97.1
Trial3_1007	52	NA()	1.2	12.3093	1.6	2.3774	3.3	0.2123	2.8	0.87	1241.2	11.1	20.4	21.0	1235.8	8.3	15.1	15.7	1227.2	12.0	21.7	23.0	1227.2	21.7	101.1
Trial3_1008	169	NA()	1.4	5.3924	0.9	12.2819	1.4	0.4806	1.1	0.77	2529.7	9.6	33.1	34.4	2626.2	5.9	17.7	18.5	2702.2	7.1	16.8	17.9	2702.2	16.8	93.6
Trial3_1009	35	NA()	1.5	12.7805	2.1	1.9528	3.7	0.1811	3.1	0.83	1072.9	10.7	18.4	18.9	1099.4	9.0	14.9	15.4	1153.1	15.6	24.1	25.2	1153.1	24.1	93.0
Trial3_1010	71	NA()	0.9	9.1848	1.1	4.5878	2.6	0.3058	2.4	0.91	1719.8	12.2	25.9	26.8	1747.1	7.9	16.7	17.4	1780.7	8.8	19.0	20.2	1780.7	19.0	96.6
Trial3_1011	192	NA()	1.6	17.8859	1.7	0.5099	3.2	0.0662	2.7	0.84	413.1	3.3	6.9	7.1	418.4	3.5	7.0	7.3	448.8	13.6	24.6	26.0	413.1	6.9	92.0
Trial3_1012	28	NA()	1.0	18.6906	3.7	0.5092	4.9	0.0691	3.3	0.66	430.5	5.4	8.3	8.5	417.9	8.6	10.5	10.7	350.2	48.6	52.9	53.6	430.5	8.3	122.9
Trial3_1013	64	NA()	0.7	13.2089	1.6	1.8471	3.3	0.1770	2.8	0.87	1050.7	9.7	17.6	18.1	1062.4	7.7	14.0	14.5	1087.3	12.2	22.1	23.4	1087.3	22.1	96.6
Trial3_1014	85	NA()	0.8	17.8891	3.4	0.2156	4.7	0.0280	3.2	0.69	177.9	2.0	3.4	3.4	198.2	3.5	4.7	4.9	448.4	34.7	40.2	41.1	177.9	3.4	NA
Trial3_1015	215	NA()	1.1	19.0082	2.0	0.3655	3.4	0.0504	2.8	0.82	317.0	2.4	5.3	5.4	316.3	2.8	5.6	5.8	312.0	16.1	26.4	27.8	317.0	5.3	NA
Trial3_1016	137	NA()	1.5	9.8558	0.8	3.8638	2.0	0.2763	1.8	0.91	1572.7	9.6	23.2	24.0	1606.1	6.3	15.6	16.3	1651.0	6.8	18.4	19.7	1651.0	18.4	95.3
Trial3_1017	18	NA()	0.8	11.5868	2.6	2.5349	4.1	0.2131	3.2	0.78	1245.4	12.1	21.0	21.6	1282.1	11.1	17.1	17.6	1345.0	21.2	27.7	28.7	1345.0	27.7	92.6
Trial3_1018	42	NA()	1.9	11.2531	1.7	2.7282	3.4	0.2228	2.9	0.87	1296.5	11.3	21.1	21.7	1336.2	8.5	15.7	16.3	1401.2	11.9	21.3	22.5	1401.2	21.3	92.5
Trial3_1019	61	NA()	0.5	8.9685	1.1	4.8238	2.7	0.3139	2.5	0.91	1759.9	12.2	26.3	27.2	1789.1	7.9	16.9	17.6	1824.0	9.1	19.0	20.2	1824.0	19.0	96.5
Trial3_1020	222	NA()	3.3	9.4642	0.8	4.4760	1.4	0.3074	1.1	0.79	1727.8	7.5	24.1	25.1	1726.5	4.7	15.5	16.2	1725.8	5.3	17.7	19.0	1725.8	17.7	100.1
Trial3_1021	121	NA()	1.3	13.0586	1.2	1.9120	2.6	0.1812	2.3	0.89	1073.4	7.7	16.8	17.4	1085.3	5.9	13.2	13.8	1110.2	8.5	20.3	21.6	1110.2	20.3	96.7
Trial3_1022	25	NA()	0.6	9.7859	1.9	3.3998	3.6	0.2414	3.1	0.85	1394.0	11.8	22.4	23.0	1504.4	9.4	16.8	17.4	1664.2	13.6	21.8	22.9	1664.2	21.8	83.8
Trial3_1023	31	NA()	1.4	11.0684	1.9	2.9377	3.5	0.2359	3.0	0.85	1365.5	11.4	21.9	22.5	1391.7	8.9	16.1	16.7	1432.9	13.7	22.3	23.4	1432.9	22.3	95.3
Trial3_1024	127	NA()	1.9	12.2478	1.1	2.3251	2.4	0.2066	2.2	0.90	1210.9	8.2	18.6	19.2	1220.0	6.0	13.9	14.5	1237.1	7.8	19.7	21.0	1237.1	19.7	97.9

Trial3_1025	27	NA()	1.6	13.9427	2.5	1.6965	4.1	0.1716	3.2	0.78	1021.1	9.7	17.3	17.8	1007.2	9.1	14.5	15.0	978.1	19.8	27.3	28.4	978.1	27.3	104.4
Trial3_1026	87	NA()	0.8	18.3381	2.7	0.4740	4.1	0.0631	3.1	0.76	394.3	3.8	7.0	7.2	394.0	4.7	7.4	7.7	393.0	22.9	30.8	32.0	394.3	7.0	NA
Trial3_1027	267	NA()	10.7	13.0881	0.8	1.8830	1.7	0.1788	1.5	0.87	1060.5	4.8	15.6	16.2	1075.1	3.8	12.3	12.9	1105.7	5.6	19.2	20.7	1105.7	19.2	95.9
Trial3_1028	214	NA()	4.0	13.4979	1.0	1.7263	2.1	0.1691	1.8	0.89	1007.0	5.8	15.3	15.8	1018.4	4.5	12.3	12.8	1043.8	6.6	19.7	21.2	1043.8	19.7	96.5
Trial3_1029	71	NA()	3.0	13.6820	1.6	1.6978	3.2	0.1686	2.8	0.87	1004.1	8.8	16.6	17.1	1007.7	7.1	13.4	13.9	1016.4	12.0	22.1	23.5	1016.4	22.1	98.8
Trial3_1030	100	NA()	1.8	13.4940	1.4	1.7376	2.9	0.1701	2.6	0.88	1012.8	8.2	16.4	16.9	1022.6	6.4	13.1	13.6	1044.4	9.7	21.0	22.4	1044.4	21.0	97.0
Trial3_1031	55	NA()	1.9	8.3707	1.1	5.6458	2.7	0.3429	2.4	0.91	1900.7	12.5	27.9	28.8	1923.1	7.9	17.2	18.0	1948.2	9.2	18.9	20.1	1948.2	18.9	97.6
Trial3_1032	61	NA()	1.2	5.3007	1.0	13.0562	2.2	0.5022	2.0	0.90	2623.1	12.0	34.8	36.0	2683.7	7.4	18.3	19.1	2730.5	9.2	17.8	18.8	2730.5	17.8	96.1
Trial3_1033	91	NA()	1.8	10.8616	1.1	3.2129	2.6	0.2532	2.3	0.90	1455.0	10.7	22.4	23.2	1460.3	7.2	15.5	16.2	1468.8	8.3	19.4	20.7	1468.8	19.4	99.1
Trial3_1034	74	NA()	2.9	8.1327	1.0	6.3849	2.3	0.3768	2.1	0.91	2061.2	12.4	29.5	30.5	2030.2	7.6	17.3	18.1	1999.6	8.9	18.6	19.8	1999.6	18.6	103.1
Trial3_1035	233	NA()	2.7	12.6906	0.9	2.1394	1.8	0.1970	1.6	0.88	1159.2	6.3	17.2	17.9	1161.6	4.6	13.1	13.7	1167.1	6.0	19.2	20.6	1167.1	19.2	99.3
Trial3_1036	80	NA()	0.6	5.1725	1.0	13.8485	1.9	0.5198	1.7	0.86	2698.1	11.4	35.3	36.6	2739.4	7.2	18.3	19.1	2770.7	9.1	17.6	18.7	2770.7	17.6	97.4
Trial3_1037	31	NA()	1.7	11.2043	1.9	2.8291	3.6	0.2300	3.0	0.85	1334.5	12.0	21.9	22.5	1363.3	9.2	16.2	16.8	1409.5	13.6	22.3	23.4	1409.5	22.3	94.7
Trial3_1038	61	NA()	1.4	13.3193	1.7	1.8304	3.3	0.1769	2.9	0.86	1050.0	9.6	17.5	18.0	1056.4	7.7	13.9	14.5	1070.6	12.5	22.3	23.6	1070.6	22.3	98.1
Trial3_1039	86	NA()	2.1	19.6339	3.5	0.2286	4.8	0.0326	3.2	0.67	206.6	2.3	3.8	3.9	209.0	4.1	5.3	5.4	237.8	43.4	48.3	49.1	206.6	3.8	NA
Trial3_1040	153	NA()	2.1	7.9920	0.9	5.9975	1.7	0.3478	1.4	0.85	1924.0	9.7	27.0	28.0	1975.5	6.0	16.5	17.3	2030.5	6.3	17.5	18.8	2030.5	17.5	94.8
Trial3_1041	490	NA()	0.8	19.9266	1.8	0.2032	3.1	0.0294	2.5	0.82	186.7	0.3	2.8	2.9	187.8	1.1	3.2	3.4	203.6	15.1	26.1	27.6	186.7	2.8	NA
Trial3_1042	62	NA()	1.9	12.6769	1.6	2.1001	3.2	0.1932	2.8	0.87	1138.6	9.8	18.6	19.2	1148.9	7.5	14.3	14.9	1169.2	11.3	21.4	22.7	1169.2	21.4	97.4
Trial3_1043	84	NA()	1.2	12.4601	1.3	2.2112	2.9	0.1999	2.6	0.89	1174.8	9.4	18.8	19.4	1184.6	7.1	14.3	14.8	1203.3	9.9	20.7	22.0	1203.3	20.7	97.6
Trial3_1044	137	NA()	1.7	13.3531	1.2	1.8402	2.6	0.1783	2.3	0.89	1057.7	7.6	16.6	17.2	1059.9	5.8	13.0	13.6	1065.5	8.3	20.3	21.7	1065.5	20.3	99.3
Trial3_1045	204	NA()	1.5	11.1733	0.9	3.1012	1.7	0.2514	1.5	0.86	1445.8	7.4	21.0	21.7	1433.0	5.0	14.5	15.2	1414.8	5.8	18.5	19.9	1414.8	18.5	102.2
Trial3_1046	575	NA()	5.6	5.3330	0.8	10.7116	1.3	0.4145	1.0	0.78	2235.4	2.4	28.7	29.9	2498.4	1.6	16.5	17.4	2720.4	1.8	15.3	16.5	2720.4	15.3	82.2
Trial3_1047	76	NA()	0.8	10.6842	1.2	3.2873	2.7	0.2548	2.5	0.90	1463.4	11.1	22.7	23.5	1478.0	7.6	15.8	16.4	1500.0	9.1	19.6	20.9	1500.0	19.6	97.6
Trial3_1048	130	NA()	1.3	10.8533	0.9	3.2201	2.2	0.2536	2.0	0.90	1456.9	9.7	22.0	22.7	1462.0	6.5	15.2	15.8	1470.2	7.2	18.9	20.3	1470.2	18.9	99.1
Trial3_1049	22	NA()	1.5	9.6308	1.9	4.0548	3.6	0.2834	3.1	0.85	1608.2	13.5	25.5	26.2	1645.3	10.0	17.6	18.2	1693.7	14.3	22.2	23.3	1693.7	22.2	95.0
Trial3_1050	162	NA()	1.8	13.7765	1.1	1.7127	2.4	0.1712	2.2	0.89	1018.7	7.2	16.0	16.5	1013.3	5.5	12.6	13.2	1002.4	7.9	20.3	21.7	1002.4	20.3	101.6
Trial3_1051	37	NA()	0.7	11.0754	1.7	2.9716	3.4	0.2388	2.9	0.86	1380.5	11.9	22.3	23.0	1400.4	8.8	16.1	16.7	1431.7	12.4	21.5	22.7	1431.7	21.5	96.4
Trial3_1052	221	NA()	2.1	18.3528	1.9	0.3975	3.3	0.0529	2.7	0.83	332.5	2.4	5.5	5.7	339.8	2.8	5.9	6.1	391.2	14.5	25.3	26.7	332.5	5.5	NA
Trial3_1053	67	NA()	1.6	16.8951	2.5	0.7318	4.0	0.0897	3.1	0.78	553.8	6.0	10.0	10.3	557.6	6.7	10.1	10.4	574.0	23.4	30.8	31.9	553.8	10.0	96.5
Trial3_1054	633	NA()	7.1	13.3899	0.8	1.8359	1.3	0.1784	1.0	0.78	1058.1	2.3	15.0	15.6	1058.4	1.9	11.8	12.4	1060.0	3.6	18.9	20.4	1060.0	18.9	99.8
Trial3_1055	61	NA()	1.1	5.1664	1.0	13.7417	2.2	0.5151	2.0	0.89	2678.5	12.4	35.4	36.7	2732.1	7.8	18.5	19.3	2772.6	9.9	18.0	19.1	2772.6	18.0	96.6
Trial3_1056	62	NA()	1.6	9.0784	1.1	4.8638	2.7	0.3204	2.4	0.91	1791.6	12.8	26.9	27.8	1796.0	8.2	17.0	17.7	1801.9	9.5	19.3	20.5	1801.9	19.3	99.4
Trial3_1057	23	NA()	0.4	5.3709	1.2	12.6983	3.0	0.4949	2.7	0.91	2591.7	14.3	35.4	36.6	2657.5	9.1	19.0	19.8	2708.8	11.5	19.1	20.1	2708.8	19.1	95.7
Trial3_1058	39	NA()	1.3	12.9480	2.0	1.8974	3.7	0.1783	3.1	0.83	1057.5	10.3	18.0	18.5	1080.2	8.6	14.6	15.1	1127.2	14.9	23.6	24.8	1127.2	23.6	93.8

Trial 4: CP-40 Element2 TC							Isotope ratios					Apparent ages (Ma)													
Analysis	U	206Pb	U/Th	206Pb*	±	207Pb*	±	206Pb*	±	error	206Pb*	±1s(a)	±1s(b)	±1s(c)	207Pb*	±1s(a)	±1s(b)	±1s(c)	206Pb*	±1s(a)	±1s(b)	±1s(c)	Best age	±	Conc
	(ppm)	204Pb		207Pb*	(%)	235U*	(%)	238U	(%)	corr.	238U*	(Ma)	(Ma)	(Ma)	235U	(Ma)	(Ma)	(Ma)	207Pb*	(Ma)	(Ma)	(Ma)	(Ma)	(Ma)	(%)
Trial4_001	23	NA()	1.7	12.9592	3.1	2.0058	4.8	0.1886	3.6	0.76	1113.8	10.2	20.6	21.1	1117.5	11.0	17.7	18.1	1125.5	25.5	33.1	34.0	1125.5	33.1	99.0
Trial4_002	330	NA()	2.3	12.3656	0.8	2.2878	1.6	0.2053	1.4	0.86	1203.6	4.0	19.7	20.2	1208.5	3.1	14.8	15.4	1218.3	4.8	21.4	22.7	1218.3	21.4	98.8
Trial4_003	223	NA()	24.0	8.8919	0.9	5.1590	1.6	0.3329	1.3	0.84	1852.2	7.4	29.2	30.0	1845.9	4.6	18.0	18.7	1839.6	5.3	19.9	21.1	1839.6	19.9	100.7
Trial4_004	41	NA()	1.6	10.1463	1.8	3.7751	3.7	0.2779	3.3	0.88	1580.9	11.6	27.2	27.8	1587.5	8.6	18.6	19.2	1597.0	12.9	23.6	24.7	1597.0	23.6	99.0
Trial4_005	131	NA()	1.0	19.2777	3.6	0.2510	5.1	0.0351	3.6	0.71	222.5	2.0	4.3	4.4	227.4	3.4	5.4	5.5	279.9	31.1	39.5	40.5	222.5	4.3	NA
Trial4_006	108	NA()	2.2	12.6164	1.5	2.1334	3.2	0.1953	2.8	0.89	1150.0	8.2	20.2	20.7	1159.7	6.4	15.6	16.1	1178.7	9.9	23.2	24.4	1178.7	23.2	97.6
Trial4_007	37	NA()	1.0	5.0083	1.1	14.6451	3.0	0.5322	2.8	0.93	2750.7	11.8	41.0	42.2	2792.5	7.6	20.9	21.6	2823.5	9.8	19.9	20.9	2823.5	19.9	97.4
Trial4_008	156	NA()	2.2	13.2893	1.3	1.7602	2.9	0.1697	2.6	0.89	1010.6	6.7	17.7	18.2	1030.9	5.3	14.3	14.8	1075.2	8.3	22.9	24.1	1075.2	22.9	94.0
Trial4_009	287	NA()	1.4	20.2416	2.9	0.1850	4.5	0.0272	3.4	0.75	172.8	1.0	3.1	3.2	172.3	2.1	3.9	4.0	167.0	28.3	37.6	38.6	172.8	3.1	NA
Trial4_010	58	NA()	1.3	11.1550	1.7	3.0194	3.6	0.2444	3.1	0.88	1409.5	10.5	24.5	25.2	1412.5	7.7	17.4	18.0	1418.0	11.2	23.2	24.3	1418.0	23.2	99.4
Trial4_011	54	NA()	0.9	5.2382	1.0	13.7379	2.7	0.5222	2.5	0.93	2708.3	11.1	40.3	41.5	2731.8	7.1	20.6	21.4	2750.0	9.1	19.7	20.7	2750.0	19.7	98.5
Trial4_012	34	NA()	0.8	11.2670	2.2	3.0242	4.0	0.2472	3.4	0.84	1424.2	10.6	24.8	25.4	1413.7	9.3	18.2	18.7	1398.9	17.1	26.5	27.5	1398.9	26.5	101.8
Trial4_013	123	NA()	0.6	5.1314	0.9	14.1996	1.7	0.5287	1.5	0.85	2735.9	9.0	40.1	41.3	2763.1	5.7	20.2	21.0	2783.8	7.3	18.8	19.9	2783.8	18.8	98.3
Trial4_014	142	NA()	4.7	13.6023	1.4	1.6448	3.1	0.1623	2.7	0.88	969.7	6.4	17.0	17.5	987.5	5.4	14.0	14.5	1028.2	9.5	23.5	24.7	1028.2	23.5	94.3
Trial4_015	33	NA()	1.1	13.1127	2.7	1.9001	4.4	0.1808	3.5	0.80	1071.3	9.3	19.6	20.1	1081.1	9.3	16.5	16.9	1102.0	20.5	29.5	30.5	1102.0	29.5	97.2
Trial4_016	88	NA()	3.6	13.0385	1.7	1.8902	3.5	0.1788	3.1	0.87	1060.5	7.7	18.8	19.3	1077.6	6.4	15.0	15.5	1113.3	11.2	24.0	25.2	1113.3	24.0	95.3
Trial4_017	216	NA()	1.5	16.2364	1.6	0.8255	3.2	0.0973	2.8	0.86	598.3	3.4	10.6	10.9	611.1	3.7	10.1	10.5	659.9	11.4	25.4	26.7	598.3	10.6	90.7
Trial4_018	602	NA()	1.9	9.6837	0.8	4.1073	1.3	0.2886	1.0	0.78	1634.5	2.6	25.4	26.2	1655.8	1.9	16.8	17.5	1683.6	2.6	19.8	21.0	1683.6	19.8	97.1
Trial4_019	78	NA()	0.8	5.9101	0.9	10.3645	2.5	0.4445	2.3	0.93	2370.6	10.5	36.3	37.4	2467.9	6.5	20.0	20.8	2549.8	7.6	19.3	20.4	2549.8	19.3	93.0
Trial4_020	203	NA()	1.1	17.6687	2.0	0.5518	3.7	0.0707	3.1	0.83	440.6	2.7	8.0	8.2	446.2	3.4	8.1	8.4	475.9	15.1	27.9	29.2	440.6	8.0	92.6
Trial4_021	308	NA()	2.6	9.3082	0.8	4.4349	1.3	0.2995	1.0	0.78	1689.0	4.5	26.4	27.2	1718.9	3.1	17.3	17.9	1756.3	3.9	19.8	21.0	1756.3	19.8	96.2
Trial4_022	84	NA()	2.2	12.5489	1.6	2.2805	3.4	0.2077	3.0	0.88	1216.3	8.4	21.2	21.7	1206.3	6.5	15.8	16.4	1189.3	10.2	23.3	24.5	1189.3	23.3	102.3
Trial4_023	193	NA()	1.4	20.1988	3.0	0.2553	4.6	0.0374	3.4	0.75	236.8	1.6	4.4	4.5	230.8	2.7	5.0	5.2	171.9	26.0	35.9	37.0	236.8	4.4	NA
Trial4_024	78	NA()	2.2	14.1195	2.0	1.5401	3.8	0.1578	3.2	0.85	944.4	7.4	17.1	17.5	946.5	6.6	14.2	14.7	952.4	13.4	25.5	26.7	952.4	25.5	99.2
Trial4_025	130	NA()	1.3	9.2882	1.0	4.6903	2.3	0.3161	2.1	0.91	1770.7	8.7	28.5	29.3	1765.5	5.6	18.0	18.7	1760.2	6.6	20.5	21.7	1760.2	20.5	100.6
Trial4_026	392	NA()	1.8	18.1182	1.5	0.5315	2.9	0.0699	2.5	0.86	435.4	1.1	7.5	7.7	432.8	1.8	7.5	7.7	420.0	10.2	25.8	27.2	435.4	7.5	103.7
Trial4_027	187	NA()	0.8	9.3987	0.9	4.7952	1.8	0.3270	1.6	0.88	1823.9	7.6	28.8	29.7	1784.1	4.8	17.9	18.5	1738.6	5.7	20.3	21.4	1738.6	20.3	104.9
Trial4_028	64	NA()	0.7	10.5843	1.5	3.3021	3.4	0.2536	3.0	0.89	1457.0	9.9	24.9	25.5	1481.6	7.3	17.6	18.1	1517.7	10.4	22.6	23.7	1517.7	22.6	96.0
Trial4_029	98	NA()	1.1	11.6109	1.3	2.8311	3.1	0.2385	2.7	0.90	1379.0	8.8	23.5	24.1	1363.8	6.3	16.6	17.2	1341.0	8.7	22.2	23.4	1341.0	22.2	102.8
Trial4_030	235	NA()	1.3	8.8370	0.8	5.1220	1.5	0.3284	1.2	0.82	1830.8	6.3	28.6	29.5	1839.8	4.0	17.9	18.5	1850.8	4.8	19.8	20.9	1850.8	19.8	98.9
Trial4_031	49	NA()	0.9	5.3444	1.0	12.6434	2.9	0.4903	2.7	0.93	2572.0	11.5	38.9	40.0	2653.5	7.2	20.6	21.3	2716.9	9.0	19.7	20.6	2716.9	19.7	94.7
Trial4_032	142	NA()	2.2	9.4242	0.9	4.5318	2.2	0.3099	2.0	0.91	1740.2	8.9	28.2	29.0	1736.8	5.8	18.0	18.6	1733.6	6.7	20.6	21.8	1733.6	20.6	100.4
Trial4_033	217	NA()	1.3	5.3623	0.9	12.5963	1.4	0.4901	1.1	0.77	2571.1	7.1	37.8	39.0	2649.9	4.4	19.8	20.5	2711.4	5.4	18.3	19.4	2711.4	18.3	94.8
Trial4_034	214	NA()	1.3	12.8227	1.1	1.9968	2.4	0.1858	2.2	0.89	1098.5	5.7	18.6	19.1	1114.4	4.5	14.6	15.1	1146.6	6.8	22.1	23.4	1146.6	22.1	95.8
Trial4_035	71	NA()	1.1	10.0502	1.4	3.9206	3.2	0.2859	2.9	0.90	1621.0	10.5	27.2	27.9	1617.9	7.1	18.0	18.7	1614.7	9.2	21.8	22.9	1614.7	21.8	100.4
Trial4_036	59	NA()	1.3	11.1475	1.6	3.1149	3.5	0.2519	3.1	0.89	1448.5	10.6	25.1	25.7	1436.4	7.7	17.5	18.1	1419.3	11.1	23.1	24.2	1419.3	23.1	102.1
Trial4_037	43	NA()	0.8	12.8623	2.3	1.9859	4.2	0.1853	3.5	0.83	1096.1	10.0	20.3	20.8	1110.7	8.7	16.3	16.8	1140.4	16.3	26.7	27.7	1140.4	26.7	96.1
Trial4_038	86	NA()	1.6	13.3589	1.7	1.8032	3.6	0.1748	3.1	0.87	1038.4	8.7	18.9	19.4	1046.6	7.0	15.1	15.6	1064.7	11.7	24.3	25.5	1064.7	24.3	97.5
Trial4_039	227	NA()	0.4	19.5021	2.4	0.3378	4.1	0.0478	3.3	0.80	301.0	2.2	5.6	5.8	295.5	3.1	6.1	6.3	253.3	21.3	32.4	33.6	301.0	5.6	NA
Trial4_040	13	NA()	3.1	12.4342	3.7	1.9201	4.9	0.1732	3.2	0.66	1029.9	11.1	20.0	20.5	1088.1	16.5	21.5	21.8	1207.4	43.0	47.9	48.4	1207.4	47.9	85.3
Trial4_041	310	NA()	1.1	8.8771	0.8	4.7454	1.3	0.3057	1.1	0.79	1719.3	5.4	27.0	27.8	1775.3	3.5	17.5	18.2	1842.6	4.1	19.6	20.8	1842.6	19.6	93.3
Trial4_042	75	NA()	1.1	8.6703	1.2	5.3972	2.9	0.3395	2.6	0.91	1884.5	11.3	30.8	31.6	1884.4	7.2	19.0	19.6	1885.1	8.7	21.0	22.1	1885.1	21.0	100.0
Trial4_043	74	NA()	0.6	5.0554	1.0	14.8901	2.3	0.5462	2.1	0.91	2809.3	10.5	41.3	42.5	2808.2	6.8	20.7	21.4	2808.2	9.0	19.5	20.5	2808.2	19.5	100.0
Trial4_044	30	NA()	0.9	7.1209	1.5	7.4208	3.6	0.3834	3.2	0.90	2092.2	13.1	33.9	34.8	2163.5	8.6	20.3	20.9	2232.5	10.7	21.3	22.3	2232.5	21.3	93.7
Trial4_045	530	NA()	4.0	13.6276	0.8	1.7123	1.3	0.1693	1.0	0.77	1008.3	0.7	16.4	16.9	1013.1	1.4	13.2	13.7	1024.5	4.2	21.9	23.2	1024.5	21.9	98.4
Trial4_046	51	NA()	1.9	11.2414	1.8	2.9431	3.7	0.2401	3.2	0.87	1387.0	10.7	24.3	25.0	1393.1	8.2	17.6	18.1	1403.2	12.7	24.0	25.1	1403.2	24.0	98.8
Trial4_047	71	NA()	1.7	10.9624	1.6	2.7378	3.5	0.2178	3.1	0.89	1270.1	9.9	22.5	23.1	1338.8	7.7	17.1	17.6	1451.2	10.9	22.9	24.1	1451.2	22.9	87.5
Trial4_048	33	NA()	1.2	13.7625	2.8	1.7710	4.6	0.1769	3.6	0.78	1049.8	9.7	19.6	20.0	1034.9	9.9	16.6	17.0	1004.5	23.3	31.7	32.7	1004.5	31.7	104.5
Trial4_049	118	NA()	1.5	13.3625	1.5	1.9186	3.2	0.1860	2.8	0.88	1099.8	8.1	19.5	20.0	1087.6	6.3	15.1	15.6	1064.1	10.3	23.7	24.9	1064.1	23.7	103.4
Trial4_050	62	NA()	1.4	13.6609	2.1	1.6543	4.0	0.1640	3.4	0.84	978.9	8.5	18.1	18.5	991.2	7.7	15.1	15.5	1019.5	15.4	26.5	27.6	1019.5	26.5	96.0
Trial4_051	62	NA()	8.3	9.9538	1.5	3.9399	3.3	0.2846	3.0	0.90	1614.3	11.3	27.4	28.2	1621.9	7.8	18.3	18.9	1632.7	10.2	22.2	23.3	1632.7	22.2	98.9
Trial4_052	204	NA()	0.7	18.1709	2.1	0.4884	3.8	0.0644	3.2	0.83	402.3	2.9	7.4	7.6	403.8	3.5	7.7	7.9	413.5	16.6	28.9	30.2	402.3	7.4	97.3
Trial4_053	61	NA()	1.8	13.1496	2.0	1.9750	3.8	0.1884	3.3	0.85	1112.9	9.4	20.2	20.7	1107.0	8.0	16.0	16.4	1096.3	15.0	26.0	27.1	1096.3	26.0	101.5
Trial4_054	162	NA()	2.1	13.1806	1.3	1.8885	2.8	0.1806	2.5	0.89	1070.3	7.1	18.7	19.2	1077.1	5.5	14.7	15.2	1091.6	8.1	22.7	24.0	1091.6	22.7	98.1
Trial4_055	374	NA()	2.9	5.6457	0.8	11.4248	1.3	0.4680	1.0	0.78	2474.8	4.3	36.3	37.4	2558.5	2.6	19.3	20.1	2626.1	3.2	17.9	19.0	2626.1	17.9	94.2
Trial4_056	291	NA()	3.3	13.7051	1.0	1.7325	2.1	0.1723	1.8	0.88	1024.7	4.4	17.2	17.7	1020.7	3.6	13.7	14.2	1013.0	6.0	22.3	23.6	1013.0	22.3	101.2
Trial4_057	76	NA()	0.9	5.5359	1.0	13.0136	2.3	0.5227	2.1	0.91	2710.7	10.8	40.3	41.4	2680.6	6.8	20.5	21.2	2658.7	8.8	19.7	20.7	2658.7	19.7	102.0
Trial4_058	28	NA()	1.4	12.2834	2.9	2.1863	4.6	0.1949	3.6	0.78	1147.7	10.4	21.1	21.6	1176.7	10.9	18.0	18.4	1231.4	23.8	31.6	32.5	1231.4	31.6	93.2
Trial4_059	18	NA()	1.9	13.0624	3.7	1.8898	4.9	0.1791	3.2	0.65	1062.1	10.4	20.1	20.5	1077.5	12.4	18.4	18.8	1109.6	30.8	37.4	38.2	1109.6	37.4	95.7
Trial4_060	42	NA()	2.2	13.0255	2.5	1.8660	4.3	0.1764	3.5	0.81	1047.0	9.3	19.3	19.8	1069.1	9.4	16.5	16.9	1115.3	20.9	29.8	30.7	1115.3	29.8	93.9
Trial4_061	44	NA()	1.3	12.5621	2.3	2.0468	4.1	0.1866	3.5	0.83	1102.7	9.6	20.2	20.7	1131.2	8.6	16.4	16.9	1187.2	16.5	26.7	27.7	1187.2	26.7	92.9
Trial4_062	79	NA()	2.1	17.8506	2.9	0.6817	4.6	0.0883	3.5	0.77	545.5	5.2	10.5	10.8	527.8	6.1	10.4	10.7	453.2	24.6	34.1	35.1	545.5	10.5	120.4
Trial4_063	211	NA()	2.3	17.1883	1.8	0.6874	3.4	0.0857	2.9	0.85	530.3	3.4	9.6	9.8	531.3	3.7	9.3	9.6	536.5	13.0	26.6	27.9	530.3	9.6	98.8
Trial4_064	13	NA()	1.3	12.5344	4.0	2.3236	5.1	0.2113	3.2	0.63	1235.9	12.0	23.1	23.6	1219.5	14.1	20.3	20.7	1191.6	33.2	39.3	40.0	1191.6	39.3	103.7
Trial4_065	384	NA()	1.5	19.4186	2.1	0.2856	3.7	0.0402	3.0	0.82	254.3	0.9	4.5	4.6	255.1	1.7	4.9	5.1	263.1	15.9	29.1	30.4	254.3	4.5	NA
Trial4_066	143	NA()	2.5	7.9354	0.9	6.5256	2.0	0.3757	1.8	0.90	2056.3	9.6	32.3	33.2	2049.4	5.8	19.0	19.6	2043.1	6.7	19.9	21.0	2043.1	19.9	100.6
Trial4_067	92	NA()	0.8	13.5225	1.7	1.7799	3.5	0.1746	3.1	0.87	1037.6	8.0	18.6	19.1	1038.1	6.5	14.8	15.3	1040.1	11.2	24.2	25.4	1040.1	24.2	99.8
Trial4_068	137	NA()	1.7	12.8466	1.3	2.1587	2.9	0.2012	2.6	0.89	1181.9	8.0	20.6	21.1	1167.8	5.9	15.4	15.9	1142.8	8.4	22.7	23.9	1142.8	22.7	103.4
Trial4_069	105	NA()	0.9	8.8500	1.0	5.3707	2.5	0.3449	2.3	0.91	1910.1	10.8	30.9	31.8	1880.2	6.7	18.8	19.4	1848.1	7.7	20.7	21.8	1848.1	20.7	103.4
Trial4_070	28	NA()	1.2	14.0035	3.2	1.6150	4.9	0.1641	3.6	0.75	979.5	9.5	18.5	18.9	976.1	10.7	16.7	17.1	969.2	27.4	34.9	35.8	969.2	34.9	101.1
Trial4_071	47	NA()	0.9	5.5424	1.1	12.4682	2.9	0.5014	2.7	0.93	2619.9	12.3	39.7	40.8	2640.3	7.6	20.7	21.4	2656.8	9.5	20.0	21.0	2656.8	20.0	98.6
Trial4_072	386	NA()	2.5	12.3547	0.8	2.1174	1.6	0.1898	1.4	0.85	1120.4	3.1	18.3	18.8	1154.5	2.6	14.4	14.9	1220.0	4.8	21.4	22.7	1220.0	21.4	91.8
Trial4_073	268	NA()	1.5	18.2580	1.9	0.4952	3.5	0.0656	2.9	0.84	409.6	2.4	7.4	7.6	408.4	3.1	7.6	7.8	402.8	15.6	28.4	29.7	409.6	7.4	101.7
Trial4_074	54	NA()	0.6	10.7836	1.7	3.2611	3.6	0.2552	3.2	0.88	1465.0	10.8	25.4	26.0	1471.8	8.0	17.8	18.4	1482.4	11.6	23.2	24.3	1482.4	23.2	98.8
Trial4_075	771	NA()	10.6	13.2791	0.8	1.8990	1.4	0.1830	1.1	0.80	1083.2	4.8	18.1	18.7	1080.7	3.5	14.1	14.6	1076.7	4.0	21.7	23.0	1076.7	21.7	100.6
Trial4_076	100	NA()	1.0	16.4707	2.4	0.8495	4.1	0.1015	3.3	0.82	623.3	5.2	11.6	11.9	624.4	5.7	11.1	11.5	629.1	18.6	29.4	30.5	623.3	11.6	99.1
Trial4_077	78	NA()	1.4	18.9735	3.9	0.2573	5.0	0.0354	3.2	0.64	224.4	2.3	4.5	4.6	232.5	5.5	6.9	7.1	316.1	55.1	60.2	60.8	224.4	4.5	NA
Trial4_078	29	NA()	0.9	13.8005	3.0	1.7457	4.7	0.1748	3.6	0.77	1038.5	9.3	19.2	19.7	1025.6	9.9	16.5	17.0	998.9	24.2	32.4	33.3	998.9	32.4	104.0
Trial4_079	671	NA()	4.1	8.1586	0.8	5.6145	1.3	0.3324	1.0	0.78	1849.9	3.9	28.4	29.3	1918.3	2.5	17.8	18.5	1994.0	2.7	19.1	20.2	1994.0	19.1	92.8
Trial4_080	79	NA()	0.9	16.6645	2.7	0.8129	4.4	0.0983	3.5	0.79	604.4	5.4	11.5	11.7	604.1	6.4	11.3	11.6	603.8	22.9	32.5	33.5	604.4	11.5	100.1
Trial4_081	273	NA()	2.1	18.2336	1.8	0.5172	3.4	0.0684	2.9	0.84	426.6	2.1	7.5	7.8	423.3	2.7	7.6	7.8	405.8	12.9	27.0	28.3	426.6	7.5	105.1
Trial4_082	153	NA()	0.9	16.5738	2.0	0.7978	3.7	0.0959	3.1	0.85	590.6	4.3	10.8	11.1	595.6	4.6	10.3	10.7	615.6	14.8	27.3	28.5	590.6	10.8	95.9
Trial4_083	149	NA()	2.6	5.0843	0.9	15.0146	1.5	0.5539	1.2	0.79	2841.4	8.7	41.2	42.4	2816.2	5.4	20.2	21.0	2798.9	7.0	18.7	19.7	2798.9	18.7	101.5
Trial4_084	63	NA()	0.8	5.4448	1.0	13.4276	2.5	0.5305	2.4	0.92	2743.5	11.5	40.8	42.0	2710.2	7.1	20.6	21.3	2686.2	9.0	19.7	20.7	2686.2	19.7	102.1
Trial4_085	101	NA()	0.5	8.9642	1.1	5.1040	2.6	0.3320	2.4	0.92	1848.0	10.8	30.2	31.0	1836.8	6.8	18.7	19.3	1824.9	7.8	20.8	21.9	1824.9	20.8	101.3
Trial4_086	298	NA()	3.7	9.2001	0.8	4.5891	1.3	0.3063	1.0	0.78	1722.7	5.9	27.1	28.0	1747.3	3.8	17.5	18.2	1777.6	4.3	19.8	21.0	1777.6	19.8	96.9
Trial4_087	61	NA()	1.2	9.3021	1.4	4.8424	3.2	0.3268	2.9	0.91	1823.1	12.0	30.3	31.1	1792.3	7.7	18.9	19.5	1757.5	9.6	21.6	22.7	1757.5	21.6	103.7
Trial4_088	24	NA()	1.4	13.2567	3.2	1.9331	4.9	0.1859	3.7	0.75	1099.3	10.5	20.6	21.1	1092.6	11.5	17.9	18.3	1080.1	27.5	34.8	35.7	1080.1	34.8	101.8
Trial4_089	197	NA()	2.5	9.5152	0.9	4.0683	2.0	0.2809	1.8	0.90	1595.8	8.1	26.1	26.8	1648.0	5.3	17.5	18.2	1716.0	5.7	20.3	21.5	1716.0	20.3	93.0
Trial4_090	130	NA()	0.8	8.8482	0.9	5.2519	2.3	0.3372	2.1	0.91	1873.1	10.1	30.2	31.1	1861.1	6.3	18.6	19.2	1848.5	7.0	20.4	21.6	1848.5	20.4	101.3
Trial4_091	300	NA()	1.3	18.9204	1.9	0.4500	3.5	0.0618	2.9	0.84	386.4	2.0	6.9	7.1	377.3	2.5	6.9	7.2	322.5	13.5	27.6	28.9	386.4	6.9	NA
Trial4_092	229	NA()	1.2	18.4897	2.0	0.5128	3.6	0.0688	3.0	0.84	428.9	2.7	7.8	8.0	420.3	3.1	7.7	8.0	374.5	14.3	27.8	29.1	428.9	7.8	114.5
Trial4_093	404	NA()	3.2	13.2806	0.8	1.9001	1.5	0.1831	1.3	0.84	1083.9	2.4	17.7	18.2	1081.1	2.3	13.8	14.4	1076.5	4.8	21.8	23.1	1076.5	21.8	100.7
Trial4_094	101	NA()	0.8	19.3392	3.9	0.2847	5.4	0.0400	3.7	0.68	252.5	2.5	5.0	5.1	254.4	4.3	6.3	6.4	272.5	37.4	44.6	45.4	252.5	5.0	NA
Trial4_095	51	NA()	1.3	11.2885	1.8	3.0551	3.7	0.2502	3.2	0.87	1439.7	10.8	25.1	25.7	1421.5	8.1	17.6	18.2	1395.2	12.2	23.7	24.8	1395.2	23.7	103.2
Trial4_096	111	NA()	1.7	13.2202	1.5	1.8884	3.3	0.1811	2.9	0.88	1073.2	7.8	19.0	19.5	1077.0	6.3	15.0	15.5	1085.6	10.3	23.6	24.8	1085.6	23.6	98.9
Trial4_097	69	NA()	2.2	6.2485	1.0	10.0458	2.6	0.4555	2.5	0.93	2419.5	11.9	37.3	38.3	2439.0	7.2	20.2	20.9	2456.1	8.6	19.9	20.9	2456.1	19.9	98.5

Trial4_098	75	NA()	1.7	12.5085	1.7	2.1670	3.6	0.1967	3.1	0.88	1157.5	9.2	20.7	21.2	1170.5	7.3	16.0	16.5	1195.6	11.8	24.0	25.2	1195.6	24.0	96.8
Trial4_099	16	NA()	1.5	12.3790	3.6	2.2236	4.8	0.1997	3.2	0.66	1173.8	11.3	22.0	22.5	1188.5	13.1	19.4	19.8	1216.2	30.3	36.8	37.5	1216.2	36.8	96.5
Trial4_100	260	NA()	1.9	8.6965	0.8	5.2398	1.4	0.3306	1.2	0.81	1841.5	6.8	28.9	29.7	1859.1	4.2	18.0	18.7	1879.7	4.7	19.7	20.9	1879.7	19.7	98.0
Trial4_101	178	NA()	1.3	13.5156	1.2	1.8574	2.7	0.1822	2.4	0.89	1078.7	6.7	18.7	19.2	1066.1	5.2	14.5	15.0	1041.1	8.0	22.9	24.1	1041.1	22.9	103.6
Trial4_102	78	NA()	0.9	5.4370	1.0	13.2182	2.3	0.5215	2.1	0.91	2705.4	11.1	40.3	41.4	2695.4	6.8	20.5	21.2	2688.6	8.6	19.5	20.5	2688.6	19.5	100.6
Trial4_103	151	NA()	0.9	8.7595	0.9	5.0638	2.1	0.3218	1.9	0.91	1798.7	9.7	29.2	30.0	1830.1	6.1	18.4	19.1	1866.7	6.6	20.2	21.4	1866.7	20.2	96.4
Trial4_104	48	NA()	1.1	5.3733	1.1	13.4959	2.9	0.5262	2.7	0.93	2725.3	12.4	40.9	42.0	2715.0	7.6	20.8	21.5	2708.0	9.7	20.0	21.0	2708.0	20.0	100.6
Trial4_105	45	NA()	1.5	13.7659	2.5	1.6840	4.3	0.1682	3.5	0.81	1002.2	9.2	18.7	19.1	1002.5	9.0	15.9	16.3	1004.0	20.3	29.6	30.6	1004.0	29.6	99.8
Trial4_106	37	NA()	1.6	7.6239	1.5	6.9982	3.5	0.3871	3.1	0.91	2109.5	13.3	34.2	35.1	2111.2	8.5	20.1	20.8	2113.6	10.7	21.4	22.5	2113.6	21.4	99.8
Trial4_107	103	NA()	1.0	10.8486	1.3	3.3302	2.9	0.2621	2.6	0.90	1500.8	10.3	25.6	26.3	1488.2	7.0	17.5	18.1	1471.0	8.9	22.0	23.2	1471.0	22.0	102.0
Trial4_108	57	NA()	1.4	11.3694	1.8	2.8584	3.7	0.2358	3.2	0.88	1364.8	11.0	24.2	24.8	1371.0	8.3	17.5	18.0	1381.5	12.5	23.9	25.0	1381.5	23.9	98.8
Trial4_109	73	NA()	0.8	13.0180	1.9	1.9439	3.7	0.1836	3.2	0.87	1086.7	9.2	19.8	20.3	1096.3	7.5	15.7	16.1	1116.5	12.8	24.7	25.9	1116.5	24.7	97.3
Trial4_110	22	NA()	2.0	10.0671	2.4	3.8934	4.3	0.2844	3.5	0.83	1613.5	13.2	28.3	29.0	1612.3	11.0	19.9	20.4	1611.6	18.7	27.2	28.1	1611.6	27.2	100.1
Trial4_111	232	NA()	13.5	13.4191	1.1	1.7389	2.5	0.1693	2.2	0.89	1008.3	5.8	17.4	17.8	1023.1	4.6	14.0	14.5	1055.6	7.2	22.5	23.8	1055.6	22.5	95.5
Trial4_112	314	NA()	2.0	13.6657	1.0	1.6624	2.1	0.1648	1.8	0.87	983.6	3.9	16.5	16.9	994.3	3.3	13.4	13.9	1018.8	6.1	22.3	23.6	1018.8	22.3	96.5
Trial4_113	87	NA()	0.8	11.0588	1.4	3.0595	3.2	0.2455	2.9	0.90	1415.2	10.0	24.4	25.0	1422.6	7.0	17.2	17.8	1434.5	9.1	22.2	23.3	1434.5	22.2	98.7
Trial4_114	43	NA()	1.6	12.3402	2.2	2.3142	4.1	0.2072	3.4	0.84	1213.9	10.3	21.9	22.5	1216.6	8.7	16.9	17.4	1222.3	15.8	26.2	27.2	1222.3	26.2	99.3
Trial4_115	125	NA()	1.3	8.7892	1.0	5.1109	2.4	0.3259	2.2	0.91	1818.7	10.1	29.5	30.4	1837.9	6.3	18.5	19.2	1860.6	7.1	20.4	21.6	1860.6	20.4	97.7
Trial4_116	330	NA()	1.7	13.5702	1.0	1.7410	2.0	0.1714	1.7	0.87	1020.0	3.6	16.9	17.4	1023.8	3.0	13.6	14.1	1033.0	5.6	22.2	23.5	1033.0	22.2	98.7
Trial4_117	50	NA()	0.6	5.1644	1.0	14.6660	2.8	0.5496	2.6	0.93	2823.4	11.7	41.7	42.9	2793.8	7.3	20.8	21.5	2773.3	9.4	19.8	20.8	2773.3	19.8	101.8
Trial4_118	73	NA()	1.3	12.7255	1.8	1.9822	3.7	0.1830	3.2	0.87	1083.5	8.7	19.5	20.0	1109.5	7.3	15.6	16.1	1161.6	12.5	24.5	25.6	1161.6	24.5	93.3
Trial4_119	121	NA()	2.8	9.3622	1.0	4.4803	2.5	0.3044	2.3	0.91	1712.9	9.9	28.2	28.9	1727.3	6.4	18.2	18.8	1745.7	7.3	20.7	21.9	1745.7	20.7	98.1
Trial4_120	105	NA()	1.6	13.6990	1.6	1.7969	3.4	0.1786	3.0	0.88	1059.4	8.2	19.0	19.5	1044.3	6.7	14.9	15.4	1013.9	11.7	24.5	25.6	1013.9	24.5	104.5
Trial4_121	74	NA()	1.4	13.2263	1.9	1.8820	3.7	0.1806	3.2	0.86	1070.3	8.9	19.4	19.9	1074.8	7.3	15.4	15.9	1084.7	12.5	24.7	25.8	1084.7	24.7	98.7
Trial4_122	42	NA()	0.9	6.1273	1.2	10.7272	3.1	0.4769	2.9	0.92	2513.9	12.6	38.6	39.7	2499.8	7.7	20.5	21.2	2489.1	9.7	20.3	21.3	2489.1	20.3	101.0
Trial4_123	60	NA()	1.1	10.9071	1.6	3.0924	3.5	0.2447	3.1	0.89	1411.3	10.6	24.6	25.2	1430.8	7.7	17.5	18.1	1460.8	10.5	22.7	23.9	1460.8	22.7	96.6
Trial4_124	121	NA()	1.9	13.2725	1.5	1.9221	3.2	0.1851	2.8	0.89	1094.8	7.8	19.3	19.8	1088.8	6.1	15.0	15.5	1077.7	9.5	23.3	24.6	1077.7	23.3	101.6
Trial4_125	14	NA()	186.8	13.6389	3.8	1.6688	4.9	0.1651	3.2	0.65	985.3	10.1	18.9	19.4	996.7	16.1	20.7	21.0	1022.8	46.2	51.0	51.6	1022.8	51.0	96.3
Trial4_126	127	NA()	1.1	9.9345	1.0	4.0093	2.5	0.2890	2.3	0.91	1636.6	9.3	27.0	27.7	1636.1	6.1	17.7	18.3	1636.3	7.1	20.9	22.1	1636.3	20.9	100.0
Trial4_127	298	NA()	3.0	13.4579	1.0	1.8850	2.0	0.1841	1.8	0.87	1089.2	4.0	18.0	18.5	1075.8	3.3	14.0	14.5	1049.8	5.8	22.2	23.5	1049.8	22.2	103.7
Trial4_128	111	NA()	2.6	18.5329	3.2	0.4066	4.8	0.0547	3.5	0.74	343.1	3.0	6.6	6.7	346.4	4.7	7.7	7.8	369.3	30.3	38.5	39.5	343.1	6.6	NA
Trial4_129	40	NA()	0.5	10.9646	2.0	3.0674	3.9	0.2440	3.4	0.86	1407.7	10.7	24.6	25.2	1424.6	8.5	17.8	18.4	1450.8	13.6	24.4	25.4	1450.8	24.4	97.0
Trial4_130	122	NA()	1.7	13.1372	1.4	1.9581	3.1	0.1866	2.8	0.89	1103.2	7.4	19.2	19.8	1101.2	5.8	14.9	15.4	1098.2	9.2	23.1	24.3	1098.2	23.1	100.5
Trial4_131	53	NA()	1.3	13.7207	2.3	1.7437	4.1	0.1736	3.4	0.83	1031.9	8.4	18.7	19.2	1024.8	7.9	15.4	15.8	1010.7	17.0	27.4	28.5	1010.7	27.4	102.1
Trial4_132	349	NA()	9.6	12.5414	0.8	2.1092	1.7	0.1919	1.5	0.87	1131.9	3.3	18.5	19.0	1151.8	2.8	14.4	14.9	1190.5	4.9	21.5	22.8	1190.5	21.5	95.1
Trial4_133	29	NA()	1.0	13.8959	3.2	1.6106	4.9	0.1624	3.6	0.75	970.1	8.9	18.1	18.6	974.4	10.8	16.7	17.1	984.9	28.5	35.8	36.6	984.9	35.8	98.5
Trial4_134	354	NA()	1.7	8.8452	0.8	4.8304	1.3	0.3100	1.0	0.78	1740.8	4.0	27.0	27.9	1790.2	2.7	17.4	18.1	1849.1	3.4	19.5	20.7	1849.1	19.5	94.1
Trial4_135	41	NA()	1.4	6.2201	1.2	10.4968	3.1	0.4737	2.9	0.92	2500.0	12.3	38.4	39.4	2479.6	7.6	20.4	21.2	2463.7	9.4	20.3	21.3	2463.7	20.3	101.5
Trial4_136	100	NA()	1.8	11.1483	1.3	3.1080	3.0	0.2514	2.7	0.90	1445.8	9.0	24.4	25.1	1434.7	6.3	17.0	17.5	1419.1	8.3	21.9	23.1	1419.1	21.9	101.9
Trial4_137	36	NA()	1.5	13.0753	2.6	1.9052	4.4	0.1808	3.5	0.81	1071.1	9.3	19.6	20.1	1082.9	8.7	16.2	16.6	1107.7	18.0	27.8	28.9	1107.7	27.8	96.7
Trial4_138	438	NA()	1.9	13.4311	0.8	1.8020	1.5	0.1756	1.3	0.84	1043.0	1.5	16.9	17.5	1046.2	1.8	13.5	14.0	1053.8	4.7	21.9	23.2	1053.8	21.9	99.0
Trial4_139	239	NA()	3.1	13.7458	1.1	1.7738	2.4	0.1769	2.1	0.88	1050.1	4.9	17.7	18.2	1035.9	4.0	13.9	14.4	1007.0	6.8	22.6	23.8	1007.0	22.6	104.3
Trial4_140	61	NA()	0.3	8.8295	1.3	5.3111	3.1	0.3403	2.9	0.91	1887.9	11.5	30.9	31.7	1870.7	7.4	19.0	19.7	1852.3	9.2	21.3	22.4	1852.3	21.3	101.9
Trial4_141	131	NA()	1.0	6.5040	0.9	9.9896	1.9	0.4714	1.6	0.88	2489.9	9.7	37.5	38.6	2433.8	5.7	19.8	20.5	2388.1	6.9	19.3	20.4	2388.1	19.3	104.3
Trial4_142	44	NA()	1.2	13.1145	2.4	1.9149	4.2	0.1822	3.5	0.83	1079.1	9.1	19.7	20.1	1086.3	8.3	16.0	16.5	1101.7	16.9	27.1	28.2	1101.7	27.1	97.9
Trial4_143	70	NA()	2.4	12.5701	1.8	2.1628	3.7	0.1973	3.2	0.87	1160.6	8.6	20.5	21.0	1169.2	7.0	15.8	16.3	1186.0	11.8	24.0	25.2	1186.0	24.0	97.9
Trial4_144	22	NA()	0.4	5.5135	1.4	13.0337	3.6	0.5214	3.2	0.91	2705.2	13.1	40.9	42.0	2682.1	8.2	21.0	21.7	2665.4	10.5	20.5	21.4	2665.4	20.5	101.5
Trial4_145	67	NA()	1.5	8.7750	1.3	5.1874	3.1	0.3303	2.8	0.91	1839.8	10.8	30.1	30.9	1850.6	7.0	18.8	19.4	1863.5	8.5	21.0	22.1	1863.5	21.0	98.7
Trial4_146	181	NA()	1.0	8.3752	0.9	5.5283	1.8	0.3360	1.6	0.89	1867.2	7.6	29.4	30.3	1905.0	4.8	18.3	18.9	1947.3	5.3	19.7	20.8	1947.3	19.7	95.9
Trial4_147	449	NA()	2.8	13.2428	0.8	1.8578	1.5	0.1785	1.2	0.82	1058.8	1.1	17.1	17.7	1066.2	1.6	13.6	14.2	1082.2	4.5	21.7	23.1	1082.2	21.7	97.8
Trial4_148	40	NA()	2.1	10.0308	1.8	4.0060	3.7	0.2916	3.3	0.88	1649.4	11.1	27.8	28.5	1635.4	8.1	18.5	19.1	1618.4	12.0	23.1	24.2	1618.4	23.1	101.9
Trial4_149	95	NA()	1.5	11.2396	1.3	3.1194	3.0	0.2544	2.7	0.90	1461.1	9.0	24.6	25.3	1437.5	6.2	16.9	17.5	1403.5	8.4	22.0	23.2	1403.5	22.0	104.1
Trial4_150	234	NA()	2.9	9.6141	0.8	4.3279	1.7	0.3019	1.4	0.86	1700.8	6.1	26.9	27.7	1698.7	4.0	17.4	18.0	1696.9	4.8	20.1	21.3	1696.9	20.1	100.2
Trial4_151	27	NA()	1.1	6.8624	1.6	8.3213	3.6	0.4143	3.3	0.90	2234.8	12.8	35.5	36.5	2266.6	8.3	20.3	21.0	2296.3	10.5	21.0	22.0	2296.3	21.0	97.3
Trial4_152	86	NA()	2.2	10.5650	1.4	3.1916	3.2	0.2447	2.9	0.90	1410.9	9.1	24.0	24.7	1455.1	6.6	17.2	17.8	1521.1	8.8	21.9	23.0	1521.1	21.9	92.8
Trial4_153	17	NA()	1.1	6.7332	1.9	9.0243	4.0	0.4409	3.5	0.87	2354.6	14.0	37.3	38.3	2340.5	9.6	21.0	21.7	2328.9	13.3	22.5	23.4	2328.9	22.5	101.1
Trial4_154	399	NA()	2.6	11.7400	0.8	2.5132	1.4	0.2141	1.1	0.81	1250.6	2.0	20.0	20.6	1275.9	2.0	15.0	15.6	1319.6	4.0	20.9	22.2	1319.6	20.9	94.8
Trial4_155	136	NA()	1.1	9.8908	1.0	4.0736	2.4	0.2924	2.2	0.91	1653.3	8.6	27.0	27.7	1649.0	5.6	17.6	18.2	1644.5	6.6	20.8	21.9	1644.5	20.8	100.5
Trial4_156	79	NA()	1.2	11.1228	1.4	3.2148	3.2	0.2595	2.9	0.90	1487.1	9.6	25.2	25.8	1460.7	6.8	17.2	17.8	1423.5	9.4	22.3	23.5	1423.5	22.3	104.5
Trial4_157	157	NA()	2.1	11.6863	1.1	2.5876	2.6	0.2194	2.4	0.90	1278.8	7.1	21.5	22.1	1297.1	5.2	15.9	16.5	1328.5	7.2	21.8	23.0	1328.5	21.8	96.3
Trial4_158	212	NA()	1.6	14.1252	1.3	1.5377	2.7	0.1576	2.4	0.89	943.4	5.2	16.2	16.7	945.6	4.3	13.3	13.8	951.5	7.6	23.0	24.3	951.5	23.0	99.2
Trial4_159	247	NA()	1.3	9.3101	0.8	4.3212	1.6	0.2919	1.4	0.86	1651.1	6.4	26.3	27.1	1697.4	4.2	17.4	18.1	1755.9	4.7	20.0	21.2	1755.9	20.0	94.0
Trial4_160	78	NA()	1.4	13.4077	1.9	1.8753	3.7	0.1824	3.2	0.86	1080.3	8.4	19.3	19.8	1072.4	7.0	15.3	15.7	1057.3	12.8	24.9	26.1	1057.3	24.9	102.2
Trial4_161	69	NA()	1.5	5.4167	0.9	13.4542	2.5	0.5288	2.3	0.92	2736.4	10.9	40.6	41.7	2712.1	6.6	20.5	21.2	2694.8	8.2	19.4	20.4	2694.8	19.4	101.5
Trial4_162	164	NA()	1.9	18.1856	2.3	0.5357	4.0	0.0707	3.3	0.81	440.3	3.2	8.1	8.3	435.6	4.0	8.3	8.5	411.7	18.9	30.3	31.5	440.3	8.1	106.9
Trial4_163	203	NA()	2.2	12.5784	1.1	2.2140	2.4	0.2021	2.2	0.89	1186.4	6.1	19.9	20.5	1185.5	4.6	15.0	15.6	1184.7	6.6	22.0	23.2	1184.7	22.0	100.1
Trial4_164	112	NA()	2.7	13.6372	1.6	1.7471	3.4	0.1729	3.0	0.88	1027.9	7.5	18.2	18.7	1026.1	6.1	14.6	15.0	1023.0	10.5	23.9	25.2	1023.0	23.9	100.5
Trial4_165	23	NA()	0.3	6.2335	1.6	9.9712	3.7	0.4510	3.3	0.90	2399.7	13.7	37.7	38.7	2432.1	8.8	20.9	21.5	2460.1	11.2	21.1	22.1	2460.1	21.1	97.5
Trial4_166	499	NA()	0.5	12.5854	0.8	2.0212	1.3	0.1846	1.0	0.77	1091.9	0.8	17.6	18.2	1122.6	1.5	14.0	14.5	1183.6	4.0	21.3	22.6	1183.6	21.3	92.3
Trial4_167	198	NA()	2.1	9.0741	0.9	4.9848	1.8	0.3282	1.6	0.88	1829.7	7.7	29.0	29.8	1816.7	4.8	18.0	18.6	1802.7	5.4	20.0	21.2	1802.7	20.0	101.5
Trial4_168	58	NA()	2.0	12.0788	2.0	2.1415	3.9	0.1877	3.3	0.86	1108.9	8.8	19.9	20.4	1162.3	7.9	16.2	16.7	1264.2	14.4	25.2	26.3	1264.2	25.2	87.7
Trial4_169	101	NA()	0.7	17.9697	2.9	0.5539	4.5	0.0722	3.5	0.77	449.5	4.0	8.6	8.8	447.5	5.4	9.1	9.4	438.4	25.9	35.1	36.1	449.5	8.6	102.5
Trial4_170	124	NA()	0.9	10.7835	1.2	3.3646	2.7	0.2633	2.5	0.90	1506.5	9.0	25.2	25.9	1496.2	6.1	17.2	17.8	1482.4	7.7	21.5	22.7	1482.4	21.5	101.6
Trial4_171	38	NA()	1.5	12.2410	2.3	2.3549	4.2	0.2092	3.5	0.83	1224.3	10.1	22.0	22.5	1229.0	8.7	17.0	17.5	1238.2	16.2	26.4	27.4	1238.2	26.4	98.9
Trial4_172	53	NA()	1.4	3.0411	1.0	33.4705	2.4	0.7386	2.2	0.92	3565.3	8.4	48.7	50.2	3594.6	6.0	21.1	21.9	3611.6	8.1	18.2	19.1	3611.6	18.2	98.7
Trial4_173	55	NA()	2.7	9.4138	1.5	4.6690	3.4	0.3189	3.0	0.90	1784.5	11.6	29.7	30.5	1761.7	7.7	18.8	19.4	1735.6	9.9	21.8	22.9	1735.6	21.8	102.8
Trial4_174	374	NA()	1.9	13.5667	0.9	1.7909	1.7	0.1763	1.5	0.85	1046.7	2.6	17.1	17.7	1042.2	2.4	13.6	14.1	1033.5	5.1	22.1	23.4	1033.5	22.1	101.3
Trial4_175	58	NA()	2.1	12.7057	2.1	1.9445	3.9	0.1793	3.3	0.85	1063.0	8.7	19.2	19.7	1096.6	7.6	15.7	16.2	1164.7	14.2	25.4	26.5	1164.7	25.4	91.3
Trial4_176	125	NA()	1.8	12.7399	1.4	2.2351	3.0	0.2066	2.7	0.89	1210.7	7.9	20.9	21.5	1192.1	5.9	15.5	16.1	1159.4	8.7	22.8	24.0	1159.4	22.8	104.4
Trial4_177	74	NA()	1.5	12.8651	1.8	2.0723	3.6	0.1934	3.2	0.87	1140.0	8.7	20.3	20.8	1139.7	7.1	15.7	16.2	1140.0	12.1	24.3	25.5	1140.0	24.3	100.0
Trial4_178	31	NA()	1.1	13.0383	2.8	1.9081	4.6	0.1805	3.6	0.78	1069.8	9.5	19.7	20.2	1083.9	9.5	16.6	17.1	1113.3	21.0	29.9	30.8	1113.3	29.9	96.1
Trial4_179	84	NA()	1.2	9.8430	1.3	3.7558	3.1	0.2682	2.8	0.91	1531.9	10.0	25.9	26.6	1583.4	6.9	17.8	18.4	1653.4	8.5	21.4	22.5	1653.4	21.4	92.6
Trial4_180	61	NA()	1.2	10.7934	1.6	3.2153	3.5	0.2518	3.1	0.88	1447.8	10.3	25.0	25.6	1460.9	7.7	17.7	18.2	1480.7	11.4	23.1	24.2	1480.7	23.1	97.8
Trial4_181	209	NA()	1.3	8.6728	0.8	5.0220	1.8	0.3160	1.5	0.88	1770.3	7.4	28.1	28.9	1823.0	4.7	18.0	18.7	1884.6	5.2	19.8	21.0	1884.6	19.8	93.9
Trial4_182	27	NA()	1.4	6.0153	1.4	11.3300	3.5	0.4945	3.2	0.91	2590.2	13.1	39.6	40.7	2550.7	8.2	20.8	21.5	2520.2	10.6	20.7	21.7	2520.2	20.7	102.8
Trial4_183	52	NA()	0.5	20.8053	5.5	0.1890	6.3	0.0285	3.2	0.51	181.4	2.1	3.8	3.8	175.8	7.6	8.3	8.3	102.5	108.0	110.9	111.2	181.4	3.8	NA
Trial4_184	87	NA()	2.1	12.6565	1.6	2.1991	3.4	0.2020	3.0	0.88	1185.8	8.6	20.8	21.4	1180.8	6.6	15.8	16.3	1172.4	10.3	23.3	24.6	1172.4	23.3	101.1
Trial4_185	57	NA()	2.0	12.4072	2.0	2.1583	3.9	0.1943	3.3	0.86	1144.7	9.1	20.5	21.0	1167.7	7.6	16.1	16.6	1211.6	13.2	24.7	25.8	1211.6	24.7	94.5
Trial4_186	206	NA()	2.3	13.5931	1.2	1.6927	2.7	0.1670	2.4	0.89	995.3	5.8	17.2	17.6	1005.8	4.7	13.9	14.4	1029.6	7.6	22.8	24.1	1029.6	22.8	96.7
Trial4_187	185	NA()	1.6	13.9794	1.3	1.6634	2.8	0.1687	2.5	0.89	1005.1	6.2	17.5	17.9	994.7	5.0	13.9	14.4	972.7	8.5	23.2	24.5	972.7	23.2	103.3
Trial4_188	163	NA()	2.0	10.9047	1.0	3.0191	2.5	0.2389	2.2	0.90	1380.9	7.7	23.1	23.7	1412.4	5.4	16.5	17.1	1461.2	6.5	21.2	22.4	1461.2	21.2	94.5
Trial4_189	223	NA()	3.2	13.5258	1.1	1.8717	2.4	0.1837	2.1	0.89	1087.1	5.5	18.4	18.9	1071.1	4.3	14.2	14.7	1039.6	6.9	22.5	23.8	1039.6	22.5	104.6
Trial4_190	193	NA()	1.8	17.9060	2.2	0.4694	3.9	0.0610	3.2	0.82	381.6	2.4	6.9	7.1	390.7	3.2	7.4	7.6	446.3	16.6	28.8	30.0	381.6	6.9	NA
Trial4_191	150	NA()	2.2	13.2889	1.3	1.9770	2.9	0.1906	2.6	0.89	1124.8	6.4	19.2	19.7	1107.7	5.1	14.7	15.2	1075.2	8.5	22.9	24.2	1075.2	22.9	104.6
Trial4_192	110	NA()	1.2	12.6949	1.5	2.1657	3.2	0.1995	2.8	0.89	1172.6	7.4	20.2	20.7	1170.1	5.8	15.4	15.9	1166.4	9.4	23.0	24.2	1166.4	23.0	100.5
Trial4_193	151	NA()	1.3	17.9487	2.4	0.5327	4.1	0.0694	3.3	0.80	432.4	3.0	7.9	8.1	433.6	4.2	8.4	8.6	441.0	20.9	31.5	32.6	432.4	7.9	98.1
Trial4_194	60	NA()	1.6	4.8296	0.9	16.0384	2.5	0.5620	2.3	0.93	2875.0	9.7	41.8	43.0	2879.1	6.2	20.5	21.3	2882.6	8.0	19.0	20.0	2882.6	19.0	99.7
Trial4_195	57	NA()	0.9	5.5470	0.9	12.4602	2.7	0.5015	2.5	0.94	2620.3	9.6	38.9	40.1	2639.7	6.0	20.2	20.9	2655.4	7.5	19.1	20.2	2655.4	19.1	98.7
Trial4_196	130	NA()	1.9	12.6145	1.4	2.0869	3.0	0.1910	2.7	0.89	1126.8	5.9	19.1	19.6	1144.5	4.9	14.9	15.4	1179.0	8.4	22.6	23.8	1179.0	22.6	95.6
Trial4_197	34	NA()	1.7	13.6077	3.0	1.5842	4.6	0.1564	3.6	0.77	936.8	7.3	16.9	17.4	964.0	9.1	15.6	16.1	1027.4	24.2	32.3	33.2	1027.4	32.3	91.2
Trial4_198	193	NA()	1.2	11.1039	1.0	3.1071	2.1	0.2503	1.9	0.89	1440.2	5.4	23.3	23.9	1434.4	3.9	16.2	16.8	1426.7	5.7	21.0	22.3	1426.7	21.0	100.9
Trial4_199	62	NA()	1.6	13.1831	2.0	1.8946	3.8	0.1812	3.3	0.85	1073.7	7.8	19.0	19.5	1079.2	6.9	15.3	15.8	1091.2	13.6	25.3	26.4	1091.2	25.3	98.4
Trial4_200	66	NA()	1.0	10.8236	1.6	2.9513	3.5	0.2318	3.1	0.89	1343.9	8.8	23.0	23.6	1395.2	6.9	17.0	17.6	1475.4	10.4	22.7	23.8	1475.4	22.7	91.1
Trial4_201	165	NA()	2.2	5.2945	0.9	12.9734	1.5	0.4984	1.2	0.81	2606.9	7.3	38.3	39.4	2677.7	4.5	19.8	20.6	2732.4	5.4	18.3	19.3	2732.4	18.3	95.4
Trial4_202	196	NA()	1.0	8.9384	0.8	5.3019	1.7	0.3439	1.4	0.86	1905.2	6.7	29.7	30.6	1869.2	4.2	18.0	18.7	1830.1	5.1	19.9	21.1	1830.1	19.9	104.1
Trial4_203	268	NA()	1.9	11.1116	0.8	3.1513	1.6	0.2541	1.4	0.86	1459.5	5.0	23.4	24.1	1445.3	3.5	16.2	16.8	1425.4	4.8	20.8	22.1	1425.4	20.8	102.4
Trial4_204	19	NA()	2.2	12.6248	3.4	2.0602	5.0	0.1887	3.7	0.73	1114.4	10.3	20.7	21.2	1135.7	12.2	18.6	19.0	1177.4	29.3	36.0	36.8	1177.4	36.0	94.7
Trial4_205	211	NA()	3.7	7.8303	0.9	6.8738	1.5	0.3905	1.2	0.82	2125.3	7.7	32.7	33.6	2095.3	4.7	18.7	19.5	2066.7	5.4	19.5	20.6	2066.7	19.5	102.8
Trial4_206	76	NA()	0.8	18.9471	3.8	0.2666	4.9	0.0366	3.2	0.65	232.0	2.4	4.7	4.8	239.9	5.4	6.9	7.0	319.3	51.6	57.0	57.6	232.0	4.7	NA
Trial4_207	54	NA()	2.6	9.1759	1.4	4.6824	3.4	0.3117	3.0	0.90	1749.3	11.4	29.2	30.0	1764.1	7.7	18.8	19.4	1782.4	9.9	21.7	22.8	1782.4	21.7	98.1
Trial4_208	109	NA()	2.2	13.5494	1.6	1.6639	3.4	0.1636	3.0	0.88	976.7	7.2	17.4	17.9	994.9	6.0	14.3	14.8	1036.1	10.2	23.8	25.0	1036.1	23.8	94.3
Trial4_209	188	NA()	3.2	11.3582	1.0	2.8044	2.3	0.2311	2.1	0.90	1340.4	6.8	22.3	22.9	1356.7	4.8	16.1	16.7	1383.4	6.2	21.3	22.5	1383.4	21.3	96.9
Trial4_210	207	NA()	1.1	13.4624	1.2	1.8312	2.5	0.1789	2.3	0.89	1060.8	5.5	18.0	18.5	1056.7	4.4	14.2	14.7	1049.1	7.3	22.6	23.9	1049.1	22.6	101.1
Trial4_211	65	NA()	2.3	13.7023	2.0	1.7243	3.9	0.1714	3.3	0.85	1020.0	8.1	18.4	18.9	1017.6	7.0	14.9	15.4	1013.4	13.8	25.5	26.7	1013.4	25.5	100.7
Trial4_212	98	NA()	2.1	19.0370	3.8	0.2726	5.3	0.0377	3.7	0.69	238.3	2.2	4.7	4.8	244.8	5.5	7.1	7.2	308.5	53.1	58.4	59.0	238.3	4.7	NA
Trial4_213	113	NA()	2.0	9.3276	1.0	4.5556	2.6	0.3083	2.3	0.91	1732.5	9.9	28.4	29.2	1741.2	6.3	18.2	18.8	1752.5	7.3	20.7	21.9	1752.5	20.7	98.9
Trial4_214	29	NA()	0.8	13.2527	3.0	1.7686	4.7	0.1701	3.6	0.77	1012.5	9.3	18.9	19.3	1034.0	10.1	16.7	17.1	1080.7	24.1	32.2	33.1	1080.7	32.2	93.7
Trial4_215	81	NA()	1.4	11.0232	1.4	3.1892	3.2	0.2551	2.9	0.90	1464.6	10.0	25.0	25.7	1454.5	6.9	17.3	17.9	1440.7	9.1	22.2	23.3	1440.7	22.2	101.7
Trial4_216	115	NA()	2.2	17.5198	2.5	0.6726	4.2	0.0855	3.4	0.80	528.9	4.4	9.9	10.2	522.3	5.7	10.1	10.4	494.5	24.1	33.6	34.7	528.9	9.9	106.9
Trial4_217	42	NA()	1.5	5.5854	1.1	12.1103	3.1	0.4908	2.8	0.93	2574.1	12.3	39.2	40.3	2613.0	7.6	20.6	21.4	2644.0	9.3	19.9	20.9	2644.0	19.9	97.4
Trial4_218	391	NA()	3.8	13.3183	0.9	1.7845	1.7	0.1724	1.5	0.86	1025.6	2.7	16.8	17.3	1039.8	2.5	13.6	14.1	1070.8	5.0	21.9	23.2	1070.8	21.9	95.8
Trial4_219	103	NA()	0.8	13.3558	1.6	1.9435	3.3	0.1883	2.9	0.88	1112.4	8.5	19.8	20.3	1096.2	6.6	15.2	15.7	1065.1	10.7	23.9	25.1	1065.1	23.9	104.4
Trial4_220	196	NA()	2.2	19.6750	3.0	0.2607	4.6	0.0372	3.5	0.75	235.6	1.9	4.5	4.6	235.2	3.1	5.3	5.4	232.9	28.1	37.2	38.3	235.6	4.5	NA
Trial4_221	53	NA()	0.7	16.2271	3.2	0.8298	4.9	0.0977	3.6	0.75	600.9	6.1	11.8	12.0	613.5	7.6	12.1	12.4	661.1	26.9	35.2	36.2	600.9	11.8	90.9
Trial4_222	33	NA()	0.9	6.1195	1.3	10.2956	3.3	0.4572	3.1	0.92	2427.0	13.4	37.9	38.9	2461.7	8.3	20.7	21.4	2491.3	10.2	20.6	21.6	2491.3	20.6	97.4
Trial4_223	30	NA()	0.8	10.8304	2.2	3.2205	4.1	0.2531	3.5	0.84	1454.4	12.0	25.8	26.4	1462.1	9.5	18.5	19.0	1474.2	15.3	25.3	26.3	1474.2	25.3	98.7
Trial4_224	50	NA()	1.3	17.3445	3.7	0.6548	5.3	0.0824	3.7	0.71	510.5	5.7	10.3	10.6	511.4	7.6	11.2	11.5	516.7	32.7	40.2	41.0	510.5	10.3	98.8
Trial4_225	113	NA()	4.9	17.1037	2.3	0.7862	4.0	0.0976	3.3	0.82	600.2	5.7	11.5	11.8	589.0	5.6	10.8	11.1	547.3	17.1	28.8	30.0	600.2	11.5	109.7
Trial4_226	158	NA()	3.0	13.2821	1.3	1.8971	2.9	0.1828	2.6	0.89	1082.4	8.0	19.2	19.7	1080.1	6.0	14.9	15.4	1076.3	8.4	22.9	24.1	1076.3	22.9	100.6
Trial4_227	129	NA()	1.4	16.8354	2.2	0.7923	3.9	0.0968	3.3	0.83	595.6	5.4	11.3	11.6	592.5	5.4	10.7	11.0	581.8	16.1	28.1	29.3	595.6	11.3	102.4
Trial4_228	224	NA()	3.5	13.5764	1.1	1.8199	2.5	0.1793	2.2	0.89	1063.0	6.6	18.4	18.9	1052.6	5.1	14.3	14.8	1032.1	7.5	22.7	24.0	1032.1	22.7	103.0
Trial4_229	33	NA()	1.3	7.7216	1.6	6.4215	3.6	0.3598	3.3	0.90	1981.1	14.1	33.0	33.9	2035.2	9.0	20.1	20.8	2091.3	10.8	21.6	22.6	2091.3	21.6	94.7
Trial4_230	125	NA()	1.7	5.6269	0.9	12.7773	1.8	0.5217	1.5	0.85	2706.3	10.9	40.2	41.4	2663.4	6.6	20.4	21.1	2631.7	8.4	19.5	20.5	2631.7	19.5	102.8
Trial4_231	29	NA()	2.9	10.1039	2.1	3.7571	4.1	0.2754	3.5	0.85	1568.4	13.3	27.8	28.5	1583.6	9.9	19.2	19.7	1604.8	14.5	24.5	25.5	1604.8	24.5	97.7
Trial4_232	138	NA()	1.1	19.9109	3.6	0.2536	5.1	0.0366	3.6	0.71	232.0	2.3	4.6	4.7	229.5	3.7	5.6	5.7	205.4	34.1	42.0	43.0	232.0	4.6	NA
Trial4_233	285	NA()	2.1	13.0549	1.0	2.0524	2.0	0.1944	1.7	0.88	1145.2	5.8	19.3	19.8	1133.1	4.3	14.6	15.2	1110.8	5.9	22.0	23.3	1110.8	22.0	103.1
Trial4_234	232	NA()	0.8	11.5774	0.9	2.8764	2.0	0.2416	1.8	0.89	1395.2	7.4	23.2	23.9	1375.7	5.0	16.2	16.8	1346.6	6.0	21.3	22.6	1346.6	21.3	103.6
Trial4_235	151	NA()	2.2	12.0255	1.2	2.5677	2.7	0.2240	2.4	0.90	1303.2	8.9	22.5	23.1	1291.5	6.2	16.2	16.8	1272.9	7.7	22.1	23.3	1272.9	22.1	102.4

Trial4_236	292	NA()	3.3	13.3122	1.0	1.7390	2.2	0.1680	1.9	0.88	1000.9	5.0	17.0	17.5	1023.1	4.0	13.8	14.3	1071.7	6.1	22.2	23.5	1071.7	22.2	93.4
Trial4_237	266	NA()	0.9	18.0299	1.8	0.5283	3.4	0.0691	2.9	0.85	430.8	2.8	7.8	8.0	430.7	3.1	7.8	8.1	430.9	12.9	27.0	28.3	430.8	7.8	100.0
Trial4_238	168	NA()	1.3	13.7398	1.3	1.7234	2.9	0.1718	2.6	0.89	1022.1	7.3	18.1	18.6	1017.3	5.6	14.3	14.8	1007.8	8.3	23.1	24.3	1007.8	23.1	101.4
Trial4_239	35	NA()	1.3	12.9640	2.6	1.9866	4.4	0.1869	3.6	0.81	1104.4	11.0	20.9	21.4	1111.0	9.8	17.0	17.4	1124.7	19.3	28.7	29.6	1124.7	28.7	98.2
Trial4_240	123	NA()	1.5	9.7423	1.0	4.1982	2.5	0.2968	2.3	0.91	1675.3	11.3	28.2	29.0	1673.7	7.2	18.3	18.9	1672.5	7.8	21.1	22.3	1672.5	21.1	100.2
Trial4_241	60	NA()	1.8	13.6811	2.1	1.7722	4.0	0.1759	3.4	0.85	1044.7	10.0	19.6	20.1	1035.3	8.3	15.7	16.1	1016.5	15.4	26.4	27.5	1016.5	26.4	102.8
Trial4_242	137	NA()	1.5	13.5618	1.4	1.8498	3.1	0.1820	2.7	0.89	1078.0	8.7	19.4	19.9	1063.4	6.6	15.0	15.5	1034.2	9.6	23.5	24.8	1034.2	23.5	104.2
Trial4_243	112	NA()	2.0	9.1840	1.0	4.7584	2.6	0.3171	2.4	0.91	1775.5	11.8	29.7	30.4	1777.6	7.4	18.7	19.3	1780.8	8.0	20.9	22.1	1780.8	20.9	99.7
Trial4_244	46	NA()	1.5	12.2014	2.1	2.4617	4.0	0.2179	3.4	0.85	1271.0	11.7	23.3	23.9	1260.9	9.1	17.4	17.9	1244.5	14.6	25.4	26.5	1244.5	25.4	102.1
Trial4_245	60	NA()	3.4	9.6738	1.4	4.3930	3.3	0.3084	3.0	0.90	1732.6	13.1	29.7	30.4	1711.0	8.4	18.9	19.5	1685.5	10.1	22.0	23.1	1685.5	22.0	102.8
Trial4_246	225	NA()	1.3	5.3437	0.9	13.0080	1.4	0.5044	1.1	0.77	2632.5	8.6	38.8	40.0	2680.2	5.1	20.0	20.7	2717.1	6.1	18.5	19.6	2717.1	18.5	96.9
Trial4_247	91	NA()	1.0	16.1460	2.4	0.8745	4.2	0.1025	3.4	0.82	628.8	6.2	12.2	12.5	638.0	6.3	11.6	11.9	671.9	18.1	29.0	30.2	628.8	12.2	93.6
Trial4_248	115	NA()	4.3	9.3898	1.1	4.3947	2.6	0.2994	2.4	0.91	1688.4	11.2	28.3	29.1	1711.3	7.1	18.4	19.0	1740.3	7.7	20.9	22.0	1740.3	20.9	97.0
Trial4_249	80	NA()	1.2	16.8112	2.7	0.8035	4.4	0.0980	3.5	0.80	602.7	6.1	11.8	12.1	598.8	6.3	11.2	11.5	584.9	19.5	30.2	31.3	602.7	11.8	103.1
Trial4_250	113	NA()	1.8	13.4511	1.6	1.8217	3.3	0.1778	2.9	0.88	1055.0	8.7	19.1	19.6	1053.3	6.8	15.1	15.5	1050.8	10.9	24.0	25.2	1050.8	24.0	100.4
Trial4_251	59	NA()	1.1	13.4149	2.1	1.8631	3.9	0.1814	3.3	0.85	1074.4	9.2	19.7	20.1	1068.1	7.8	15.6	16.1	1056.3	14.7	25.9	27.0	1056.3	25.9	101.7
Trial4_252	38	NA()	0.8	8.7352	1.6	5.1015	3.6	0.3233	3.2	0.89	1806.0	12.8	30.4	31.2	1836.4	8.7	19.4	20.1	1871.7	11.2	22.2	23.2	1871.7	22.2	96.5
Trial4_253	29	NA()	2.0	13.4929	3.0	1.8766	4.7	0.1837	3.6	0.77	1087.3	10.2	20.3	20.8	1072.8	10.6	17.2	17.6	1044.5	24.8	32.7	33.6	1044.5	32.7	104.1
Trial4_254	87	NA()	2.5	13.6664	1.8	1.8079	3.6	0.1793	3.1	0.87	1063.0	8.6	19.2	19.7	1048.3	6.9	15.1	15.5	1018.7	12.0	24.6	25.8	1018.7	24.6	104.4
Trial4_255	48	NA()	1.8	7.8302	1.3	6.7467	3.2	0.3833	3.0	0.91	2091.8	12.6	33.7	34.6	2078.8	7.9	19.8	20.4	2066.7	9.6	21.0	22.1	2066.7	21.0	101.2
Trial4_256	170	NA()	5.4	12.7367	1.2	2.1704	2.7	0.2006	2.4	0.90	1178.5	7.2	20.2	20.8	1171.6	5.3	15.2	15.7	1159.9	7.4	22.3	23.6	1159.9	22.3	101.6
Trial4_257	24	NA()	1.6	13.4601	3.3	1.8459	4.9	0.1803	3.7	0.74	1068.5	10.1	20.0	20.4	1062.0	11.4	17.7	18.1	1049.5	28.2	35.4	36.2	1049.5	35.4	101.8
Trial4_258	68	NA()	0.6	5.3394	0.9	12.9581	2.5	0.5020	2.4	0.93	2622.5	11.6	39.5	40.6	2676.6	7.0	20.5	21.3	2718.5	8.3	19.4	20.4	2718.5	19.4	96.5
Trial4_259	150	NA()	1.7	13.2879	1.3	1.9012	2.9	0.1833	2.6	0.89	1085.0	6.7	18.7	19.3	1081.5	5.3	14.6	15.1	1075.4	8.5	22.9	24.2	1075.4	22.9	100.9
Trial4_260	34	NA()	1.1	10.2874	2.1	3.3173	4.0	0.2476	3.4	0.85	1426.2	10.9	24.9	25.6	1485.1	8.7	18.2	18.7	1571.2	13.3	23.9	25.0	1571.2	23.9	90.8
Trial4_261	116	NA()	1.5	13.5301	1.6	1.7176	3.3	0.1686	2.9	0.88	1004.5	7.0	17.7	18.2	1015.1	5.7	14.3	14.8	1039.0	9.6	23.5	24.7	1039.0	23.5	96.7
Trial4_262	40	NA()	0.6	5.6503	1.1	12.5957	3.0	0.5164	2.8	0.93	2683.9	11.6	40.2	41.3	2649.9	7.1	20.6	21.3	2624.8	9.1	19.8	20.8	2624.8	19.8	102.3
Trial4_263	109	NA()	1.3	18.5176	2.8	0.5397	4.4	0.0725	3.4	0.77	451.3	3.6	8.5	8.7	438.2	4.9	8.8	9.1	371.1	25.0	34.6	35.6	451.3	8.5	121.6
Trial4_264	245	NA()	3.1	12.8125	1.0	2.0900	2.2	0.1943	2.0	0.89	1144.6	4.8	19.0	19.5	1145.5	3.8	14.6	15.1	1148.1	6.0	21.9	23.2	1148.1	21.9	99.7
Trial4_265	98	NA()	1.7	19.2279	3.8	0.3280	5.2	0.0458	3.6	0.69	288.4	2.6	5.6	5.7	288.0	4.5	6.8	7.0	285.7	35.1	42.7	43.6	288.4	5.6	NA
Trial4_266	104	NA()	2.1	10.2688	1.2	3.8575	2.8	0.2874	2.5	0.91	1628.6	9.2	26.8	27.6	1604.8	6.1	17.6	18.2	1574.6	7.6	21.3	22.5	1574.6	21.3	103.4
Trial4_267	55	NA()	2.1	5.4735	0.9	12.9349	2.7	0.5137	2.5	0.94	2672.5	10.3	39.7	40.8	2674.9	6.3	20.3	21.0	2677.5	7.9	19.2	20.3	2677.5	19.2	99.8
Trial4_268	170	NA()	1.4	13.5909	1.3	1.7793	2.8	0.1755	2.5	0.89	1042.1	5.3	17.7	18.2	1037.9	4.5	14.0	14.6	1029.9	8.2	23.0	24.2	1029.9	23.0	101.2
Trial4_269	89	NA()	2.0	13.0505	1.7	1.9852	3.5	0.1880	3.0	0.88	1110.5	7.2	19.3	19.8	1110.5	6.0	15.1	15.6	1111.4	10.7	23.7	24.9	1111.4	23.7	99.9
Trial4_270	75	NA()	0.9	5.6138	0.9	12.6133	2.4	0.5138	2.2	0.92	2672.8	9.6	39.5	40.7	2651.2	5.9	20.2	20.9	2635.5	7.4	19.1	20.1	2635.5	19.1	101.4
Trial4_271	127	NA()	2.2	12.7650	1.4	2.1380	3.0	0.1980	2.7	0.89	1164.7	6.2	19.7	20.2	1161.2	5.0	15.0	15.5	1155.5	8.3	22.6	23.9	1155.5	22.6	100.8
Trial4_272	61	NA()	1.4	11.2443	1.6	3.0402	3.5	0.2480	3.1	0.88	1428.4	8.6	24.0	24.7	1417.8	6.5	17.0	17.5	1402.7	10.2	22.8	23.9	1402.7	22.8	101.8
Trial4_273	132	NA()	1.9	9.8408	1.0	3.9246	2.5	0.2802	2.3	0.91	1592.5	7.2	25.8	26.5	1618.8	5.0	17.3	17.9	1653.9	6.4	20.7	21.8	1653.9	20.7	96.3
Trial4_274	140	NA()	3.8	9.6315	0.9	4.4999	2.2	0.3145	2.0	0.91	1762.7	7.3	28.0	28.8	1731.0	4.8	17.7	18.3	1693.6	6.0	20.5	21.6	1693.6	20.5	104.1
Trial4_275	109	NA()	1.5	10.5990	1.1	3.4611	2.9	0.2662	2.7	0.93	1521.4	10.3	27.2	27.9	1518.4	6.9	17.8	18.3	1515.1	8.0	19.2	20.5	1515.1	19.2	100.4
Trial4_276	107	NA()	1.0	9.6944	1.0	4.1619	2.8	0.2928	2.6	0.93	1655.3	10.7	29.2	29.9	1666.5	7.0	18.4	19.0	1681.6	7.8	18.8	20.1	1681.6	18.8	98.4
Trial4_277	127	NA()	1.4	5.1191	0.9	14.2408	1.8	0.5290	1.6	0.86	2737.0	10.3	42.7	43.9	2765.9	6.2	20.7	21.4	2787.7	7.7	17.0	18.1	2787.7	17.0	98.2
Trial4_278	430	NA()	5.5	12.2310	0.8	2.3701	1.3	0.2103	1.0	0.78	1230.7	2.5	21.0	21.6	1233.6	2.1	15.0	15.6	1239.8	3.9	18.6	20.0	1239.8	18.6	99.3
Trial4_279	132	NA()	0.4	13.4443	1.3	1.7371	3.2	0.1695	3.0	0.92	1009.1	7.7	19.0	19.4	1022.4	6.1	14.7	15.2	1051.8	9.3	20.9	22.2	1051.8	20.9	95.9
Trial4_280	153	NA()	2.6	12.7987	1.1	2.0673	2.9	0.1920	2.7	0.92	1132.1	7.9	20.8	21.3	1138.0	5.8	15.4	15.9	1150.2	7.9	20.0	21.4	1150.2	20.0	98.4
Trial4_281	51	NA()	2.6	11.4707	1.6	2.9345	3.8	0.2442	3.4	0.91	1408.7	11.4	26.2	26.8	1390.9	8.3	17.8	18.3	1364.4	11.9	21.4	22.7	1364.4	21.4	103.2
Trial4_282	93	NA()	0.8	5.1349	0.9	13.9324	2.2	0.5191	2.0	0.91	2695.4	11.4	42.5	43.6	2745.1	6.8	20.8	21.6	2782.6	8.2	17.3	18.4	2782.6	17.3	96.9
Trial4_283	132	NA()	0.6	5.2123	0.9	14.2048	1.7	0.5372	1.4	0.84	2771.8	10.3	43.1	44.3	2763.5	6.2	20.7	21.4	2758.1	7.8	17.1	18.2	2758.1	17.1	100.5
Trial4_284	212	NA()	2.4	12.9529	1.0	1.6980	2.7	0.1596	2.5	0.92	954.5	6.0	17.5	18.0	1007.8	4.9	14.1	14.6	1126.4	7.2	19.8	21.2	1126.4	19.8	84.7
Trial4_285	133	NA()	1.2	13.1976	1.3	1.8506	3.2	0.1772	2.9	0.92	1051.7	8.0	19.7	20.2	1063.6	6.2	15.0	15.5	1089.0	9.0	20.6	22.0	1089.0	20.6	96.6
Trial4_286	153	NA()	1.3	5.1997	0.9	12.9961	1.7	0.4903	1.4	0.84	2572.1	10.0	40.7	41.8	2679.4	5.9	20.5	21.2	2762.1	6.7	16.6	17.8	2762.1	16.6	93.1
Trial4_287	178	NA()	1.6	11.1145	0.9	3.1095	2.3	0.2508	2.2	0.93	1442.5	8.2	25.4	26.0	1435.1	5.5	16.9	17.5	1424.9	6.5	18.9	20.2	1424.9	18.9	101.2
Trial4_288	33	NA()	1.0	17.9534	4.0	0.5207	5.2	0.0682	3.2	0.63	425.5	4.9	9.1	9.2	427.6	10.2	12.5	12.7	440.4	59.7	63.1	63.7	425.5	9.1	96.6
Trial4_289	39	NA()	1.2	5.2288	1.0	14.3363	3.2	0.5439	3.0	0.95	2799.8	12.8	44.1	45.3	2772.2	7.8	21.2	21.9	2752.9	9.8	18.1	19.2	2752.9	18.1	101.7
Trial4_290	61	NA()	2.1	13.7824	2.0	1.6665	4.1	0.1667	3.6	0.87	993.6	8.7	19.2	19.6	995.8	8.8	15.8	16.3	1001.6	20.5	27.8	28.9	1001.6	27.8	99.2
Trial4_291	202	NA()	3.7	18.1429	1.8	0.5678	3.7	0.0747	3.3	0.88	464.7	3.3	9.0	9.2	456.6	3.5	8.4	8.7	417.0	13.6	24.8	26.2	464.7	9.0	111.4
Trial4_292	90	NA()	2.4	18.2654	2.8	0.4954	4.7	0.0657	3.8	0.80	409.9	4.0	8.4	8.6	408.6	5.2	8.7	8.9	401.9	26.6	33.7	34.8	409.9	8.4	102.0
Trial4_293	110	NA()	1.4	12.7845	1.3	2.0421	3.3	0.1894	3.0	0.92	1118.3	8.2	20.8	21.3	1129.7	6.3	15.5	16.0	1152.4	9.2	20.6	21.9	1152.4	20.6	97.0
Trial4_294	38	NA()	1.6	5.3773	1.0	13.4778	3.2	0.5259	3.1	0.95	2724.0	12.9	43.3	44.4	2713.7	7.8	21.1	21.8	2706.8	9.7	18.1	19.2	2706.8	18.1	100.6
Trial4_295	71	NA()	1.0	10.9593	1.3	3.1686	3.4	0.2520	3.2	0.92	1448.6	10.2	26.2	26.8	1449.5	7.2	17.6	18.1	1451.7	9.3	19.9	21.2	1451.7	19.9	99.8
Trial4_296	212	NA()	3.1	10.8982	0.8	3.0269	2.1	0.2394	2.0	0.92	1383.3	6.7	24.1	24.7	1414.4	4.7	16.5	17.1	1462.4	5.6	18.5	19.9	1462.4	18.5	94.6
Trial4_297	117	NA()	1.0	19.7738	3.4	0.1986	4.7	0.0285	3.2	0.68	181.1	1.7	3.7	3.8	183.9	3.3	4.8	4.9	221.3	39.1	44.6	45.4	181.1	3.7	NA
Trial4_298	37	NA()	1.3	13.5279	2.4	1.6979	4.4	0.1667	3.8	0.85	993.7	9.0	19.4	19.8	1007.7	8.6	15.8	16.3	1039.3	18.6	26.4	27.5	1039.3	26.4	95.6
Trial4_299	41	NA()	1.0	5.4917	1.0	12.9852	3.1	0.5174	3.0	0.95	2688.3	12.5	42.8	43.9	2678.6	7.6	21.0	21.7	2672.0	9.4	18.0	19.1	2672.0	18.0	100.6
Trial4_300	134	NA()	1.2	13.7759	1.3	1.7428	3.2	0.1742	2.9	0.91	1035.2	7.2	19.2	19.6	1024.5	5.7	14.6	15.0	1002.5	9.4	21.0	22.4	1002.5	21.0	103.3
Trial4_301	691	NA()	1.1	19.9935	1.4	0.2793	3.0	0.0405	2.7	0.89	256.0	1.1	4.8	4.9	250.1	1.4	4.8	5.0	195.8	10.8	24.1	25.7	256.0	4.8	NA
Trial4_302	90	NA()	1.7	14.0504	1.6	1.6215	3.7	0.1653	3.3	0.90	986.2	7.9	18.8	19.2	978.6	6.5	14.6	15.1	962.4	11.9	22.3	23.7	962.4	22.3	102.5
Trial4_303	235	NA()	2.8	13.1219	1.0	1.9206	2.4	0.1829	2.2	0.92	1082.6	5.6	19.4	19.8	1088.3	4.3	14.5	15.0	1100.6	6.5	19.6	21.1	1100.6	19.6	98.4
Trial4_304	80	NA()	2.4	13.3982	1.6	1.7750	3.7	0.1726	3.4	0.90	1026.2	8.4	19.5	20.0	1036.3	7.0	15.2	15.7	1058.8	12.4	22.4	23.7	1058.8	22.4	96.9
Trial4_305	332	NA()	2.1	10.1564	0.8	3.5444	1.4	0.2612	1.2	0.82	1496.0	4.5	25.2	25.9	1537.2	3.2	16.8	17.4	1595.1	4.0	17.8	19.1	1595.1	17.8	93.8
Trial4_306	28	NA()	1.7	5.2642	1.1	13.8037	3.5	0.5273	3.3	0.94	2729.9	13.4	43.5	44.6	2736.3	8.2	21.3	22.0	2741.8	10.2	18.3	19.4	2741.8	18.3	99.6
Trial4_307	91	NA()	1.4	11.2649	1.2	2.9251	3.3	0.2391	3.0	0.92	1382.0	9.6	25.1	25.7	1388.4	6.8	17.1	17.7	1399.2	8.9	19.9	21.2	1399.2	19.9	98.8
Trial4_308	130	NA()	0.7	10.7066	1.0	3.2866	2.7	0.2553	2.6	0.93	1465.9	8.9	26.0	26.6	1477.9	6.0	17.3	17.9	1496.0	7.2	18.9	20.3	1496.0	18.9	98.0
Trial4_309	187	NA()	1.4	20.0034	2.6	0.2800	4.5	0.0406	3.6	0.81	256.8	1.9	5.1	5.2	250.7	2.9	5.4	5.6	194.6	24.4	32.6	33.8	256.8	5.1	NA
Trial4_310	136	NA()	1.3	19.8210	3.0	0.3001	4.8	0.0432	3.8	0.78	272.4	2.4	5.5	5.6	266.5	3.8	6.2	6.3	215.9	31.2	37.9	38.9	272.4	5.5	NA
Trial4_311	54	NA()	1.5	8.8542	1.2	4.9312	3.4	0.3168	3.2	0.93	1774.1	11.8	31.2	31.9	1807.6	7.8	19.2	19.8	1847.2	9.4	19.2	20.4	1847.2	19.2	96.0
Trial4_312	87	NA()	3.6	12.3778	1.4	2.3813	3.4	0.2139	3.1	0.92	1249.4	9.1	23.0	23.5	1237.0	6.8	16.3	16.9	1216.4	10.0	20.8	22.1	1216.4	20.8	102.7
Trial4_313	53	NA()	2.7	13.5243	2.0	1.7445	4.1	0.1712	3.6	0.88	1018.7	8.7	19.6	20.0	1025.1	7.6	15.4	15.9	1039.8	14.6	23.8	25.0	1039.8	23.8	98.0
Trial4_314	102	NA()	1.1	17.4535	2.3	0.6990	4.3	0.0885	3.6	0.84	546.8	4.9	10.9	11.2	538.2	5.6	10.3	10.6	502.9	21.2	29.4	30.6	546.8	10.9	108.7
Trial4_315	152	NA()	2.6	10.0415	0.9	4.0188	2.4	0.2928	2.2	0.93	1655.6	9.3	28.7	29.4	1638.0	5.9	17.9	18.5	1616.4	6.6	18.5	19.8	1616.4	18.5	102.4
Trial4_316	73	NA()	1.2	19.8248	3.8	0.2671	5.0	0.0384	3.2	0.64	243.1	2.7	5.2	5.3	240.4	5.6	7.2	7.3	215.4	54.9	58.9	59.6	243.1	5.2	NA
Trial4_317	67	NA()	0.9	10.9717	1.4	3.1269	3.5	0.2489	3.2	0.92	1433.0	10.9	26.3	26.9	1439.3	7.7	17.7	18.3	1449.6	9.9	20.2	21.5	1449.6	20.2	98.9
Trial4_318	71	NA()	1.1	12.3081	1.5	2.2700	3.7	0.2027	3.3	0.91	1189.9	9.7	22.4	22.9	1203.0	7.4	16.4	16.9	1227.4	10.9	21.2	22.5	1227.4	21.2	96.9
Trial4_319	87	NA()	4.6	9.6960	1.1	4.0021	3.1	0.2816	2.9	0.93	1599.2	11.0	28.5	29.2	1634.6	7.3	18.4	19.0	1681.3	8.4	19.1	20.3	1681.3	19.1	95.1
Trial4_320	26	NA()	1.1	11.4058	2.3	2.7781	4.4	0.2299	3.8	0.85	1334.1	11.8	25.4	25.9	1349.7	10.4	18.7	19.2	1375.4	19.0	26.1	27.1	1375.4	26.1	97.0
Trial4_321	109	NA()	1.5	9.4142	1.0	4.3994	2.7	0.3005	2.6	0.94	1693.9	10.9	29.8	30.5	1712.2	6.9	18.5	19.2	1735.6	7.4	18.5	19.8	1735.6	18.5	97.6
Trial4_322	87	NA()	1.9	20.1008	3.7	0.2552	4.9	0.0372	3.2	0.65	235.6	2.6	5.0	5.1	230.8	5.1	6.7	6.8	183.3	51.9	56.2	56.9	235.6	5.0	NA
Trial4_323	208	NA()	3.9	13.4076	1.0	1.8888	2.6	0.1837	2.4	0.92	1087.4	6.6	19.7	20.2	1077.2	4.9	14.7	15.2	1057.3	7.0	19.9	21.4	1057.3	19.9	102.8
Trial4_324	55	NA()	0.8	13.0711	1.8	2.0691	4.0	0.1962	3.6	0.89	1155.1	10.4	22.2	22.7	1138.6	8.2	16.4	16.9	1108.3	13.9	23.1	24.4	1108.3	23.1	104.2
Trial4_325	196	NA()	3.1	13.8116	1.1	1.7509	2.7	0.1755	2.5	0.92	1042.2	6.8	19.2	19.6	1027.5	5.2	14.4	14.9	997.3	7.5	20.3	21.7	997.3	20.3	104.5
Trial4_326	49	NA()	0.7	5.5731	1.0	12.6703	3.0	0.5124	2.8	0.95	2666.7	13.5	42.8	43.9	2655.5	7.9	21.1	21.8	2647.6	9.6	18.1	19.2	2647.6	18.1	100.7
Trial4_327	150	NA()	1.8	13.5929	1.2	1.7915	3.1	0.1767	2.8	0.92	1048.9	7.5	19.5	20.0	1042.4	5.8	14.7	15.2	1029.6	8.6	20.6	22.0	1029.6	20.6	101.9
Trial4_328	184	NA()	1.6	13.4697	1.1	1.8799	2.7	0.1837	2.5	0.92	1087.3	7.0	19.9	20.4	1074.0	5.3	14.8	15.3	1048.0	7.7	20.2	21.6	1048.0	20.2	103.8
Trial4_329	128	NA()	1.4	20.1085	3.0	0.3034	4.8	0.0443	3.8	0.78	279.2	2.7	5.7	5.9	269.0	3.7	6.2	6.3	182.4	29.0	36.1	37.2	279.2	5.7	NA
Trial4_330	81	NA()	1.3	20.0023	3.6	0.2673	4.8	0.0388	3.2	0.66	245.3	2.7	5.2	5.3	240.5	5.0	6.7	6.8	194.7	47.6	52.2	53.0	245.3	5.2	NA
Trial4_331	112	NA()	2.3	13.0904	1.3	2.0626	3.3	0.1959	3.0	0.92	1153.3	8.7	21.5	22.0	1136.5	6.5	15.6	16.1	1105.4	9.3	20.7	22.1	1105.4	20.7	104.3
Trial4_332	33	NA()	1.5	5.6021	1.1	12.3219	3.4	0.5009	3.2	0.94	2617.5	13.8	42.3	43.4	2629.2	8.3	21.2	21.9	2639.0	10.1	18.4	19.5	2639.0	18.4	99.2
Trial4_333	32	NA()	3.2	9.2458	1.7	4.5216	3.9	0.3033	3.6	0.91	1707.9	13.2	30.9	31.6	1735.0	9.2	19.6	20.2	1768.6	12.4	21.0	22.1	1768.6	21.0	96.6
Trial4_334	89	NA()	1.2	17.9784	3.5	0.2660	4.7	0.0347	3.2	0.68	219.9	2.3	4.7	4.8	239.5	4.3	6.2	6.3	437.3	38.3	43.5	44.3	219.9	4.7	NA
Trial4_335	46	NA()	1.3	11.1347	1.6	3.2152	3.8	0.2598	3.5	0.91	1488.6	12.1	27.5	28.1	1460.8	8.6	18.2	18.8	1421.5	12.0	21.4	22.6	1421.5	21.4	104.7
Trial4_336	151	NA()	1.0	11.5952	1.1	2.1887	3.0	0.1841	2.8	0.93	1089.6	7.8	20.2	20.7	1177.5	6.1	15.7	16.2	1343.6	7.6	19.4	20.8	1343.6	19.4	81.1
Trial4_337	185	NA()	2.1	10.6717	0.9	3.2006	2.3	0.2478	2.1	0.93	1427.3	7.6	25.0	25.6	1457.3	5.2	16.9	17.5	1502.2	6.0	18.5	19.9	1502.2	18.5	95.0
Trial4_338	142	NA()	1.4	11.3115	1.0	3.0373	2.7	0.2493	2.5	0.93	1434.8	8.6	25.4	26.0	1417.0	5.8	16.9	17.5	1391.3	7.2	19.2	20.5	1391.3	19.2	103.1
Trial4_339	83	NA()	1.0	11.7603	1.3	2.7093	3.4	0.2312	3.1	0.92	1340.7	9.7	24.5	25.1	1331.0	7.0	16.9	17.5	1316.3	9.6	20.4	21.7	1316.3	20.4	101.9
Trial4_340	27	NA()	1.0	13.5061	2.8	1.6904	4.8	0.1657	3.9	0.81	988.2	9.4	19.5	19.9	1004.9	9.4	16.3	16.7	1042.6	21.4	28.4	29.5	1042.6	28.4	94.8
Trial4_341	531	NA()	8.7	18.0975	1.2	0.5154	2.6	0.0677	2.3	0.89	422.2	0.6	7.6	7.8	422.1	1.4	7.3	7.6	422.6	8.7	22.4	24.0	422.2	7.6	99.9
Trial4_342	114	NA()	1.1	2.9399	0.9	34.7092	1.5	0.7404	1.2	0.79	3572.1	7.0	51.5	52.9	3630.4	5.3	21.2	21.9	3663.4	7.1	15.9	16.9	3663.4	15.9	97.5
Trial4_343	241	NA()	2.8	13.4073	1.0	1.6974	2.5	0.1651	2.3	0.92	985.2	4.8	17.6	18.1	1007.6	3.9	13.8	14.3	1057.4	6.6	19.8	21.2	1057.4	19.8	93.2
Trial4_344	295	NA()	2.4	17.4483	1.4	0.6689	3.1	0.0847	2.8	0.90	524.0	2.3	9.6	9.9	520.0	2.7	8.9	9.2	503.6	10.8	23.1	24.6	524.0	9.6	104.1
Trial4_345	30	NA()	1.0	13.7017	2.6	1.7532	4.6	0.1743	3.8	0.82	1035.7	9.3	20.1	20.5	1028.3	9.5	16.5	16.9	1013.5	22.3	29.1	30.1	1013.5	29.1	102.2
Trial4_346	303	NA()	2.0	11.3134	0.8	2.8038	1.7	0.2302	1.5	0.87	1335.4	4.4	22.9	23.5	1356.6	3.3	15.9	16.5	1391.0	4.6	18.4	19.8	1391.0	18.4	96.0
Trial4_347	93	NA()	0.8	5.4909	0.9	13.4010	2.2	0.5339	2.0	0.90	2757.9	10.4	43.0	44.1	2708.3	6.2	20.6	21.3	2672.3	7.7	17.2	18.3	2672.3	17.2	103.2
Trial4_348	242	NA()	2.5	13.5176	1.0	1.8574	2.4	0.1822	2.2	0.91	1078.8	4.6	19.1	19.6	1066.0	3.8	14.2	14.7	1040.8	6.6	19.9	21.3	1040.8	19.9	103.7
Trial4_349	250	NA()	3.4	9.4306	0.8	4.4267	1.6	0.3029	1.4	0.86	1705.7	5.8	28.5	29.2	1717.4	3.8	17.6	18.3	1732.4	4.5	17.6	18.9	1732.4	17.6	98.5
Trial4_350	180	NA()	1.8	13.9439	1.2	1.5830	2.9	0.1602	2.7	0.92	957.7	5.4	17.4	17.8	963.5	4.5	13.7	14.2	977.9	8.1	20.5	22.0	977.9	20.5	97.9
Trial4_351	172	NA()	3.5	9.3820	0.9	4.4301	2.1	0.3016	2.0	0.92	1699.1	7.2	28.7	29.4	1718.0	4.7	17.8	18.5	1741.8	5.5	17.9	19.2	1741.8	17.9	97.5
Trial4_352	40	NA()	0.6	10.9717	1.7	3.1126	3.9	0.2478	3.5	0.90	1427.1	10.0	25.8	26.4	1435.8	7.7	17.7	18.3	1449.6	12.1	21.4	22.6	1449.6	21.4	98.4
Trial4_353	43	NA()	0.9	5.4998	1.0	12.7575	3.1	0.5091	3.0	0.95	2652.8	11.3	42.0	43.1	2661.9	6.8	20.7	21.4	2669.6	8.4	17.5	18.6	2669.6	17.5	99.4
Trial4_354	37	NA()	1.5	13.3642	2.3	1.9024	4.3	0.1845	3.7	0.85	1091.4	8.7	20.6	21.1	1081.9	7.9	15.9	16.4	1063.9	16.3	24.8	25.9	1063.9	24.8	102.6
Trial4_355	158	NA()	1.3	19.6994	3.0	0.2585	4.8	0.0369	3.7	0.78	233.9	1.7	4.6	4.7	233.4	3.2	5.4	5.5	230.1	30.9	37.6	38.6	233.9	4.6	NA
Trial4_356	150	NA()	1.4	19.6558	3.0	0.2837	4.8	0.0405	3.7	0.78	255.7	1.9	5.0	5.1	253.6	3.6	5.9	6.0	235.2	33.1	39.4	40.3	255.7	5.0	NA
Trial4_357	121	NA()	2.0	9.3794	0.9	4.7488	2.5	0.3232	2.3	0.94	1805.3	8.7	30.6	31.3	1775.9	5.5	18.3	18.9	1742.3	6.5	18.2	19.5	1742.3	18.2	103.6
Trial4_358	633	NA()	3.8	13.7705	0.8	1.7417	1.3	0.1740	1.1	0.79	1034.3	3.4	18.1	18.6	1024.1	2.6	13.7	14.2	1003.3	4.1	19.2	20.7	1003.3	19.2	103.1
Trial4_359	65	NA()	1.3	11.9358	1.6	2.4341	3.7	0.2108	3.4	0.91	1233.1	8.7	22.6	23.2	1252.7	7.0	16.5	17.0	1287.5	11.2	21.2	22.5	1287.5	21.2	95.8
Trial4_360	283	NA()	3.7	13.4186	0.9	1.6926	2.3	0.1648	2.1	0.91	983.4	3.3	17.3	17.7	1005.7	3.0	13.6	14.1	1055.7	6.0	19.6	21.1	1055.7	19.6	93.2
Trial4_361	143	NA()	2.5	9.7703	0.9	3.9989	2.4	0.2835	2.3	0.93	1608.9	7.8	27.6	28.3	1634.0	5.2	17.7	18.3	1667.2	6.2	18.2	19.5	1667.2	18.2	96.5
Trial4_362	32	NA()	0.9	13.5468	2.6	1.6996	4.6	0.1671	3.8	0.83	995.9	8.6	19.2	19.6	1008.4	8.9	16.0	16.4	1036.5	20.9	28.1	29.1	1036.5	28.1	96.1
Trial4_363	382	NA()	3.5	19.6989	1.9	0.2560	3.8	0.0366	3.2	0.86	231.7	0.6	4.3	4.4	231.4	1.6	4.6	4.7	230.1	16.5	27.0	28.4	231.7	4.3	NA
Trial4_364	60	NA()	1.4	20.0505	3.7	0.2634	4.9	0.0383	3.2	0.65	242.4	2.5	5.1	5.2	237.4	5.0	6.7	6.8	189.1	50.0	54.4	55.1	242.4	5.1	NA
Trial4_365	61	NA()	4.0	13.5822	1.8	1.8003	3.9	0.1774	3.5	0.89	1052.9	8.4	19.9	20.4	1045.6	7.1	15.3	15.8	1031.2	13.4	23.1	24.3	1031.2	23.1	102.1
Trial4_366	308	NA()	0.9	9.9090	0.8	3.9833	1.4	0.2864	1.2	0.81	1623.5	4.5	27.1	27.8	1630.8	3.1	17.2	17.8	1641.1	4.0	17.7	19.0	1641.1	17.7	98.9
Trial4_367	398	NA()	1.7	16.5513	1.1	0.8137	2.5	0.0977	2.3	0.90	601.0	1.3	10.8	11.0	604.5	2.0	9.7	10.0	618.6	8.0	21.5	23.1	601.0	10.8	97.2
Trial4_368	73	NA()	1.2	13.4575	1.7	1.6963	3.8	0.1656	3.4	0.89	988.0	7.8	18.7	19.2	1007.1	6.9	14.9	15.4	1049.9	13.3	22.9	24.2	1049.9	22.9	94.1
Trial4_369	167	NA()	2.5	13.2310	1.2	1.7789	2.9	0.1708	2.7	0.92	1016.4	6.1	18.5	19.0	1037.8	4.9	14.4	14.9	1084.0	7.9	20.2	21.6	1084.0	20.2	93.8
Trial4_370	15	NA()	1.4	14.0547	3.5	1.6494	4.8	0.1682	3.2	0.67	1002.2	9.8	19.9	20.3	989.3	13.5	18.9	19.2	961.8	37.9	42.4	43.1	961.8	42.4	104.2
Trial4_371	309	NA()	2.9	12.3720	0.8	2.0823	1.9	0.1869	1.7	0.91	1104.7	3.6	19.2	19.7	1143.0	3.0	14.6	15.1	1217.3	5.1	18.9	20.4	1217.3	18.9	90.8
Trial4_372	44	NA()	1.6	5.5234	1.0	11.3733	3.2	0.4558	3.0	0.95	2421.0	12.2	39.5	40.5	2554.2	7.3	20.7	21.4	2662.5	8.3	17.5	18.6	2662.5	17.5	90.9
Trial4_373	40	NA()	1.8	12.6317	2.1	2.0888	4.2	0.1914	3.7	0.87	1129.2	9.4	21.4	21.9	1145.1	8.4	16.6	17.0	1176.3	16.2	24.5	25.6	1176.3	24.5	96.0

Trial4_374	29	NA()	1.9	13.2856	2.6	1.7908	4.6	0.1726	3.8	0.82	1026.6	9.2	19.9	20.3	1042.1	9.3	16.4	16.8	1075.7	21.0	28.0	29.1	1075.7	28.0	95.4
Trial4_375	33	NA()	1.5	13.3420	2.4	1.8514	4.5	0.1792	3.8	0.84	1062.8	9.7	20.6	21.1	1063.9	9.2	16.5	17.0	1067.2	19.9	27.3	28.3	1067.2	27.3	99.6
Trial4_376	216	NA()	2.5	12.5579	0.9	2.2247	2.3	0.2027	2.2	0.92	1189.9	6.3	21.2	21.7	1188.8	4.6	15.3	15.8	1187.9	6.3	19.4	20.8	1187.9	19.4	100.2
Trial4_377	109	NA()	1.4	17.9737	2.5	0.5269	4.4	0.0687	3.7	0.83	428.4	3.9	8.6	8.8	429.7	4.6	8.6	8.9	437.9	20.7	29.3	30.5	428.4	8.6	97.8
Trial4_378	88	NA()	2.4	13.3407	1.5	1.8675	3.6	0.1808	3.3	0.91	1071.2	8.4	20.2	20.7	1069.6	6.7	15.3	15.8	1067.4	10.9	21.6	22.9	1067.4	21.6	100.4
Trial4_379	179	NA()	2.4	4.5680	0.9	16.9205	1.4	0.5608	1.1	0.77	2870.1	8.4	43.9	45.1	2930.3	5.0	20.6	21.3	2972.7	6.1	16.1	17.3	2972.7	16.1	96.5
Trial4_380	396	NA()	1.5	18.6921	1.4	0.5038	3.0	0.0683	2.7	0.89	426.1	1.2	7.8	8.0	414.3	1.9	7.3	7.6	350.0	10.5	23.4	24.9	426.1	7.8	121.8
Trial4_381	73	NA()	1.7	8.2298	1.0	5.9891	3.0	0.3576	2.8	0.94	1971.0	12.0	33.8	34.6	1974.3	7.4	19.5	20.2	1978.5	8.3	18.5	19.7	1978.5	18.5	99.6
Trial4_382	71	NA()	1.6	13.2622	1.6	1.9544	3.8	0.1881	3.4	0.90	1110.9	9.1	21.1	21.5	1100.0	7.2	15.7	16.2	1079.3	11.7	22.0	23.3	1079.3	22.0	102.9
Trial4_383	19	NA()	2.1	13.2740	3.2	1.8906	4.6	0.1821	3.2	0.70	1078.4	10.7	21.3	21.8	1077.8	12.4	18.6	19.0	1077.5	30.7	35.9	36.7	1077.5	35.9	100.1
Trial4_384	74	NA()	1.0	16.6432	2.4	0.8284	4.4	0.1000	3.7	0.84	614.6	5.9	12.4	12.7	612.7	6.5	11.6	11.9	606.6	21.5	29.4	30.6	614.6	12.4	101.3
Trial4_385	454	NA()	2.3	19.4317	1.6	0.3018	3.4	0.0426	3.0	0.88	268.7	0.6	4.9	5.1	267.8	1.5	5.1	5.3	261.6	13.5	25.2	26.7	268.7	4.9	NA
Trial4_386	101	NA()	1.6	17.7472	2.5	0.5527	4.5	0.0712	3.7	0.83	443.2	4.3	9.1	9.2	446.8	5.1	9.1	9.3	466.0	21.9	30.0	31.2	443.2	9.1	95.1
Trial4_387	82	NA()	1.5	12.7201	1.5	2.1249	3.6	0.1961	3.3	0.91	1154.4	9.5	21.8	22.3	1156.9	7.3	16.1	16.6	1162.5	11.0	21.4	22.7	1162.5	21.4	99.3
Trial4_388	711	NA()	4.2	10.9783	0.8	3.1483	1.3	0.2508	1.0	0.79	1442.5	4.0	24.4	25.0	1444.6	2.7	16.2	16.9	1448.4	3.1	17.9	19.3	1448.4	17.9	99.6
Trial4_389	39	NA()	1.8	13.8426	2.4	1.6360	4.5	0.1643	3.8	0.85	980.7	9.5	19.4	19.8	984.2	8.8	15.8	16.2	992.7	18.8	26.6	27.7	992.7	26.6	98.8
Trial4_390	27	NA()	1.2	11.0893	2.0	3.2493	4.2	0.2614	3.7	0.88	1497.2	12.8	28.0	28.6	1469.0	9.6	18.8	19.3	1429.3	14.8	23.0	24.2	1429.3	23.0	104.8
Trial4_391	61	NA()	1.4	16.6297	2.9	0.7309	4.8	0.0882	3.8	0.80	544.9	5.7	11.2	11.5	557.1	7.0	11.3	11.6	608.4	26.5	33.2	34.3	544.9	11.2	89.6
Trial4_392	271	NA()	2.1	11.0052	0.8	3.1525	1.8	0.2517	1.6	0.88	1447.4	6.3	24.9	25.6	1445.6	4.3	16.6	17.2	1443.8	5.0	18.4	19.7	1443.8	18.4	100.3
Trial4_393	55	NA()	1.7	11.8216	1.6	2.6572	3.8	0.2279	3.4	0.90	1323.6	11.0	24.8	25.4	1316.7	8.1	17.3	17.9	1306.2	11.7	21.5	22.7	1306.2	21.5	101.3
Trial4_394	49	NA()	1.4	20.4835	4.2	0.2761	5.3	0.0410	3.2	0.61	259.3	3.0	5.6	5.7	247.6	6.8	8.2	8.3	139.2	67.4	70.8	71.4	259.3	5.6	NA
Trial4_395	70	NA()	1.4	6.1603	0.9	10.0975	2.8	0.4513	2.6	0.94	2401.2	12.3	39.3	40.2	2443.7	7.2	20.5	21.2	2480.1	8.2	17.6	18.8	2480.1	17.6	96.8
Trial4_396	42	NA()	2.8	9.2545	1.4	4.7776	3.7	0.3208	3.4	0.92	1793.7	12.7	31.8	32.5	1781.0	8.4	19.4	20.0	1766.9	10.6	20.0	21.1	1766.9	20.0	101.5
Trial4_397	156	NA()	0.6	5.4718	0.9	13.3358	1.6	0.5295	1.3	0.82	2739.2	9.4	42.6	43.7	2703.7	5.6	20.4	21.1	2678.0	6.8	16.8	17.9	2678.0	16.8	102.3
Trial4_398	198	NA()	2.1	9.5790	0.9	4.3888	2.0	0.3050	1.8	0.90	1716.3	8.2	29.2	30.0	1710.2	5.2	18.0	18.6	1703.7	5.6	18.0	19.3	1703.7	18.0	100.7
Trial4_399	66	NA()	2.2	5.9018	0.9	11.3451	2.7	0.4858	2.6	0.94	2552.6	12.3	41.1	42.1	2551.9	7.2	20.7	21.4	2552.1	8.5	17.7	18.8	2552.1	17.7	100.0
Trial4_400	73	NA()	2.1	11.2930	1.3	3.0854	3.4	0.2528	3.2	0.92	1453.0	10.7	26.5	27.1	1429.1	7.5	17.6	18.2	1394.4	10.1	20.5	21.7	1394.4	20.5	104.2
Trial4_401	53	NA()	0.8	10.8712	1.5	3.2972	3.7	0.2601	3.4	0.91	1490.3	11.3	27.2	27.8	1480.4	8.0	18.0	18.6	1467.1	10.8	20.7	21.9	1467.1	20.7	101.6
Trial4_402	76	NA()	0.9	7.9235	1.0	6.5907	2.9	0.3789	2.7	0.94	2071.2	11.9	35.0	35.9	2058.1	7.2	19.7	20.3	2045.8	8.1	18.3	19.5	2045.8	18.3	101.2
Trial4_403	143	NA()	1.3	13.1314	1.2	1.9647	3.1	0.1872	2.8	0.92	1106.2	7.5	20.3	20.8	1103.5	5.7	15.1	15.6	1099.1	8.4	20.4	21.8	1099.1	20.4	100.6
Trial4_404	118	NA()	0.7	6.0760	0.9	11.0527	2.0	0.4873	1.8	0.89	2558.9	10.7	40.7	41.8	2527.6	6.2	20.3	21.0	2503.3	7.4	17.2	18.4	2503.3	17.2	102.2
Trial4_405	48	NA()	1.1	11.2594	1.6	3.0128	3.8	0.2461	3.5	0.91	1418.5	11.1	26.2	26.7	1410.9	8.1	17.8	18.4	1400.1	11.9	21.4	22.6	1400.1	21.4	101.3
Trial4_406	152	NA()	1.0	17.6562	1.9	0.6536	3.9	0.0837	3.4	0.87	518.4	4.1	10.1	10.4	510.7	4.4	9.4	9.7	477.4	16.5	26.3	27.6	518.4	10.1	108.6
Trial4_407	147	NA()	2.4	9.4097	0.8	4.6147	2.3	0.3151	2.1	0.93	1765.6	9.6	30.3	31.0	1751.9	6.0	18.3	19.0	1736.4	6.5	18.2	19.5	1736.4	18.2	101.7
Trial4_408	223	NA()	3.0	18.5690	1.9	0.4413	3.9	0.0595	3.3	0.87	372.3	2.4	7.2	7.3	371.1	3.0	7.1	7.4	364.9	16.0	26.3	27.7	372.3	7.2	NA
Trial4_409	72	NA()	2.1	6.7894	0.9	8.5171	2.8	0.4196	2.6	0.95	2258.6	11.9	37.4	38.3	2287.8	7.1	20.2	20.9	2314.7	8.1	17.8	19.0	2314.7	17.8	97.6
Trial4_410	55	NA()	1.0	11.0916	1.5	3.0906	3.7	0.2487	3.3	0.91	1431.9	10.8	26.2	26.8	1430.4	7.7	17.7	18.3	1428.9	10.6	20.6	21.9	1428.9	20.6	100.2
Trial4_411	74	NA()	1.7	11.6162	1.4	2.7137	3.5	0.2287	3.2	0.92	1327.8	9.9	24.4	25.0	1332.2	7.3	17.1	17.6	1340.1	10.3	20.6	21.9	1340.1	20.6	99.1
Trial4_412	39	NA()	1.0	5.6249	1.0	12.5264	3.2	0.5113	3.0	0.95	2662.0	12.7	42.5	43.6	2644.7	7.7	21.0	21.7	2632.3	9.6	18.2	19.2	2632.3	18.2	101.1
Trial4_413	20	NA()	1.1	12.7583	3.0	1.9991	4.9	0.1851	3.9	0.79	1094.6	10.7	21.6	22.0	1115.2	11.7	18.3	18.7	1156.5	27.0	32.7	33.5	1156.5	32.7	94.6
Trial4_414	108	NA()	1.5	5.2687	0.9	13.9800	2.0	0.5345	1.8	0.89	2760.2	10.5	43.1	44.2	2748.4	6.4	20.7	21.4	2740.4	7.9	17.2	18.3	2740.4	17.2	100.7
Trial4_415	68	NA()	2.2	13.2859	1.7	1.8072	3.8	0.1742	3.4	0.90	1035.3	8.9	19.9	20.3	1048.1	7.3	15.4	15.9	1075.7	12.4	22.4	23.7	1075.7	22.4	96.2
Trial4_416	428	NA()	0.8	18.4295	1.3	0.5476	2.8	0.0732	2.5	0.89	455.6	1.1	8.3	8.5	443.4	1.8	7.7	8.0	381.9	9.4	22.9	24.4	455.6	8.3	119.3
Trial4_417	56	NA()	2.5	12.7719	1.7	2.0882	3.9	0.1935	3.5	0.90	1140.4	9.9	21.8	22.3	1144.9	7.9	16.3	16.8	1154.4	12.8	22.5	23.7	1154.4	22.5	98.8
Trial4_418	767	NA()	13.5	13.6670	0.8	1.7347	1.4	0.1720	1.1	0.80	1023.3	4.3	18.1	18.6	1021.5	3.1	13.8	14.3	1018.6	3.9	19.2	20.7	1018.6	19.2	100.5
Trial4_419	41	NA()	1.7	9.5918	1.5	4.3105	3.7	0.3000	3.4	0.92	1691.3	12.7	30.4	31.1	1695.4	8.6	19.2	19.8	1701.2	11.0	20.3	21.5	1701.2	20.3	99.4
Trial4_420	138	NA()	0.9	13.7537	1.3	1.7023	3.2	0.1699	2.9	0.92	1011.5	7.5	19.0	19.4	1009.4	5.9	14.5	15.0	1005.8	9.3	21.0	22.4	1005.8	21.0	100.6
Trial4_421	55	NA()	2.0	13.7769	1.9	1.7714	4.1	0.1771	3.6	0.88	1051.0	9.4	20.4	20.8	1035.0	8.0	15.7	16.1	1002.4	15.4	24.3	25.5	1002.4	24.3	104.9
Trial4_422	103	NA()	0.7	11.4873	1.2	2.9406	3.1	0.2451	2.9	0.93	1413.2	10.1	25.7	26.3	1392.4	6.9	17.2	17.8	1361.6	8.7	19.8	21.2	1361.6	19.8	103.8
Trial4_423	87	NA()	1.6	12.9155	1.4	2.0616	3.5	0.1932	3.2	0.91	1138.7	9.3	21.5	22.0	1136.2	7.1	15.9	16.4	1132.2	10.5	21.2	22.5	1132.2	21.2	100.6
Trial4_424	646	NA()	2.2	17.7375	1.1	0.5401	2.3	0.0695	2.1	0.89	433.2	1.1	7.9	8.1	438.5	1.5	7.6	7.8	467.3	7.5	21.9	23.5	433.2	7.9	92.7
Trial4_425	187	NA()	1.6	8.4260	0.9	5.6402	1.8	0.3448	1.6	0.88	1909.9	9.0	32.0	32.9	1922.3	5.5	18.7	19.4	1936.4	5.8	17.6	18.9	1936.4	17.6	98.6
Trial4_426	98	NA()	0.8	13.2192	1.4	1.9224	3.5	0.1844	3.2	0.91	1090.9	8.7	20.6	21.1	1088.9	6.7	15.4	15.9	1085.7	10.0	21.1	22.4	1085.7	21.1	100.5
Trial4_427	64	NA()	1.1	13.9633	1.8	1.6609	4.0	0.1683	3.5	0.89	1002.6	8.9	19.4	19.9	993.7	7.4	15.1	15.6	975.1	13.6	23.2	24.5	975.1	23.2	102.8
Trial4_428	78	NA()	0.7	19.5176	4.1	0.1965	5.2	0.0278	3.2	0.62	177.0	2.0	3.8	3.9	182.2	5.1	6.1	6.2	251.5	64.9	68.3	68.9	177.0	3.8	NA
Trial4_429	174	NA()	2.1	11.1718	0.9	3.0714	2.3	0.2490	2.2	0.93	1433.2	8.6	25.4	26.0	1425.6	5.7	16.9	17.5	1415.1	6.6	18.9	20.3	1415.1	18.9	101.3
Trial4_430	24	NA()	1.1	7.7555	1.6	6.5121	3.9	0.3665	3.6	0.91	2012.7	14.4	35.2	36.0	2047.5	9.5	20.6	21.3	2083.6	12.2	20.4	21.4	2083.6	20.4	96.6
Trial4_431	15	NA()	1.9	13.2840	3.4	1.9821	4.7	0.1910	3.2	0.69	1127.0	11.5	22.4	22.9	1109.4	12.1	18.5	19.0	1076.0	28.3	33.9	34.7	1076.0	33.9	104.7
Trial4_432	85	NA()	1.0	8.5035	1.0	5.5726	2.8	0.3438	2.7	0.94	1905.1	12.0	32.9	33.7	1911.9	7.4	19.4	20.0	1920.0	8.1	18.5	19.7	1920.0	18.5	99.2
Trial4_433	61	NA()	1.8	13.6888	1.9	1.6596	4.1	0.1648	3.6	0.88	983.6	9.0	19.2	19.6	993.2	7.9	15.4	15.8	1015.4	15.3	24.2	25.4	1015.4	24.2	96.9
Trial4_434	78	NA()	1.1	11.1902	1.3	3.0697	3.4	0.2492	3.1	0.92	1434.6	10.9	26.3	26.9	1425.2	7.6	17.6	18.2	1412.0	9.6	20.2	21.4	1412.0	20.2	101.6
Trial4_435	54	NA()	1.9	12.8415	1.8	2.0940	4.0	0.1951	3.5	0.89	1149.0	10.2	22.1	22.6	1146.9	8.1	16.4	16.9	1143.6	13.3	22.7	23.9	1143.6	22.7	100.5
Trial4_436	183	NA()	2.7	12.5780	1.0	2.1036	2.7	0.1920	2.5	0.92	1132.1	6.7	20.4	20.9	1150.0	5.1	15.2	15.7	1184.7	7.1	19.6	21.0	1184.7	19.6	95.6
Trial4_437	30	NA()	1.6	10.8787	2.1	3.0938	4.2	0.2442	3.7	0.87	1408.6	11.6	26.2	26.8	1431.2	9.7	18.7	19.2	1465.8	16.7	24.3	25.3	1465.8	24.3	96.1
Trial4_438	73	NA()	1.6	14.0587	1.8	1.6432	3.9	0.1676	3.4	0.89	999.0	8.2	19.1	19.5	987.0	7.0	14.9	15.3	961.2	13.5	23.2	24.5	961.2	23.2	103.9
Trial4_439	82	NA()	2.5	9.6616	1.2	3.7535	3.2	0.2631	3.0	0.93	1505.8	10.3	27.0	27.7	1582.9	7.1	18.1	18.7	1687.8	8.2	19.0	20.2	1687.8	19.0	89.2
Trial4_440	54	NA()	1.3	11.1905	1.6	3.0080	3.7	0.2442	3.4	0.91	1408.7	10.3	25.7	26.3	1409.6	7.7	17.6	18.1	1411.9	11.3	21.0	22.3	1411.9	21.0	99.8
Trial4_441	93	NA()	1.3	10.7496	1.2	3.1272	3.2	0.2439	3.0	0.93	1407.0	9.6	25.4	26.0	1439.4	6.8	17.4	17.9	1488.4	8.4	19.4	20.7	1488.4	19.4	94.5
Trial4_442	88	NA()	1.6	13.7357	1.6	1.6420	3.7	0.1637	3.3	0.90	977.0	7.8	18.6	19.0	986.5	6.5	14.6	15.1	1008.5	11.5	22.0	23.3	1008.5	22.0	96.9
Trial4_443	94	NA()	1.0	4.9265	0.9	15.6028	2.1	0.5577	1.9	0.90	2857.3	10.4	44.2	45.3	2852.8	6.3	20.8	21.5	2850.3	7.8	17.0	18.1	2850.3	17.0	100.2
Trial4_444	64	NA()	1.9	13.2596	1.8	1.8132	3.9	0.1745	3.5	0.89	1036.6	8.6	19.8	20.2	1050.3	7.2	15.4	15.9	1079.7	12.7	22.5	23.8	1079.7	22.5	96.0
Trial4_445	118	NA()	2.8	9.2726	0.9	4.8191	2.5	0.3242	2.4	0.93	1810.4	10.1	31.0	31.8	1788.2	6.3	18.6	19.2	1763.3	7.1	18.4	19.6	1763.3	18.4	102.7
Trial4_446	27	NA()	1.8	16.7066	3.7	0.9152	4.9	0.1109	3.2	0.65	678.2	7.1	13.9	14.2	659.8	11.9	15.6	15.9	598.4	47.6	51.7	52.3	678.2	13.9	113.3
Trial4_447	99	NA()	2.3	9.9777	1.0	4.0805	2.9	0.2954	2.7	0.93	1668.6	10.3	29.2	29.9	1650.4	6.7	18.2	18.8	1628.2	7.8	18.9	20.2	1628.2	18.9	102.5
Trial4_448	141	NA()	5.3	17.7552	2.2	0.5320	4.2	0.0685	3.6	0.85	427.3	3.5	8.4	8.6	433.1	4.1	8.4	8.6	465.1	17.4	26.9	28.2	427.3	8.4	91.9
Trial4_449	40	NA()	1.4	10.4833	1.8	3.1166	4.0	0.2371	3.6	0.90	1371.5	11.0	25.5	26.0	1436.8	8.5	18.1	18.7	1535.8	12.5	21.5	22.6	1535.8	21.5	89.3
Trial4_450	35	NA()	1.4	8.2961	1.5	5.6414	3.7	0.3396	3.4	0.92	1884.7	12.8	33.0	33.8	1922.4	8.6	19.9	20.5	1964.2	10.9	19.8	20.9	1964.2	19.8	96.0
Trial4_451	90	NA()	1.2	13.5464	1.6	1.7590	3.6	0.1729	3.3	0.90	1028.0	8.1	19.4	19.9	1030.5	6.7	15.0	15.5	1036.5	11.8	22.1	23.4	1036.5	22.1	99.2
Trial4_452	170	NA()	1.5	9.4886	0.9	4.4565	2.1	0.3068	1.9	0.91	1725.1	8.7	29.5	30.2	1722.9	5.5	18.1	18.7	1721.1	6.0	18.1	19.4	1721.1	18.1	100.2
Trial4_453	123	NA()	1.3	19.2450	3.3	0.2511	5.1	0.0351	3.8	0.76	222.2	2.1	4.6	4.7	227.5	3.4	5.4	5.6	283.7	31.7	38.1	39.1	222.2	4.6	NA
Trial4_454	66	NA()	2.5	12.5951	1.7	2.0564	3.8	0.1879	3.4	0.90	1110.2	8.9	21.0	21.4	1134.4	7.3	16.0	16.5	1182.0	12.3	22.1	23.3	1182.0	22.1	93.9
Trial4_455	53	NA()	1.3	11.2361	1.6	2.9735	3.7	0.2424	3.4	0.91	1399.3	10.6	25.7	26.3	1400.9	7.8	17.6	18.2	1404.1	11.3	21.1	22.3	1404.1	21.1	99.7
Trial4_456	199	NA()	1.8	19.4882	2.4	0.3032	4.3	0.0429	3.6	0.83	270.7	1.9	5.3	5.4	268.9	2.8	5.7	5.8	254.9	21.4	30.2	31.5	270.7	5.3	NA
Trial4_457	148	NA()	1.9	10.0761	0.9	4.0074	2.4	0.2930	2.2	0.93	1656.4	8.9	28.6	29.3	1635.7	5.7	17.8	18.4	1610.0	6.5	18.5	19.8	1610.0	18.5	102.9
Trial4_458	49	NA()	1.7	13.8235	2.1	1.7175	4.2	0.1723	3.6	0.87	1024.6	8.8	19.7	20.2	1015.1	7.7	15.4	15.8	995.5	15.2	24.2	25.4	995.5	24.2	102.9
Trial4_459	29	NA()	1.1	8.7871	1.6	5.4823	3.8	0.3495	3.5	0.91	1932.4	13.1	33.7	34.5	1897.8	8.7	19.8	20.5	1861.0	11.6	20.3	21.5	1861.0	20.3	103.8
Trial4_460	16	NA()	0.8	14.1232	3.5	1.6328	4.7	0.1673	3.2	0.67	997.3	10.3	20.0	20.4	982.9	13.4	18.7	19.1	951.8	37.0	41.6	42.3	951.8	41.6	104.8
Trial4_461	25	NA()	1.7	13.3396	2.7	1.8888	4.7	0.1828	3.8	0.81	1082.4	10.3	21.2	21.6	1077.2	10.3	17.2	17.7	1067.6	23.5	30.0	31.0	1067.6	30.0	101.4
Trial4_462	24	NA()	1.5	13.2574	2.8	1.9485	4.7	0.1874	3.9	0.81	1107.5	10.5	21.6	22.1	1097.9	10.4	17.4	17.8	1080.0	23.3	29.8	30.8	1080.0	29.8	102.5
Trial4_463	90	NA()	1.5	11.0797	1.2	3.0266	3.2	0.2433	3.0	0.93	1403.9	10.0	25.5	26.1	1414.3	7.0	17.3	17.9	1430.9	8.8	19.8	21.1	1430.9	19.8	98.1
Trial4_464	99	NA()	0.6	5.1902	0.9	13.8227	2.1	0.5206	1.9	0.90	2701.6	10.9	42.5	43.6	2737.6	6.5	20.7	21.5	2765.1	7.8	17.1	18.2	2765.1	17.1	97.7
Trial4_465	53	NA()	2.2	11.5449	1.6	2.8453	3.8	0.2383	3.4	0.91	1378.1	11.0	25.6	26.1	1367.6	8.1	17.6	18.1	1352.0	11.7	21.4	22.6	1352.0	21.4	101.9
Trial4_466	77	NA()	1.4	12.8791	1.6	2.0204	3.7	0.1888	3.3	0.91	1114.9	8.8	21.0	21.5	1122.4	7.0	15.8	16.3	1137.8	11.3	21.6	22.9	1137.8	21.6	98.0
Trial4_467	157	NA()	1.0	10.9871	0.9	3.0372	2.5	0.2421	2.4	0.93	1397.8	8.2	24.8	25.4	1417.0	5.7	16.9	17.4	1446.9	6.7	18.9	20.2	1446.9	18.9	96.6
Trial4_468	508	NA()	0.5	18.1625	1.3	0.4164	3.0	0.0549	2.6	0.89	344.4	0.4	6.3	6.4	353.5	1.4	6.4	6.6	414.6	10.2	23.1	24.6	344.4	6.3	NA
Trial4_469	121	NA()	1.2	17.4999	2.1	0.6960	4.1	0.0884	3.5	0.86	545.9	4.5	10.7	11.0	536.4	4.8	9.9	10.2	497.0	16.6	26.3	27.6	545.9	10.7	109.8
Trial4_470	127	NA()	2.4	5.4238	0.9	13.0297	1.9	0.5128	1.6	0.87	2668.5	9.8	41.8	42.9	2681.8	5.8	20.4	21.2	2692.6	7.0	16.8	18.0	2692.6	16.8	99.1
Trial4_471	242	NA()	1.1	16.1140	1.3	0.9684	3.0	0.1132	2.7	0.90	691.5	3.8	12.8	13.1	687.6	3.6	11.0	11.4	676.1	9.0	21.8	23.2	691.5	12.8	102.3
Trial4_472	44	NA()	1.5	5.5942	1.0	12.1225	3.1	0.4921	3.0	0.95	2579.6	12.5	41.5	42.5	2613.9	7.5	20.9	21.6	2641.4	9.0	17.8	18.9	2641.4	17.8	97.7
Trial4_473	39	NA()	0.7	10.0825	1.6	4.0214	3.8	0.2942	3.5	0.91	1662.5	12.2	29.8	30.5	1638.5	8.5	18.9	19.5	1608.8	11.9	21.0	22.2	1608.8	21.0	103.3
Trial4_474	44	NA()	1.1	9.8706	1.5	4.0301	3.7	0.2886	3.4	0.91	1634.7	11.9	29.4	30.1	1640.3	8.3	18.8	19.4	1648.3	11.1	20.5	21.7	1648.3	20.5	99.2
Trial4_475	38	NA()	2.3	12.9370	2.3	1.9089	4.4	0.1792	3.7	0.86	1062.6	9.5	20.6	21.0	1084.2	8.8	16.4	16.9	1128.9	17.8	25.6	26.7	1128.9	25.6	94.1
Trial4_476	747	NA()	3.3	9.5966	0.8	3.4466	1.4	0.2400	1.1	0.80	1386.7	5.3	23.8	24.4	1515.1	3.6	16.7	17.4	1700.3	3.2	17.4	18.7	1700.3	17.4	81.6
Trial4_477	396	NA()	1.6	19.9606	2.3	0.1704	4.2	0.0247	3.5	0.83	157.1	0.5	2.9	3.0	159.7	1.4	3.4	3.5	199.6	20.4	29.6	30.9	157.1	2.9	NA
Trial4_478	89	NA()	0.6	13.6737	1.5	1.7820	3.6	0.1768	3.3	0.90	1049.5	8.2	19.8	20.3	1038.9	6.6	15.0	15.5	1017.6	11.2	21.8	23.2	1017.6	21.8	103.1
Trial4_479	42	NA()	2.0	12.9436	2.1	2.0277	4.2	0.1904	3.7	0.87	1123.7	9.7	21.5	22.0	1124.8	8.2	16.3	16.8	1127.9	14.9	23.7	24.9	1127.9	23.7	99.6
Trial4_480	174	NA()	1.5	18.7080	2.6	0.3043	4.5	0.0413	3.7	0.82	260.9	2.1	5.2	5.3	269.7	3.1	5.8	6.0	348.1	23.2	31.3	32.4	260.9	5.2	NA
Trial4_481	209	NA()	0.9	20.2253	3.0	0.1930	4.8	0.0283	3.7	0.78	180.0	1.3	3.6	3.6	179.2	2.5	4.2	4.3	168.9	30.8	37.6	38.7	180.0	3.6	NA
Trial4_482	155	NA()	3.2	12.4774	1.1	2.2565	2.8	0.2043	2.6	0.92	1198.4	7.5	21.7	22.2	1198.8	5.6	15.6	16.2	1200.6	7.5	19.7	21.1	1200.6	19.7	99.8
Trial4_483	49	NA()	3.1	12.9567	2.0	1.7928	4.2	0.1686	3.6	0.87	1004.1	9.0	19.5	19.9	1042.9	8.2	15.8	16.3	1125.8	16.0	24.5	25.6	1125.8	24.5	89.2
Trial4_484	107	NA()	1.2	20.1830	3.6	0.2632	5.3	0.0385	3.9	0.73	243.8	2.4	5.1	5.2	237.2	4.4	6.2	6.4	173.8	42.9	48.1	48.9	243.8	5.1	NA
Trial4_485	70	NA()	0.6	5.4831	0.9	12.9018	2.6	0.5133	2.4	0.93	2670.7	11.9	42.4	43.5	2672.5	7.0	20.8	21.5	2674.6	8.5	17.5	18.6	2674.6	17.5	99.9
Trial4_486	145	NA()	1.3	18.9999	2.9	0.2974	4.8	0.0410	3.8	0.79	259.0	2.3	5.2	5.4	264.3	3.7	6.1	6.2	313.0	29.8	36.5	37.6	259.0	5.2	NA
Trial4_487	49	NA()	2.2	9.0847	1.3	4.7553	3.5	0.3135	3.3	0.93	1757.7	12.3	31.2	31.9	1777.0	8.1	19.2	19.8	1800.6	9.8	19.5	20.7	1800.6	19.5	97.6
Trial4_488	309	NA()	3.1	12.6561	0.8	2.0939	1.9	0.1923	1.7	0.91	1133.7	4.4	19.8	20.4	1146.8	3.4	14.7	15.2	1172.5	5.3	19.1	20.5	1172.5	19.1	96.7
Trial4_489	105	NA()	0.7	5.4122	0.9	13.2717	2.1	0.5212	1.8	0.89	2704.2	10.7	42.5	43.6	2699.2	6.4	20.6	21.4	2696.1	7.7	17.1	18.3	2696.1	17.1	100.3
Trial4_490	120	NA()	0.9	11.2991	1.1	2.7991	3.0	0.2295	2.8	0.93	1331.8	9.1	24.1	24.7	1355.3	6.4	16.8	17.4	1393.4	7.8	19.4	20.8	1393.4	19.4	95.6
Trial4_491	305	NA()	0.7	19.7501	2.0	0.2838	3.9	0.0407	3.3	0.86	257.0	1.3	4.9	5.0	253.7	2.0	5.1	5.3	224.1	16.9	27.3	28.7	257.0	4.9	NA
Trial4_492	153	NA()	1.4	13.8859	1.2	1.7244	3.0	0.1737	2.8	0.92	1032.7	7.1	19.1	19.6	1017.7	5.5	14.4	14.9	986.4	8.6	20.7	22.1	986.4	20.7	104.7
Trial4_493	106	NA()	3.1	13.6173	1.5	1.6637	3.5	0.1644	3.2	0.91	981.1	7.7	18.6	19.0	994.8	6.4	14.7	15.1	1026.0	11.1	21.8	23.1	1026.0	21.8	95.6
Trial4_494	50	NA()	1.3	13.7401	2.1	1.7543	4.2	0.1749	3.6	0.87	1039.0	9.2	20.1	20.5	1028.7	8.5	15.9	16.3	1007.8	18.1	26.1	27.2	1007.8	26.1	103.1
Trial4_495	75	NA()	1.0	10.0336	1.2	3.9785	3.2	0.2896	3.0	0.93	1639.8	11.1	29.1	29.8	1629.8	7.4	18.4	19.0	1617.8	9.0	19.4	20.7	1617.8	19.4	101.4
Trial4_496	202	NA()	7.4	13.4751	1.1	1.7714	2.7	0.1732	2.5	0.92	1029.7	6.0	18.7	19.2	1035.0	4.7	14.3	14.8	1047.2	7.3	20.1	21.5	1047.2	20.1	98.3
Trial4_497	32	NA()	1.3	11.8821	2.1	2.4842	4.3	0.2142	3.7	0.87	1251.0	10.7	23.7	24.2	1267.4	9.1	17.6	18.1	1296.2	16.2	24.2	25.4	1296.2	24.2	96.5
Trial4_498	131	NA()	2.5	12.0545	1.1	2.4442	2.9	0.2138	2.7	0.93	1248.9	8.2	22.6	23.2	1255.7	5.9	16.1	16.6	1268.2	7.8	19.7	21.0	1268.2	19.7	98.5
Trial4_499	103	NA()	1.3	8.5886	0.9	5.3681	2.7	0.3345	2.5	0.94	1860.3	10.6	31.9	32.7	1879.8	6.6	19.0	19.6	1902.2	7.2	18.1	19.4	1902.2	18.1	97.8
Trial4_500	126	NA()	2.4	13.8412	1.3	1.7347	3.2	0.1742	3.0	0.91	1035.3	7.3	19.2	19.7	1021.5	5.8	14.6	15.1	992.9	9.5	21.1	22.5	992.9	21.1	104.3
Trial4_501	124	NA()	1.1	19.9173	3.2	0.2798	5.0	0.0404	3.8	0.77	255.5	2.3	5.2	5.3	250.5	3.7	5.9	6.1	204.6	32.3	38.8	39.8	255.5	5.2	NA
Trial4_502	61	NA()	2.8	13.9928	1.9	1.6866	4.0	0.1712	3.5	0.88	1019.0	8.6	19.5	20.0	1003.5	7.4	15.2	15.6	970.8	14.3	23.7	24.9	970.8	23.7	105.0
Trial4_503	118	NA()	0.9	17.1544	2.2	0.6029	4.2	0.0750	3.6	0.85	466.5	4.0	9.3	9.5	479.1	4.8	9.3	9.5	540.8	19.3	27.9	29.2	466.5	9.3	86.2
Trial4_504	67	NA()	1.6	13.0451	1.7	1.9630	3.8	0.1858	3.4	0.90	1098.6	8.9	20.8	21.3	1102.9	7.3	15.8	16.2	1112.3	12.5	22.3	23.6	1112.3	22.3	98.8
Trial4_505	34	NA()	2.1	10.3198	1.8	3.5290	4.0	0.2643	3.6	0.90	1511.6	11.8	27.7	28.3	1533.7	8.9	18.7	19.3	1565.3	13.2	21.8	22.9	1565.3	21.8	96.6
Trial4_506	43	NA()	2.7	13.6261	2.2	1.7269	4.3	0.1707	3.7	0.86	1016.2	9.0	19.7	20.1	1018.6	8.1	15.6	16.1	1024.7	16.7	25.2	26.3	1024.7	25.2	99.2
Trial4_507	111	NA()	1.7	9.5392	1.0	4.2785	2.7	0.2961	2.5	0.94	1672.2	9.9	29.1	29.9	1689.2	6.4	18.3	18.9	1711.3	7.2	18.5	19.8	1711.3	18.5	97.7
Trial4_508	76	NA()	1.5	5.5318	0.9	11.3278	2.6	0.4547	2.5	0.94	2416.0	11.5	39.2	40.2	2550.5	6.8	20.6	21.3	2660.0	7.5	17.1	18.2	2660.0	17.1	90.8
Trial4_509	145	NA()	1.1	9.9160	0.9	3.7575	2.5	0.2704	2.3	0.93	1542.6	8.7	26.9	27.6	1583.7	5.8	17.6	18.3	1639.7	6.4	18.4	19.7	1639.7	18.4	94.1
Trial4_510	33	NA()	1.0	10.8377	1.9	3.2245	4.1	0.2536	3.6	0.89	1456.8	11.5	26.8	27.4	1463.1	8.7	18.3	18.9	1472.9	13.4	22.1	23.3	1472.9	22.1	98.9
Trial4_511	56	NA()	1.7	9.1433	1.3	4.6899	3.4	0.3111	3.2	0.93	1746.3	11.4	30.7	31.4	1765.4	7.5	19.0	19.6	1788.9	9.1	19.2	20.4	1788.9	19.2	97.6

Trial4_512	279	NA()	1.4	19.4552	2.1	0.2809	4.0	0.0397	3.4	0.85	250.7	1.3	4.8	4.9	251.4	2.2	5.1	5.3	258.8	18.9	28.5	29.8	250.7	4.8	NA
Trial4_513	69	NA()	0.7	5.3265	0.9	13.2850	2.6	0.5135	2.4	0.93	2671.4	11.3	42.2	43.3	2700.1	6.8	20.8	21.5	2722.4	8.2	17.3	18.4	2722.4	17.3	98.1
Trial4_514	55	NA()	1.4	9.2542	1.3	4.6119	3.4	0.3097	3.2	0.93	1739.1	11.6	30.6	31.3	1751.4	7.7	19.0	19.6	1766.9	9.5	19.4	20.6	1766.9	19.4	98.4
Trial4_515	74	NA()	2.8	12.8915	1.6	2.0795	3.7	0.1945	3.3	0.90	1145.8	8.9	21.4	21.9	1142.1	7.0	15.9	16.4	1135.9	11.6	21.8	23.1	1135.9	21.8	100.9
Trial4_516	205	NA()	3.2	13.6988	1.0	1.7714	2.6	0.1761	2.4	0.92	1045.5	5.7	18.8	19.3	1035.0	4.5	14.2	14.7	1013.9	7.2	20.1	21.5	1013.9	20.1	103.1
Trial4_517	562	NA()	17.2	17.2861	1.0	0.7039	2.1	0.0883	1.9	0.89	545.4	0.8	9.8	10.0	541.1	1.5	8.8	9.2	524.1	7.0	21.5	23.0	545.4	9.8	104.1
Trial4_518	329	NA()	3.6	13.7629	0.9	1.6790	2.0	0.1677	1.8	0.90	999.3	3.1	17.5	18.0	1000.6	2.8	13.5	14.0	1004.4	5.6	19.6	21.1	1004.4	19.6	99.5
Trial4_519	110	NA()	2.0	8.9592	0.9	5.1835	2.5	0.3370	2.4	0.94	1872.1	10.2	31.9	32.7	1849.9	6.3	18.8	19.4	1825.9	7.1	18.3	19.5	1825.9	18.3	102.5
Trial4_520	59	NA()	1.3	11.1433	1.4	3.1255	3.6	0.2527	3.3	0.92	1452.4	10.5	26.4	27.0	1439.0	7.5	17.7	18.2	1420.0	10.3	20.5	21.8	1420.0	20.5	102.3
Trial4_521	57	NA()	2.5	13.3574	1.8	1.8569	4.0	0.1800	3.5	0.89	1066.8	8.8	20.3	20.8	1065.9	7.4	15.6	16.0	1064.9	13.3	22.9	24.2	1064.9	22.9	100.2
Trial4_522	83	NA()	1.8	13.5029	1.6	1.8838	3.6	0.1846	3.3	0.90	1091.9	8.4	20.5	21.0	1075.4	6.8	15.4	15.9	1043.0	12.0	22.2	23.5	1043.0	22.2	104.7
Trial4_523	65	NA()	1.3	11.1421	1.4	3.1776	3.5	0.2569	3.2	0.92	1473.9	10.5	26.7	27.3	1451.7	7.3	17.7	18.2	1420.2	9.9	20.3	21.6	1420.2	20.3	103.8
Trial4_524	296	NA()	3.7	14.5385	0.8	1.4065	2.3	0.1484	2.2	0.94	891.8	3.7	16.1	16.5	891.7	3.3	12.3	12.8	892.2	6.7	15.7	17.5	891.8	16.1	100.0
Trial4_525	125	NA()	5.3	8.7205	0.7	5.2769	2.4	0.3339	2.3	0.96	1857.3	9.9	31.9	32.7	1865.1	6.2	18.2	18.8	1874.7	6.8	14.2	15.7	1874.7	14.2	99.1
Trial4_526	96	NA()	1.6	12.7904	1.1	1.9536	3.4	0.1813	3.2	0.94	1074.1	8.1	20.3	20.8	1099.7	6.5	14.9	15.4	1151.5	10.3	17.1	18.7	1151.5	17.1	93.3
Trial4_527	44	NA()	2.8	13.4546	1.7	1.7789	4.1	0.1737	3.7	0.91	1032.3	9.1	20.1	20.5	1037.8	8.1	15.3	15.8	1050.3	16.1	21.3	22.6	1050.3	21.3	98.3
Trial4_528	60	NA()	0.5	5.5245	1.0	12.5758	2.8	0.5041	2.6	0.94	2631.4	11.9	42.3	43.4	2648.4	7.1	20.1	20.8	2662.1	8.5	14.2	15.6	2662.1	14.2	98.8
Trial4_529	260	NA()	2.6	13.0690	0.7	2.0200	2.2	0.1916	2.0	0.95	1129.8	5.1	20.1	20.6	1122.3	3.9	14.1	14.7	1108.6	6.0	15.0	16.8	1108.6	15.0	101.9
Trial4_530	98	NA()	2.6	13.3793	1.2	1.7988	3.4	0.1746	3.2	0.94	1037.6	7.9	19.7	20.1	1045.0	6.3	14.5	15.0	1061.6	10.4	17.3	18.9	1061.6	17.3	97.7
Trial4_531	152	NA()	0.3	8.5177	0.9	5.8269	2.1	0.3601	1.9	0.91	1982.8	9.5	33.4	34.3	1950.4	5.7	18.2	18.9	1917.0	6.3	13.9	15.5	1917.0	13.9	103.4
Trial4_532	31	NA()	1.7	5.3577	0.9	12.8332	3.4	0.4989	3.3	0.96	2609.0	13.4	42.5	43.5	2667.5	8.1	20.5	21.2	2712.8	9.7	14.9	16.2	2712.8	14.9	96.2
Trial4_533	67	NA()	1.1	5.2198	0.9	14.1692	2.7	0.5367	2.5	0.93	2769.4	11.7	43.9	45.0	2761.1	6.9	20.2	20.9	2755.7	8.3	14.0	15.4	2755.7	14.0	100.5
Trial4_534	500	NA()	0.9	19.7973	1.3	0.2982	3.2	0.0428	3.0	0.92	270.4	0.4	5.0	5.1	265.0	1.4	4.9	5.1	218.6	13.3	20.7	22.5	270.4	5.0	NA
Trial4_535	178	NA()	2.2	9.1406	0.9	4.5451	2.2	0.3014	2.0	0.92	1698.5	8.1	29.2	29.9	1739.3	5.2	17.4	18.1	1789.5	5.6	13.7	15.4	1789.5	13.7	94.9
Trial4_536	131	NA()	1.8	16.9156	1.5	0.8541	3.7	0.1048	3.4	0.92	642.7	4.8	12.5	12.7	626.9	4.9	10.6	10.9	571.4	15.0	21.2	22.7	642.7	12.5	112.5
Trial4_537	29	NA()	0.7	5.5351	0.9	13.1603	3.4	0.5285	3.3	0.96	2735.3	13.0	43.9	44.9	2691.2	7.8	20.4	21.2	2659.0	9.6	14.9	16.2	2659.0	14.9	102.9
Trial4_538	139	NA()	1.6	16.8257	1.4	0.8787	3.6	0.1073	3.4	0.92	656.9	4.7	12.6	12.9	640.3	4.6	10.6	10.9	583.0	13.4	20.1	21.7	656.9	12.6	112.7
Trial4_539	57	NA()	0.4	5.4351	0.9	13.6997	2.8	0.5403	2.7	0.94	2784.6	11.5	44.0	45.1	2729.2	6.8	20.1	20.8	2689.1	8.2	14.0	15.4	2689.1	14.0	103.5
Trial4_540	349	NA()	2.7	13.4570	0.6	1.7814	1.9	0.1739	1.7	0.94	1033.8	2.3	18.1	18.6	1038.7	2.3	13.2	13.8	1049.9	5.3	14.9	16.7	1049.9	14.9	98.5
Trial4_541	170	NA()	1.8	19.4911	2.2	0.2999	4.3	0.0424	3.7	0.86	267.8	1.9	5.3	5.4	266.3	3.4	5.8	5.9	254.6	28.8	32.8	34.0	267.8	5.3	NA
Trial4_542	51	NA()	4.3	6.2767	0.8	10.3125	3.1	0.4697	3.0	0.96	2482.1	11.9	40.5	41.5	2463.2	7.0	19.8	20.5	2448.4	8.3	14.3	15.7	2448.4	14.3	101.4
Trial4_543	87	NA()	0.9	11.8424	1.0	2.7425	3.3	0.2357	3.1	0.95	1364.1	8.9	24.7	25.3	1340.1	6.4	16.2	16.8	1302.7	9.0	16.1	17.7	1302.7	16.1	104.7
Trial4_544	101	NA()	2.2	13.3362	1.1	1.9292	3.4	0.1867	3.2	0.94	1103.4	7.5	20.5	21.0	1091.2	6.0	14.7	15.2	1068.1	10.2	17.2	18.8	1068.1	17.2	103.3
Trial4_545	221	NA()	1.7	13.4744	0.8	1.7854	2.5	0.1746	2.4	0.95	1037.1	4.8	18.6	19.1	1040.1	3.9	13.6	14.1	1047.3	6.7	15.4	17.2	1047.3	15.4	99.0
Trial4_546	158	NA()	1.8	14.2363	1.0	1.5176	3.1	0.1568	2.9	0.95	938.8	5.6	17.3	17.8	937.5	4.8	13.1	13.6	935.5	9.1	16.8	18.5	935.5	16.8	100.4
Trial4_547	88	NA()	0.8	9.1100	0.8	4.9579	2.8	0.3277	2.7	0.96	1827.3	10.0	31.5	32.3	1812.2	6.4	18.1	18.7	1795.6	7.5	14.6	16.2	1795.6	14.6	101.8
Trial4_548	68	NA()	0.9	18.7920	2.8	0.4493	4.8	0.0613	3.9	0.81	383.3	3.6	7.9	8.0	376.8	6.0	8.7	8.9	337.9	37.5	40.6	41.5	383.3	7.9	NA
Trial4_549	124	NA()	1.2	9.1827	0.7	4.6938	2.5	0.3127	2.4	0.96	1754.2	9.1	30.3	31.0	1766.1	5.8	17.7	18.4	1781.1	6.6	14.2	15.8	1781.1	14.2	98.5
Trial4_550	74	NA()	1.0	5.2437	0.9	14.2598	2.5	0.5426	2.3	0.93	2794.1	10.5	43.9	45.0	2767.1	6.3	20.0	20.8	2748.2	7.8	13.8	15.1	2748.2	13.8	101.7
Trial4_551	68	NA()	6.0	9.4311	0.9	4.6091	3.2	0.3154	3.0	0.96	1767.3	10.6	30.9	31.7	1750.9	6.9	18.1	18.7	1732.3	8.4	15.2	16.7	1732.3	15.2	102.0
Trial4_552	78	NA()	6.4	9.3547	0.9	4.6735	3.0	0.3172	2.9	0.96	1776.2	10.9	31.1	31.9	1762.5	7.0	18.1	18.8	1747.2	8.2	15.0	16.6	1747.2	15.0	101.7
Trial4_553	87	NA()	0.8	11.1757	1.0	2.8652	3.3	0.2323	3.1	0.95	1346.7	9.5	24.8	25.3	1372.8	6.9	16.6	17.1	1414.4	9.0	16.0	17.5	1414.4	16.0	95.2
Trial4_554	449	NA()	2.1	18.1966	1.0	0.5119	2.8	0.0676	2.6	0.93	421.6	0.7	7.7	7.9	419.7	1.6	7.1	7.3	410.4	9.6	18.1	20.0	421.6	7.7	102.7
Trial4_555	169	NA()	2.0	13.2680	0.9	1.8389	2.9	0.1770	2.7	0.95	1050.8	6.6	19.4	19.8	1059.5	5.1	14.1	14.6	1078.4	7.8	15.9	17.6	1078.4	15.9	97.4
Trial4_556	154	NA()	2.9	5.5307	0.9	13.2255	1.6	0.5307	1.3	0.83	2744.6	9.0	43.0	44.1	2695.9	5.4	19.6	20.4	2660.3	6.6	13.2	14.6	2660.3	13.2	103.2
Trial4_557	168	NA()	1.5	16.2149	1.2	0.9384	3.4	0.1104	3.2	0.93	675.1	4.7	12.9	13.2	672.1	4.5	10.8	11.2	662.7	11.8	18.9	20.6	675.1	12.9	101.9
Trial4_558	33	NA()	1.4	13.5863	2.1	1.7110	4.4	0.1687	3.8	0.88	1004.8	9.2	19.7	20.2	1012.7	9.8	16.1	16.5	1030.6	23.4	27.3	28.3	1030.6	27.3	97.5
Trial4_559	14	NA()	1.2	7.9140	1.7	6.4848	4.2	0.3724	3.8	0.92	2040.6	14.6	36.0	36.8	2043.8	10.7	20.6	21.2	2047.9	15.5	19.7	20.8	2047.9	19.7	99.6
Trial4_560	49	NA()	1.9	13.2734	1.6	1.8887	4.0	0.1819	3.7	0.92	1077.3	9.1	20.8	21.2	1077.1	8.0	15.5	16.0	1077.6	15.3	20.6	22.0	1077.6	20.6	100.0
Trial4_561	96	NA()	0.9	13.0226	1.1	2.0186	3.4	0.1907	3.2	0.95	1125.4	8.6	21.2	21.7	1121.8	6.6	15.1	15.6	1115.7	10.0	17.0	18.6	1115.7	17.0	100.9
Trial4_562	87	NA()	0.5	5.5086	0.9	12.9463	2.4	0.5175	2.2	0.92	2688.4	11.2	42.8	43.9	2675.7	6.7	20.0	20.7	2666.9	8.1	14.0	15.3	2666.9	14.0	100.8
Trial4_563	50	NA()	1.8	13.8922	1.7	1.5547	4.1	0.1567	3.7	0.91	938.5	8.8	18.6	19.0	952.4	8.0	14.7	15.2	985.5	16.5	21.6	23.0	985.5	21.6	95.2
Trial4_564	62	NA()	1.6	13.9225	1.5	1.6358	3.9	0.1653	3.6	0.92	985.9	8.8	19.3	19.7	984.1	7.6	14.7	15.2	981.0	14.8	20.3	21.8	981.0	20.3	100.5
Trial4_565	204	NA()	2.0	13.5136	0.8	1.7288	2.6	0.1695	2.5	0.95	1009.5	6.2	18.6	19.1	1019.3	4.9	13.8	14.3	1041.4	7.3	15.7	17.5	1041.4	15.7	96.9
Trial4_566	51	NA()	2.1	6.4929	0.9	8.6199	3.2	0.4061	3.1	0.96	2197.1	13.1	37.4	38.3	2298.7	7.9	19.9	20.6	2391.0	8.8	14.6	16.0	2391.0	14.6	91.9
Trial4_567	20	NA()	1.9	13.3849	2.4	1.8954	4.6	0.1841	3.9	0.85	1089.2	10.8	21.7	22.1	1079.5	11.4	17.5	18.0	1060.8	27.0	30.3	31.3	1060.8	30.3	102.7
Trial4_568	72	NA()	1.9	13.3790	1.3	1.8268	3.7	0.1773	3.5	0.94	1052.4	9.1	20.4	20.8	1055.1	7.3	15.0	15.5	1061.6	12.2	18.5	20.0	1061.6	18.5	99.1
Trial4_569	132	NA()	3.9	16.8045	1.5	0.7721	3.8	0.0941	3.5	0.92	580.0	5.0	11.5	11.8	581.0	5.0	10.2	10.5	585.7	14.7	21.0	22.6	580.0	11.5	99.0
Trial4_570	60	NA()	1.1	16.3067	2.0	0.8671	4.3	0.1026	3.8	0.88	629.6	6.3	12.9	13.2	634.0	6.8	11.6	12.0	650.6	21.1	25.7	27.0	629.6	12.9	96.8
Trial4_571	33	NA()	2.4	13.7665	2.0	1.6865	4.3	0.1685	3.8	0.89	1003.6	9.9	20.1	20.5	1003.5	9.5	15.9	16.4	1003.9	21.4	25.6	26.7	1003.9	25.6	100.0
Trial4_572	41	NA()	0.8	8.6215	1.1	5.4684	3.6	0.3421	3.4	0.95	1896.7	13.6	33.7	34.5	1895.7	8.7	19.2	19.9	1895.3	10.5	16.2	17.6	1895.3	16.2	100.1
Trial4_573	75	NA()	0.5	19.2006	3.7	0.2233	4.9	0.0311	3.2	0.65	197.5	2.3	4.3	4.4	204.6	4.7	6.0	6.1	289.0	51.1	53.4	54.1	197.5	4.3	NA
Trial4_574	23	NA()	1.4	13.3073	2.3	1.9281	4.5	0.1862	3.9	0.87	1100.6	10.9	21.9	22.4	1090.9	10.8	17.2	17.6	1072.5	24.1	27.8	28.8	1072.5	27.8	102.6
Trial4_575	109	NA()	3.3	13.7702	1.1	1.7097	3.4	0.1708	3.2	0.94	1016.7	8.3	19.5	20.0	1012.2	6.5	14.4	14.9	1003.4	10.4	17.4	19.0	1003.4	17.4	101.3
Trial4_576	123	NA()	1.0	13.3075	1.0	1.9023	3.2	0.1837	3.0	0.95	1087.0	8.6	20.6	21.1	1081.9	6.5	14.8	15.3	1072.4	9.5	16.8	18.4	1072.4	16.8	101.4
Trial4_577	321	NA()	2.0	19.4517	1.6	0.2615	3.8	0.0369	3.4	0.90	233.6	1.4	4.5	4.6	235.8	2.0	4.7	4.8	259.2	17.0	23.3	24.8	233.6	4.5	NA
Trial4_578	343	NA()	1.3	12.4978	0.8	2.1263	1.8	0.1928	1.6	0.89	1136.6	4.6	20.1	20.6	1157.4	3.5	14.3	14.8	1197.4	5.0	14.5	16.3	1197.4	14.5	94.9
Trial4_579	45	NA()	1.6	13.2809	1.7	1.7423	4.1	0.1679	3.7	0.91	1000.5	9.8	20.0	20.4	1024.3	8.8	15.6	16.1	1076.4	17.3	22.2	23.4	1076.4	22.2	92.9
Trial4_580	59	NA()	4.1	13.1278	1.5	1.7565	3.9	0.1673	3.6	0.92	997.3	9.4	19.7	20.2	1029.5	8.1	15.2	15.7	1099.7	14.3	19.9	21.3	1099.7	19.9	90.7
Trial4_581	264	NA()	2.5	11.9686	0.6	2.4310	2.0	0.2111	1.9	0.95	1234.8	6.3	22.0	22.6	1251.8	4.6	15.1	15.7	1282.1	5.6	14.5	16.3	1282.1	14.5	96.3
Trial4_582	66	NA()	1.5	13.8940	1.4	1.6367	3.8	0.1650	3.6	0.93	984.5	9.1	19.4	19.9	984.5	7.7	14.8	15.2	985.2	14.1	19.8	21.3	985.2	19.8	99.9
Trial4_583	429	NA()	37.8	5.0565	0.8	13.6306	1.3	0.5001	1.0	0.78	2614.3	3.7	40.5	41.6	2724.4	2.2	19.1	19.8	2807.8	2.6	11.5	13.1	2807.8	11.5	93.1
Trial4_584	62	NA()	2.3	9.1811	1.0	4.4147	3.4	0.2941	3.2	0.95	1662.0	12.5	30.2	30.9	1715.1	8.3	18.5	19.2	1781.4	9.6	15.8	17.3	1781.4	15.8	93.3
Trial4_585	234	NA()	2.9	12.7964	0.7	2.0396	2.3	0.1894	2.2	0.95	1118.0	6.6	20.4	20.9	1128.8	4.9	14.5	15.0	1150.6	6.5	15.1	16.9	1150.6	15.1	97.2
Trial4_586	65	NA()	1.3	12.2410	1.4	1.9510	3.8	0.1733	3.6	0.93	1030.2	9.8	20.4	20.8	1098.8	8.2	15.7	16.2	1238.2	12.8	18.6	20.0	1238.2	18.6	83.2
Trial4_587	119	NA()	2.8	12.9678	1.0	2.0360	3.2	0.1916	3.0	0.95	1129.9	9.2	21.5	22.0	1127.6	6.8	15.2	15.8	1124.1	9.3	16.5	18.2	1124.1	16.5	100.5
Trial4_588	205	NA()	1.2	11.1006	0.7	2.9963	2.2	0.2413	2.1	0.95	1393.6	8.5	25.0	25.6	1406.7	5.7	16.3	16.9	1427.3	6.3	14.6	16.3	1427.3	14.6	97.6
Trial4_589	497	NA()	18.6	12.9578	0.8	1.7920	1.5	0.1685	1.2	0.83	1003.8	1.7	17.5	18.0	1042.5	1.9	13.2	13.7	1125.7	4.3	14.4	16.3	1125.7	14.4	89.2
Trial4_590	132	NA()	1.4	10.6045	0.8	3.3138	2.7	0.2550	2.6	0.96	1464.1	10.4	26.7	27.3	1484.3	6.9	17.1	17.7	1514.1	7.7	15.1	16.7	1514.1	15.1	96.7
Trial4_591	186	NA()	1.8	9.6024	0.9	4.1965	2.1	0.2924	1.9	0.91	1653.4	9.6	29.0	29.7	1673.3	6.1	17.5	18.1	1699.2	6.3	14.1	15.8	1699.2	14.1	97.3
Trial4_592	63	NA()	1.9	13.0294	1.4	1.9659	3.8	0.1859	3.5	0.93	1098.9	10.1	21.5	22.0	1103.9	8.2	15.8	16.3	1114.7	13.8	19.5	20.9	1114.7	19.5	98.6
Trial4_593	300	NA()	4.2	19.2534	1.7	0.2582	3.9	0.0361	3.5	0.90	228.4	1.5	4.5	4.6	233.2	2.3	4.7	4.9	282.7	19.7	25.2	26.7	228.4	4.5	NA
Trial4_594	238	NA()	0.8	19.3482	1.8	0.2893	4.0	0.0406	3.6	0.89	256.7	2.0	5.1	5.3	258.0	2.6	5.3	5.4	271.4	19.2	24.8	26.3	256.7	5.1	NA
Trial4_595	172	NA()	1.5	13.0203	0.9	1.9326	2.8	0.1826	2.6	0.95	1081.1	7.6	20.2	20.7	1092.5	5.7	14.6	15.1	1116.1	7.7	15.7	17.5	1116.1	15.7	96.9
Trial4_596	32	NA()	2.9	14.0190	2.2	1.4282	4.5	0.1453	3.9	0.87	874.5	9.2	17.9	18.2	900.8	9.4	15.2	15.6	966.9	22.5	26.5	27.6	966.9	26.5	90.4
Trial4_597	23	NA()	0.9	13.8999	2.4	1.5860	4.6	0.1600	4.0	0.85	956.5	10.1	19.5	19.9	964.7	10.4	16.2	16.6	984.3	25.0	28.6	29.7	984.3	28.6	97.2
Trial4_598	115	NA()	0.8	5.3177	0.9	13.2483	2.0	0.5112	1.8	0.89	2661.7	11.4	42.5	43.6	2697.5	6.7	20.0	20.8	2725.2	7.9	13.8	15.2	2725.2	13.8	97.7
Trial4_599	89	NA()	1.3	12.4415	1.1	2.1605	3.4	0.1950	3.2	0.95	1148.6	9.4	21.9	22.4	1168.4	7.2	15.7	16.2	1206.2	10.4	17.1	18.6	1206.2	17.1	95.2
Trial4_600	48	NA()	1.4	13.2985	1.6	1.9211	4.0	0.1854	3.7	0.92	1096.3	10.1	21.5	21.9	1088.4	8.5	15.8	16.3	1073.8	15.7	20.9	22.3	1073.8	20.9	102.1
Trial4_601	76	NA()	2.6	13.4389	1.3	1.8749	3.7	0.1828	3.4	0.94	1082.4	9.4	20.9	21.4	1072.3	7.4	15.2	15.7	1052.7	12.3	18.5	20.1	1052.7	18.5	102.8
Trial4_602	13	NA()	0.6	10.9846	2.5	2.8642	4.1	0.2283	3.2	0.78	1325.5	13.1	26.1	26.6	1372.6	13.3	20.1	20.6	1447.4	26.4	29.4	30.3	1447.4	29.4	91.6
Trial4_603	68	NA()	1.1	11.4559	1.1	2.8163	3.5	0.2341	3.3	0.95	1356.0	10.9	25.4	26.0	1359.9	7.8	16.9	17.5	1366.9	10.3	16.8	18.3	1366.9	16.8	99.2
Trial4_604	46	NA()	1.2	12.1150	1.4	2.4462	3.9	0.2150	3.6	0.93	1255.6	11.0	24.1	24.6	1256.3	8.5	16.7	17.2	1258.4	13.2	18.8	20.2	1258.4	18.8	99.8
Trial4_605	474	NA()	1.0	18.5894	1.4	0.2590	3.4	0.0349	3.2	0.92	221.4	0.5	4.1	4.2	233.9	1.4	4.4	4.6	362.4	14.2	21.0	22.7	221.4	4.1	NA
Trial4_606	266	NA()	8.1	12.9970	0.7	1.7596	2.3	0.1659	2.2	0.95	989.7	5.0	17.9	18.4	1030.7	4.0	13.6	14.1	1119.7	6.3	15.1	16.9	1119.7	15.1	88.4
Trial4_607	333	NA()	0.8	11.2016	0.8	2.8421	1.6	0.2310	1.4	0.86	1339.7	4.5	23.2	23.8	1366.7	3.3	15.4	16.0	1410.0	4.4	13.9	15.7	1410.0	13.9	95.0
Trial4_608	47	NA()	1.2	19.8103	4.3	0.2791	5.4	0.0401	3.2	0.60	253.5	3.0	5.5	5.7	249.9	7.4	8.6	8.7	217.1	72.4	74.1	74.6	253.5	5.5	NA
Trial4_609	18	NA()	1.0	7.7985	1.4	6.7464	4.0	0.3818	3.7	0.93	2084.5	14.7	36.6	37.4	2078.7	9.9	20.3	20.9	2073.8	13.1	17.9	19.1	2073.8	17.9	100.5
Trial4_610	201	NA()	1.6	13.2548	0.8	1.9383	2.6	0.1864	2.4	0.95	1101.9	6.5	20.1	20.6	1094.4	4.9	14.3	14.8	1080.4	7.0	15.5	17.3	1080.4	15.5	102.0
Trial4_611	271	NA()	3.0	13.8466	0.7	1.6649	2.3	0.1673	2.2	0.95	997.1	4.8	18.0	18.5	995.2	3.9	13.3	13.8	992.2	6.5	15.4	17.2	992.2	15.4	100.5
Trial4_612	290	NA()	1.4	8.3939	0.8	4.9970	1.5	0.3043	1.2	0.83	1712.8	6.1	28.9	29.7	1818.8	3.9	17.4	18.1	1943.3	4.1	13.0	14.7	1943.3	13.0	88.1
Trial4_613	63	NA()	1.0	5.3690	1.0	12.9863	2.8	0.5059	2.6	0.94	2639.2	12.2	42.5	43.5	2678.7	7.2	20.2	20.9	2709.4	8.6	14.2	15.5	2709.4	14.2	97.4
Trial4_614	46	NA()	0.7	10.7843	1.2	3.3804	3.7	0.2645	3.5	0.94	1512.9	11.7	27.9	28.5	1499.9	8.3	17.8	18.3	1482.3	11.5	17.4	18.9	1482.3	17.4	102.1
Trial4_615	13	NA()	1.3	13.3936	3.5	1.8316	4.8	0.1780	3.2	0.67	1056.0	11.0	21.3	21.8	1056.8	14.2	19.4	19.7	1059.5	37.2	39.7	40.4	1059.5	39.7	99.7
Trial4_616	34	NA()	3.2	12.7394	1.8	2.0536	4.2	0.1898	3.8	0.90	1120.5	10.3	21.9	22.4	1133.5	9.1	16.4	16.9	1159.4	17.5	22.2	23.4	1159.4	22.2	96.6
Trial4_617	255	NA()	2.7	13.6971	0.8	1.7380	2.3	0.1727	2.2	0.95	1027.1	5.3	18.6	19.1	1022.7	4.2	13.6	14.1	1014.2	6.5	15.4	17.2	1014.2	15.4	101.3
Trial4_618	65	NA()	1.9	12.8460	1.4	1.9652	3.8	0.1832	3.5	0.93	1084.3	9.3	20.9	21.4	1103.7	7.6	15.5	16.0	1142.9	12.7	18.7	20.2	1142.9	18.7	94.9
Trial4_619	63	NA()	0.8	16.8719	2.0	0.8288	4.3	0.1015	3.8	0.88	623.0	6.2	12.7	13.0	613.0	6.6	11.3	11.7	577.1	21.7	26.3	27.6	623.0	12.7	108.0
Trial4_620	97	NA()	0.9	18.8500	2.5	0.4360	4.5	0.0596	3.8	0.84	373.4	3.6	7.7	7.9	367.4	5.5	8.3	8.5	331.0	33.5	37.0	38.0	373.4	7.7	NA
Trial4_621	110	NA()	1.2	12.7700	1.0	2.0901	3.2	0.1937	3.1	0.95	1141.2	8.5	21.4	21.9	1145.5	6.5	15.2	15.7	1154.7	9.3	16.6	18.2	1154.7	16.6	98.8
Trial4_622	680	NA()	3.2	13.6612	0.8	1.7446	1.3	0.1729	1.0	0.79	1028.3	2.8	18.1	18.6	1025.2	2.3	13.1	13.6	1019.5	3.9	14.5	16.4	1019.5	14.5	100.9
Trial4_623	207	NA()	0.8	19.7476	2.0	0.2622	4.2	0.0376	3.7	0.87	237.7	1.8	4.7	4.8	236.4	2.8	5.0	5.2	224.4	24.6	29.3	30.6	237.7	4.7	NA
Trial4_624	70	NA()	1.9	12.1823	1.2	2.4325	3.6	0.2150	3.4	0.94	1255.5	9.9	23.6	24.2	1252.3	7.4	16.2	16.7	1247.6	11.0	17.4	18.9	1247.6	17.4	100.6
Trial4_625	38	NA()	0.9	8.8345	1.1	5.1782	3.6	0.3319	3.4	0.95	1847.8	13.0	32.9	33.6	1849.0	8.6	19.1	19.7	1851.3	10.9	16.6	17.9	1851.3	16.6	99.8
Trial4_626	58	NA()	1.2	13.2378	1.5	1.8050	3.9	0.1734	3.6	0.93	1030.7	8.9	20.0	20.4	1047.3	7.6	15.1	15.6	1082.9	14.0	19.7	21.1	1082.9	19.7	95.2
Trial4_627	246	NA()	1.0	18.0115	1.3	0.5639	3.4	0.0737	3.1	0.93	458.4	2.8	8.8	9.0	454.0	3.2	8.0	8.3	433.2	13.3	20.3	22.0	458.4	8.8	105.8
Trial4_628	91	NA()	1.9	12.7219	1.1	2.1134	3.4	0.1951	3.2	0.95	1148.9	8.8	21.7	22.2	1153.2	6.8	15.4	15.9	1162.2	10.2	17.0	18.6	1162.2	17.0	98.9
Trial4_629	273	NA()	1.7	18.2099	1.4	0.4145	3.6	0.0548	3.3	0.92	343.8	2.0	6.6	6.8	352.1	2.6	6.5	6.7	408.7	14.4	21.1	22.7	343.8	6.6	NA
Trial4_630	56	NA()	0.7	6.3413	0.8	10.1354	2.9	0.4664	2.8	0.96	2467.6	12.6	40.5	41.5	2447.2	7.5	19.9	20.7	2431.1	8.9	14.7	16.0	2431.1	14.7	101.5
Trial4_631	234	NA()	0.9	18.4111	1.4	0.5080	3.5	0.0679	3.3	0.92	423.2	2.7	8.2	8.4	417.1	3.1	7.5	7.8	384.1	13.9	20.8	22.5	423.2	8.2	110.2
Trial4_632	54	NA()	1.0	11.1743	1.2	2.8496	3.7	0.2310	3.5	0.94	1340.0	10.7	25.1	25.7	1368.7	8.0	17.1	17.6	1414.7	11.5	17.5	18.9	1414.7	17.5	94.7
Trial4_633	46	NA()	4.0	9.4931	1.1	4.2189	3.6	0.2906	3.4	0.95	1644.5	12.1	29.8	30.5	1677.7	8.3	18.4	19.0	1720.2	10.6	16.5	17.9	1720.2	16.5	95.6
Trial4_634	66	NA()	1.7	12.5478	1.3	2.1462	3.7	0.1954	3.5	0.94	1150.6	9.5	21.9	22.4	1163.8	7.5	15.8	16.3	1189.5	11.9	18.0	19.6	1189.5	18.0	96.7
Trial4_635	92	NA()	1.2	5.5350	0.9	12.9656	2.3	0.5207	2.1	0.91	2702.2	11.1	42.9	44.0	2677.2	6.6	20.0	20.7	2659.0	8.2	14.0	15.4	2659.0	14.0	101.6
Trial4_636	79	NA()	1.7	8.6670	0.9	4.8204	3.1	0.3031	2.9	0.96	1706.9	11.6	30.5	31.2	1788.5	7.5	18.4	19.1	1885.8	8.1	14.8	16.3	1885.8	14.8	90.5
Trial4_637	156	NA()	1.5	14.0681	1.0	1.5862	3.1	0.1619	2.9	0.95	967.4	6.8	18.2	18.6	964.8	5.4	13.6	14.1	959.8	8.7	16.6	18.3	959.8	16.6	100.8
Trial4_638	70	NA()	1.1	10.9072	1.1	3.1426	3.4	0.2487	3.2	0.95	1431.8	10.8	26.4	27.0	1443.2	7.6	17.2	17.7	1460.8	9.8	16.3	17.9	1460.8	16.3	98.0
Trial4_639	117	NA()	1.5	13.2195	1.0	1.9364	3.2	0.1857	3.0	0.95	1098.2	8.2	20.7	21.1	1093.7	6.3	14.8	15.3	1085.7	9.3	16.6	18.3	1085.7	16.6	101.2
Trial4_640	30	NA()	1.7	8.1183	1.2	6.5093	3.7	0.3834	3.5	0.95	2092.3	13.6	36.2	37.1	2047.2	8.7	19.6	20.3	2002.7	11.1	16.5	17.9	2002.7	16.5	104.5
Trial4_641	39	NA()	0.6	13.3326	1.7	1.9037	4.1	0.1842	3.7	0.91	1089.7	9.7	21.2	21.7	1082.4	8.7	15.9	16.4	1068.6	17.4	22.3	23.5	1068.6	22.3	102.0
Trial4_642	151	NA()	1.5	21.4941	4.1	0.0803	5.1	0.0125	3.2	0.61	80.3	0.8	1.7	1.7	78.5	2.2	2.7	2.7	24.9	65.8	67.8	68.5	80.3	1.7	NA
Trial4_643	105	NA()	2.9	17.6677	2.0	0.5344	4.2	0.0685	3.7	0.88	427.2	3.9	8.7	8.9	434.7	4.9	8.6	8.9	476.0	22.4	27.1	28.4	427.2	8.7	89.7
Trial4_644	67	NA()	0.7	8.9993	0.9	5.0781	3.1	0.3316	3.0	0.96	1846.1	11.8	32.4	33.2	1832.5	7.5	18.6	19.2	1817.8	8.8	15.3	16.8	1817.8	15.3	101.6
Trial4_645	44	NA()	2.3	12.3094	1.6	2.1784	4.0	0.1946	3.7	0.92	1146.1	9.9	22.1	22.6	1174.2	8.5	16.3	16.8	1227.2	15.0	20.2	21.5	1227.2	20.2	93.4
Trial4_646	65	NA()	1.5	5.2751	0.9	13.7045	2.7	0.5246	2.5	0.94	2718.5	11.7	43.3	44.4	2729.5	6.9	20.2	20.9	2738.4	8.4	14.1	15.4	2738.4	14.1	99.3
Trial4_647	47	NA()	2.1	13.6391	1.7	1.7078	4.1	0.1690	3.7	0.91	1006.7	8.9	19.6	20.1	1011.5	7.9	15.0	15.5	1022.7	15.7	21.0	22.4	1022.7	21.0	98.4
Trial4_648	24	NA()	2.9	13.2448	2.2	1.9567	4.5	0.1880	3.9	0.87	1110.8	10.3	21.8	22.2	1100.7	10.2	16.9	17.4	1081.9	22.9	26.8	27.8	1081.9	26.8	102.7
Trial4_649	159	NA()	2.2	13.2478	0.9	1.9909	2.8	0.1914	2.7	0.95	1128.8	7.2	20.7	21.2	1112.4	5.4	14.6	15.1	1081.4	7.9	15.9	17.6	1081.4	15.9	104.4

Trial4_650	155	NA()	0.5	10.8920	1.8	0.4785	4.2	0.0378	3.8	0.91	239.3	2.0	4.8	5.0	397.1	4.0	7.7	7.9	1463.4	16.3	20.9	22.1	239.3	4.8	NA
Trial4_651	162	NA()	1.5	10.9601	0.7	3.1126	2.5	0.2475	2.4	0.96	1425.7	8.1	25.4	26.0	1435.8	5.6	16.3	17.0	1451.6	6.5	14.6	16.3	1451.6	14.6	98.2
Trial4_652	62	NA()	1.3	11.8032	1.3	2.3518	3.7	0.2014	3.4	0.94	1182.9	9.5	22.4	22.9	1228.1	7.6	16.2	16.7	1309.2	11.8	17.8	19.3	1309.2	17.8	90.4
Trial4_653	109	NA()	1.6	13.3198	1.1	1.9334	3.3	0.1869	3.1	0.95	1104.3	8.1	20.7	21.2	1092.7	6.2	14.8	15.3	1070.6	9.6	16.8	18.5	1070.6	16.8	103.2
Trial4_654	151	NA()	3.1	9.3112	0.7	4.4669	2.3	0.3018	2.2	0.96	1700.2	9.1	29.5	30.3	1724.8	5.8	17.6	18.2	1755.7	6.2	14.0	15.7	1755.7	14.0	96.8
Trial4_655	98	NA()	2.0	17.3937	1.8	0.7043	4.1	0.0889	3.7	0.90	548.9	4.9	11.0	11.3	541.3	5.3	9.9	10.2	510.5	19.0	24.3	25.7	548.9	11.0	107.5
Trial4_656	95	NA()	2.4	13.4017	1.2	1.8796	3.4	0.1828	3.3	0.94	1082.1	8.4	20.5	21.0	1073.9	6.5	14.8	15.3	1058.2	10.5	17.4	19.0	1058.2	17.4	102.3
Trial4_657	34	NA()	0.7	10.7887	1.4	3.3892	3.9	0.2653	3.6	0.93	1517.0	11.9	28.1	28.7	1501.9	8.8	18.0	18.6	1481.5	13.2	18.5	19.9	1481.5	18.5	102.4
Trial4_658	103	NA()	1.8	13.3410	1.1	1.8695	3.4	0.1810	3.2	0.94	1072.3	8.1	20.2	20.7	1070.3	6.4	14.7	15.2	1067.4	10.2	17.2	18.8	1067.4	17.2	100.5
Trial4_659	75	NA()	1.5	13.4305	1.3	1.8411	3.7	0.1794	3.4	0.94	1063.8	8.6	20.3	20.8	1060.2	7.0	14.9	15.4	1053.9	12.0	18.3	19.9	1053.9	18.3	100.9
Trial4_660	64	NA()	1.7	11.5162	1.2	2.8279	3.5	0.2363	3.4	0.94	1367.5	10.3	25.4	25.9	1363.0	7.5	16.8	17.4	1356.8	10.7	17.0	18.6	1356.8	17.0	100.8
Trial4_661	59	NA()	2.7	8.7094	1.0	5.0963	3.3	0.3221	3.2	0.96	1799.8	11.9	31.8	32.6	1835.5	7.8	18.7	19.3	1877.0	9.2	15.4	16.9	1877.0	15.4	95.9
Trial4_662	286	NA()	1.1	18.3306	1.2	0.5191	3.3	0.0690	3.1	0.93	430.4	2.2	8.1	8.3	424.5	2.6	7.4	7.7	394.0	12.0	19.5	21.3	430.4	8.1	109.2
Trial4_663	206	NA()	2.3	9.7574	0.9	4.1075	2.0	0.2908	1.8	0.90	1645.6	7.7	28.4	29.1	1655.8	5.0	17.1	17.7	1669.6	5.5	13.9	15.6	1669.6	13.9	98.6
Trial4_664	38	NA()	1.9	11.7335	1.5	2.7045	4.0	0.2303	3.7	0.92	1335.8	11.0	25.2	25.8	1329.7	8.9	17.3	17.8	1320.7	15.2	20.2	21.5	1320.7	20.2	101.1
Trial4_665	142	NA()	2.3	12.9786	0.9	2.0078	3.0	0.1891	2.8	0.95	1116.4	7.7	20.7	21.2	1118.1	5.8	14.8	15.3	1122.5	8.4	16.1	17.8	1122.5	16.1	99.5
Trial4_666	95	NA()	4.2	9.1067	0.8	4.9793	2.8	0.3290	2.7	0.96	1833.6	11.2	32.0	32.8	1815.8	7.0	18.3	19.0	1796.2	8.0	14.8	16.4	1796.2	14.8	102.1
Trial4_667	81	NA()	2.1	13.6536	1.3	1.7127	3.7	0.1697	3.4	0.93	1010.3	8.4	19.5	19.9	1013.3	7.2	14.7	15.2	1020.6	13.3	19.3	20.8	1020.6	19.3	99.0
Trial4_668	29	NA()	1.1	13.8776	2.2	1.6661	4.4	0.1678	3.9	0.87	999.8	9.6	19.9	20.3	995.7	9.6	15.9	16.4	987.6	22.4	26.4	27.5	987.6	26.4	101.2
Trial4_669	97	NA()	2.3	10.5061	0.9	3.6892	2.9	0.2812	2.8	0.96	1597.6	10.6	28.6	29.3	1569.0	6.9	17.4	18.0	1531.7	8.2	15.3	16.9	1531.7	15.3	104.3
Trial4_670	31	NA()	1.8	13.2740	2.0	1.8297	4.3	0.1762	3.8	0.89	1046.3	9.9	20.7	21.1	1056.2	9.1	16.0	16.4	1077.5	18.7	23.2	24.5	1077.5	23.2	97.1
Trial4_671	47	NA()	1.6	13.1579	1.6	1.8554	4.0	0.1771	3.7	0.92	1051.4	9.4	20.5	21.0	1065.4	8.3	15.6	16.0	1095.1	15.8	20.9	22.3	1095.1	20.9	96.0
Trial4_672	88	NA()	1.5	13.1885	1.2	1.9857	3.5	0.1900	3.3	0.94	1121.5	8.9	21.3	21.8	1110.7	6.9	15.2	15.7	1090.4	10.8	17.5	19.1	1090.4	17.5	102.8
Trial4_673	33	NA()	3.3	18.0416	3.9	0.5262	5.0	0.0689	3.2	0.64	429.4	4.9	9.2	9.4	429.3	9.5	11.8	12.0	429.5	54.4	56.5	57.1	429.4	9.2	100.0
Trial4_674	96	NA()	2.2	18.3181	2.2	0.5219	4.3	0.0694	3.8	0.87	432.3	4.0	8.8	9.0	426.4	5.3	8.8	9.0	395.5	26.7	30.9	32.0	432.3	8.8	109.3
Trial4_675	64	NA()	1.9	13.4793	1.4	1.7850	3.8	0.1746	3.5	0.93	1037.3	8.8	20.0	20.5	1040.0	7.4	15.0	15.4	1046.6	13.6	19.4	20.9	1046.6	19.4	99.1
Trial4_676	93	NA()	2.3	9.6049	0.8	4.5177	2.9	0.3148	2.7	0.96	1764.5	10.8	31.0	31.7	1734.2	6.9	18.0	18.6	1698.7	8.0	15.0	16.5	1698.7	15.0	103.9
Trial4_677	136	NA()	1.0	9.0810	0.7	4.7549	2.4	0.3133	2.3	0.96	1757.0	9.7	30.5	31.2	1777.0	6.1	17.9	18.5	1801.4	6.6	14.1	15.8	1801.4	14.1	97.5
Trial4_678	161	NA()	2.2	13.2911	0.9	1.8239	2.9	0.1759	2.8	0.95	1044.5	6.8	19.4	19.8	1054.1	5.3	14.2	14.7	1074.9	8.0	15.9	17.7	1074.9	15.9	97.2
Trial4_679	23	NA()	2.1	13.3393	2.3	1.8847	4.6	0.1824	3.9	0.86	1080.2	10.3	21.3	21.8	1075.7	10.6	17.0	17.4	1067.6	24.4	28.1	29.1	1067.6	28.1	101.2
Trial4_680	183	NA()	1.5	13.6558	0.9	1.6839	2.8	0.1668	2.7	0.95	994.7	6.1	18.3	18.8	1002.5	4.9	13.6	14.1	1020.3	7.9	16.0	17.8	1020.3	16.0	97.5
Trial4_681	26	NA()	3.4	9.6191	1.5	4.3266	4.0	0.3020	3.7	0.93	1701.1	12.5	30.7	31.4	1698.5	9.2	18.9	19.5	1696.0	13.5	18.6	19.8	1696.0	18.6	100.3
Trial4_682	151	NA()	1.0	8.8064	0.9	5.3546	2.2	0.3422	2.0	0.92	1897.0	9.0	32.2	33.0	1877.6	5.5	18.0	18.7	1857.0	6.2	13.9	15.5	1857.0	13.9	102.2
Trial4_683	74	NA()	2.3	13.4008	1.3	1.9040	3.6	0.1851	3.4	0.93	1095.0	8.5	20.7	21.2	1082.5	6.9	15.0	15.5	1058.4	12.4	18.6	20.1	1058.4	18.6	103.5
Trial4_684	78	NA()	0.9	12.7873	1.2	2.1023	3.5	0.1951	3.3	0.94	1148.7	8.7	21.6	22.1	1149.5	6.8	15.4	15.9	1152.0	11.0	17.5	19.1	1152.0	17.5	99.7
Trial4_685	47	NA()	2.4	12.9385	1.6	1.9205	4.0	0.1803	3.7	0.91	1068.6	8.5	20.4	20.8	1088.2	8.0	15.6	16.1	1128.7	16.6	21.5	22.8	1128.7	21.5	94.7
Trial4_686	71	NA()	2.0	16.8304	1.9	0.8722	4.2	0.1065	3.7	0.89	652.4	5.5	12.9	13.2	636.7	6.4	11.4	11.7	582.4	21.9	26.5	27.8	652.4	12.9	112.0
Trial4_687	157	NA()	1.0	20.1665	3.3	0.1863	5.1	0.0273	3.9	0.76	173.4	1.3	3.5	3.6	173.4	2.6	4.1	4.2	175.7	34.0	37.6	38.7	173.4	3.5	NA
Trial4_688	437	NA()	3.3	13.6425	0.6	1.6250	1.6	0.1609	1.5	0.93	961.5	1.0	16.8	17.3	979.9	1.7	12.7	13.2	1022.3	4.9	14.7	16.6	1022.3	14.7	94.1
Trial4_689	109	NA()	0.8	9.9864	0.8	4.0795	2.7	0.2956	2.6	0.96	1669.5	8.9	29.0	29.8	1650.2	5.8	17.3	18.0	1626.6	7.1	14.6	16.3	1626.6	14.6	102.6
Trial4_690	220	NA()	2.9	5.3169	0.8	14.0817	1.3	0.5433	1.0	0.78	2797.1	5.6	43.0	44.1	2755.2	3.3	19.3	20.0	2725.4	4.1	12.1	13.6	2725.4	12.1	102.6
Trial4_691	144	NA()	1.0	5.5005	0.9	11.7816	1.8	0.4702	1.6	0.88	2484.5	7.7	39.5	40.5	2587.2	4.5	19.3	20.0	2669.4	5.1	12.5	14.0	2669.4	12.5	93.1
Trial4_692	99	NA()	0.9	5.0926	0.9	14.3042	2.1	0.5286	2.0	0.91	2735.4	8.8	42.8	43.9	2770.1	5.2	19.7	20.4	2796.2	6.2	12.8	14.3	2796.2	12.8	97.8
Trial4_693	165	NA()	2.4	13.2756	0.9	1.9113	2.8	0.1841	2.7	0.95	1089.4	5.3	19.6	20.1	1085.0	4.4	14.0	14.6	1077.2	7.9	15.9	17.6	1077.2	15.9	101.1
Trial4_694	206	NA()	15.0	13.3525	0.8	1.8071	2.6	0.1751	2.5	0.95	1040.0	4.9	18.7	19.2	1048.0	4.1	13.7	14.2	1065.6	7.2	15.6	17.3	1065.6	15.6	97.6
Trial4_695	242	NA()	1.3	10.6207	0.8	3.2042	2.0	0.2469	1.8	0.90	1422.6	5.1	24.5	25.2	1458.2	3.7	15.9	16.5	1511.2	5.0	13.9	15.7	1511.2	13.9	94.1
Trial4_696	194	NA()	1.9	8.6600	0.8	5.5496	1.8	0.3487	1.6	0.88	1928.5	7.0	32.1	32.9	1908.3	4.3	17.8	18.4	1887.3	5.0	13.4	15.0	1887.3	13.4	102.2
Trial4_697	112	NA()	7.2	13.5442	1.1	1.7169	3.3	0.1687	3.2	0.94	1005.1	6.5	18.7	19.1	1014.9	5.5	14.0	14.5	1036.9	10.1	17.2	18.9	1036.9	17.2	96.9
Trial4_698	279	NA()	3.1	17.5099	1.2	0.5391	3.3	0.0685	3.1	0.93	427.1	1.8	8.0	8.2	437.8	2.5	7.5	7.8	495.8	12.1	19.4	21.2	427.1	8.0	86.1
Trial4_699	36	NA()	0.7	10.8436	1.4	3.2380	3.9	0.2548	3.6	0.93	1463.0	10.3	26.7	27.3	1466.3	8.1	17.5	18.1	1471.9	12.9	18.4	19.7	1471.9	18.4	99.4
Trial4_700	10	NA()	1.1	13.8122	5.9	1.7184	6.7	0.1722	3.2	0.47	1024.3	10.2	20.5	20.9	1015.4	33.6	36.0	36.2	997.2	104.1	105.0	105.3	997.2	105.0	102.7
Trial4_701	50	NA()	1.3	13.6638	1.6	1.7809	4.0	0.1766	3.6	0.92	1048.2	8.0	19.9	20.3	1038.5	7.3	14.9	15.4	1019.1	15.2	20.6	22.0	1019.1	20.6	102.9
Trial4_702	176	NA()	0.3	13.4417	0.9	1.8290	2.8	0.1784	2.6	0.95	1058.1	5.4	19.1	19.6	1055.9	4.4	13.9	14.4	1052.2	7.6	15.8	17.6	1052.2	15.8	100.6
Trial4_703	482	NA()	0.9	18.4871	1.1	0.3845	3.0	0.0516	2.8	0.93	324.2	0.4	6.0	6.1	330.4	1.4	5.8	6.0	374.9	10.9	18.9	20.8	324.2	6.0	NA
Trial4_704	89	NA()	1.3	18.7780	2.2	0.4742	4.4	0.0646	3.8	0.86	403.6	3.3	8.1	8.3	394.1	4.5	8.0	8.2	339.7	24.8	29.3	30.6	403.6	8.1	118.8
Trial4_705	77	NA()	1.7	16.3261	2.1	0.6706	4.3	0.0794	3.8	0.88	492.8	4.1	9.8	10.0	521.1	5.5	9.8	10.1	648.1	22.3	26.7	28.0	492.8	9.8	76.0
Trial4_706	251	NA()	3.0	9.2611	0.8	4.6644	1.6	0.3134	1.3	0.85	1757.6	5.8	29.5	30.2	1760.9	3.7	17.1	17.8	1765.6	4.4	13.3	15.0	1765.6	13.3	99.5
Trial4_707	218	NA()	2.5	8.9436	0.8	4.4124	1.9	0.2863	1.7	0.90	1623.2	6.4	27.7	28.4	1714.7	4.3	17.1	17.8	1829.0	4.8	13.3	15.0	1829.0	13.3	88.7
Trial4_708	344	NA()	2.1	20.6589	2.4	0.1078	4.4	0.0162	3.7	0.84	103.3	0.4	2.0	2.0	103.9	1.3	2.4	2.4	119.1	29.2	33.3	34.6	103.3	2.0	NA
Trial4_709	60	NA()	1.0	17.9429	2.6	0.5733	4.7	0.0746	3.9	0.83	464.0	4.4	9.5	9.7	460.1	6.6	9.9	10.1	441.7	33.2	36.5	37.5	464.0	9.5	105.1
Trial4_710	78	NA()	1.7	17.8806	1.9	0.7798	4.2	0.1012	3.7	0.89	621.3	5.3	12.3	12.6	585.4	5.8	10.6	10.9	449.4	21.1	26.1	27.4	621.3	12.3	138.2
Trial4_711	58	NA()	1.2	4.4157	0.9	16.9616	2.8	0.5434	2.6	0.94	2797.9	10.8	44.0	45.1	2932.7	6.5	20.3	21.0	3027.2	7.8	13.5	14.8	3027.2	13.5	92.4
Trial4_712	282	NA()	4.4	13.6390	0.7	1.7448	2.2	0.1727	2.1	0.94	1026.8	4.3	18.4	18.8	1025.3	3.6	13.4	13.9	1022.8	6.3	15.3	17.1	1022.8	15.3	100.4
Trial4_713	64	NA()	3.1	11.2153	1.1	2.9258	3.5	0.2381	3.3	0.95	1376.8	10.2	25.4	26.0	1388.6	7.5	16.9	17.4	1407.7	10.5	16.8	18.3	1407.7	16.8	97.8
Trial4_714	114	NA()	0.7	5.7621	0.9	11.5926	2.1	0.4847	1.9	0.90	2547.6	9.9	40.8	41.8	2572.1	5.9	19.6	20.4	2592.1	7.0	13.4	14.9	2592.1	13.4	98.3
Trial4_715	92	NA()	2.5	10.4637	0.9	3.4101	3.1	0.2589	2.9	0.96	1484.3	9.5	26.7	27.3	1506.7	6.6	17.0	17.6	1539.3	8.2	15.3	16.9	1539.3	15.3	96.4
Trial4_716	400	NA()	1.6	11.6764	1.1	0.4945	3.3	0.0419	3.1	0.95	264.6	0.7	4.9	5.1	407.9	1.8	7.0	7.2	1330.1	9.0	16.1	17.7	264.6	4.9	NA
Trial4_717	134	NA()	2.6	13.4159	1.0	1.8343	3.1	0.1786	2.9	0.95	1059.1	7.0	19.6	20.1	1057.8	5.5	14.3	14.8	1056.1	9.0	16.6	18.3	1056.1	16.6	100.3
Trial4_718	51	NA()	1.4	9.5466	1.1	4.0414	3.5	0.2799	3.4	0.95	1591.1	11.0	28.7	29.4	1642.6	7.7	18.0	18.6	1709.9	9.8	16.0	17.5	1709.9	16.0	93.1
Trial4_719	62	NA()	1.3	15.0774	2.1	0.8177	4.4	0.0895	3.8	0.88	552.3	5.3	11.3	11.5	606.8	6.6	11.3	11.6	816.7	22.0	26.3	27.4	552.3	11.3	67.6
Trial4_720	164	NA()	1.9	13.3920	0.9	1.8087	2.9	0.1758	2.8	0.95	1043.7	6.8	19.3	19.8	1048.6	5.3	14.1	14.6	1059.7	8.2	16.1	17.8	1059.7	16.1	98.5
Trial4_721	13	NA()	0.8	5.8052	1.4	11.5161	4.0	0.4851	3.8	0.94	2549.4	15.8	42.6	43.6	2565.9	10.1	21.3	21.9	2579.7	13.0	17.3	18.5	2579.7	17.3	98.8
Trial4_722	157	NA()	2.0	13.7830	1.0	1.7017	3.0	0.1702	2.8	0.95	1013.1	6.8	18.9	19.3	1009.2	5.4	13.9	14.4	1001.5	8.5	16.4	18.1	1001.5	16.4	101.2
Trial4_723	152	NA()	0.9	18.7343	1.8	0.4471	4.0	0.0608	3.6	0.90	380.4	3.1	7.6	7.8	375.3	3.7	7.3	7.5	344.9	18.9	24.5	26.0	380.4	7.6	NA
Trial4_724	201	NA()	1.4	19.2734	2.0	0.2801	4.2	0.0392	3.7	0.88	247.7	1.9	4.9	5.0	250.7	2.7	5.2	5.4	280.4	21.6	26.7	28.1	247.7	4.9	NA
Trial4_725	201	NA()	0.8	18.3337	1.5	0.5211	3.7	0.0693	3.4	0.92	432.0	3.0	8.4	8.6	425.9	3.5	7.8	8.0	393.6	15.1	21.6	23.3	432.0	8.4	109.8
Trial4_726	65	NA()	2.2	11.9475	1.3	2.2853	3.7	0.1981	3.4	0.94	1165.2	9.3	22.1	22.6	1207.8	7.5	16.0	16.5	1285.6	11.6	17.7	19.3	1285.6	17.7	90.6
Trial4_727	247	NA()	9.4	5.6129	0.8	11.2324	1.4	0.4575	1.1	0.78	2428.3	7.1	38.7	39.7	2542.6	4.1	19.1	19.9	2635.8	4.5	12.3	13.8	2635.8	12.3	92.1
Trial4_728	81	NA()	1.6	7.1862	0.8	6.2175	3.0	0.3242	2.9	0.97	1810.2	11.1	31.7	32.4	2006.9	7.2	18.9	19.6	2216.7	7.2	13.9	15.4	2216.7	13.9	81.7
Trial4_729	395	NA()	2.4	9.1586	0.8	4.6354	1.3	0.3080	1.0	0.78	1731.1	3.2	28.7	29.5	1755.7	2.2	16.9	17.6	1785.9	3.1	12.9	14.7	1785.9	12.9	96.9
Trial4_730	184	NA()	1.4	18.0149	1.5	0.5620	3.7	0.0735	3.4	0.91	456.9	3.3	8.9	9.1	452.8	3.8	8.3	8.5	432.8	16.5	22.5	24.1	456.9	8.9	105.6
Trial4_731	635	NA()	3.8	16.2413	1.1	0.3325	3.0	0.0392	2.8	0.94	247.8	0.7	4.6	4.7	291.5	1.4	5.2	5.5	659.2	9.5	17.5	19.4	247.8	4.6	NA
Trial4_732	64	NA()	1.0	9.5774	1.0	3.9701	3.4	0.2759	3.2	0.95	1570.7	11.5	28.6	29.3	1628.1	7.9	18.0	18.7	1704.0	9.4	15.8	17.3	1704.0	15.8	92.2
Trial4_733	125	NA()	1.4	16.8769	1.6	0.7252	3.9	0.0888	3.5	0.91	548.4	4.8	11.0	11.2	553.7	5.0	9.9	10.2	576.4	16.1	22.0	23.5	548.4	11.0	95.2
Trial4_734	57	NA()	1.1	11.2373	1.2	3.0004	3.6	0.2446	3.4	0.94	1410.8	11.1	26.3	26.9	1407.7	8.0	17.2	17.8	1403.9	10.9	17.1	18.6	1403.9	17.1	100.5
Trial4_735	222	NA()	1.6	11.1961	0.6	3.1440	2.0	0.2554	1.9	0.95	1466.3	7.3	25.7	26.4	1443.5	4.9	16.2	16.8	1410.9	5.7	14.4	16.1	1410.9	14.4	103.9
Trial4_736	62	NA()	0.8	10.9809	1.1	3.3087	3.5	0.2636	3.3	0.95	1508.4	11.4	27.7	28.4	1483.1	7.8	17.5	18.0	1448.0	10.2	16.6	18.1	1448.0	16.6	104.2
Trial4_737	18	NA()	2.1	11.6577	2.2	2.6018	4.5	0.2201	3.9	0.87	1282.3	11.8	24.8	25.3	1301.1	10.9	18.3	18.8	1333.2	20.9	24.7	25.8	1333.2	24.7	96.2
Trial4_738	60	NA()	0.7	13.3362	1.4	1.8343	3.8	0.1775	3.6	0.93	1053.3	8.8	20.3	20.7	1057.8	7.4	15.1	15.6	1068.1	13.4	19.3	20.8	1068.1	19.3	98.6
Trial4_739	211	NA()	2.6	12.0025	0.7	2.3453	2.4	0.2043	2.3	0.95	1198.1	6.3	21.5	22.0	1226.1	4.7	15.0	15.6	1276.6	6.3	14.8	16.6	1276.6	14.8	93.9
Trial4_740	34	NA()	1.3	5.1698	0.9	13.1423	3.3	0.4930	3.2	0.97	2583.6	13.1	42.1	43.1	2689.9	7.8	20.4	21.2	2771.6	9.1	14.5	15.8	2771.6	14.5	93.2
Trial4_741	69	NA()	0.2	4.0311	1.0	21.6141	2.4	0.6322	2.2	0.92	3158.3	10.2	48.0	49.3	3166.5	6.5	20.5	21.3	3172.5	8.4	13.8	15.0	3172.5	13.8	99.6
Trial4_742	81	NA()	1.9	13.7232	1.3	1.7862	3.6	0.1779	3.4	0.93	1055.3	8.3	20.1	20.5	1040.4	6.8	14.7	15.2	1010.3	12.3	18.6	20.2	1010.3	18.6	104.5
Trial4_743	263	NA()	2.3	13.5990	0.7	1.7220	2.3	0.1699	2.2	0.95	1011.7	4.6	18.2	18.6	1016.8	3.7	13.4	13.9	1028.7	6.4	15.3	17.2	1028.7	15.3	98.3
Trial4_744	93	NA()	1.8	13.0170	1.1	1.9384	3.5	0.1831	3.3	0.94	1083.8	8.4	20.5	21.0	1094.5	6.6	14.9	15.5	1116.6	10.4	17.2	18.8	1116.6	17.2	97.1
Trial4_745	203	NA()	1.6	16.9072	1.5	0.5391	3.7	0.0661	3.4	0.92	412.8	2.7	8.0	8.2	437.8	3.4	7.9	8.2	572.5	14.8	21.1	22.6	412.8	8.0	72.1
Trial4_746	160	NA()	1.6	12.8377	0.9	2.0101	2.8	0.1872	2.7	0.95	1106.4	6.6	20.2	20.7	1118.9	5.1	14.5	15.0	1144.2	7.6	15.6	17.4	1144.2	15.6	96.7
Trial4_747	95	NA()	1.3	19.4599	3.4	0.2832	5.2	0.0400	3.9	0.76	252.7	2.4	5.2	5.3	253.2	4.1	6.1	6.2	258.2	35.5	38.9	39.8	252.7	5.2	NA
Trial4_748	111	NA()	1.8	13.6061	1.1	1.6929	3.4	0.1671	3.2	0.94	996.3	6.9	18.7	19.1	1005.9	5.7	14.0	14.5	1027.6	10.0	17.1	18.8	1027.6	17.1	97.0
Trial4_749	124	NA()	1.3	13.3444	1.0	1.7949	3.2	0.1738	3.0	0.95	1033.0	6.8	19.2	19.6	1043.6	5.5	14.2	14.7	1066.8	9.1	16.5	18.3	1066.8	16.5	96.8
Trial4_750	45	NA()	1.2	4.6698	1.0	17.1607	3.0	0.5815	2.8	0.95	2954.8	11.1	45.9	47.1	2943.9	6.8	20.4	21.1	2937.1	8.7	14.1	15.4	2937.1	14.1	100.6
Trial4_751	335	NA()	0.7	20.8238	2.0	0.1798	4.1	0.0272	3.5	0.87	172.8	0.7	3.3	3.4	167.9	1.7	3.5	3.7	100.3	24.6	29.5	30.9	172.8	3.3	NA
Trial4_752	24	NA()	1.5	7.8620	1.3	6.8168	3.8	0.3889	3.6	0.94	2117.6	13.2	36.4	37.2	2087.9	8.7	19.7	20.4	2059.5	11.5	16.7	18.0	2059.5	16.7	102.8
Trial4_753	52	NA()	1.8	13.3768	1.6	1.6913	4.0	0.1642	3.7	0.91	979.8	8.1	18.9	19.3	1005.2	7.5	14.8	15.3	1062.0	15.6	20.9	22.3	1062.0	20.9	92.3
Trial4_754	77	NA()	1.1	17.3392	2.1	0.6694	4.3	0.0842	3.8	0.88	521.2	4.5	10.4	10.7	520.3	5.5	9.9	10.1	517.3	22.5	27.1	28.3	521.2	10.4	100.8
Trial4_755	142	NA()	2.7	12.5046	0.9	2.1257	3.0	0.1929	2.8	0.95	1136.9	6.8	20.7	21.2	1157.2	5.3	14.8	15.3	1196.2	8.0	15.8	17.4	1196.2	15.8	95.0
Trial4_756	101	NA()	1.0	11.1891	0.9	3.1070	3.0	0.2522	2.9	0.95	1450.1	8.9	26.0	26.6	1434.4	6.2	16.6	17.2	1412.1	8.1	15.5	17.1	1412.1	15.5	102.7
Trial4_757	63	NA()	1.7	8.9236	0.9	4.9370	3.2	0.3197	3.1	0.96	1788.1	10.8	31.3	32.0	1808.6	7.1	18.3	19.0	1833.1	8.6	15.2	16.7	1833.1	15.2	97.5
Trial4_758	81	NA()	1.9	12.4842	1.2	2.1984	3.5	0.1991	3.3	0.94	1170.7	8.3	21.8	22.3	1180.5	6.6	15.4	16.0	1199.5	10.5	17.2	18.8	1199.5	17.2	97.6
Trial4_759	49	NA()	1.5	12.8124	1.5	1.9987	3.9	0.1858	3.6	0.92	1098.6	8.8	20.9	21.4	1115.1	7.7	15.6	16.1	1148.1	14.5	19.9	21.3	1148.1	19.9	95.7
Trial4_760	47	NA()	1.1	10.7460	1.3	3.2277	3.7	0.2517	3.5	0.94	1447.1	10.0	26.3	27.0	1463.8	7.5	17.2	17.8	1489.0	11.1	17.1	18.6	1489.0	17.1	97.2
Trial4_761	26	NA()	2.6	13.3298	2.2	1.8144	4.5	0.1755	3.9	0.87	1042.3	9.1	20.3	20.7	1050.7	9.6	16.3	16.7	1069.1	22.6	26.5	27.6	1069.1	26.5	97.5
Trial4_762	172	NA()	1.4	19.1059	2.1	0.3057	4.2	0.0424	3.7	0.87	267.5	1.9	5.3	5.4	270.8	3.0	5.6	5.8	300.3	23.8	28.5	29.8	267.5	5.3	NA
Trial4_763	128	NA()	2.0	13.1464	1.0	1.9673	3.1	0.1877	2.9	0.95	1108.7	6.9	20.4	20.9	1104.4	5.5	14.5	15.1	1096.8	8.9	16.4	18.1	1096.8	16.4	101.1
Trial4_764	239	NA()	1.9	9.7486	0.8	4.1254	1.8	0.2918	1.5	0.88	1650.6	6.0	28.0	28.7	1659.4	4.0	16.8	17.5	1671.3	4.7	13.6	15.3	1671.3	13.6	98.8
Trial4_765	60	NA()	1.6	10.0988	1.1	3.7846	3.4	0.2773	3.2	0.95	1577.9	10.4	28.3	28.9	1589.5	7.3	17.6	18.2	1605.8	9.6	16.0	17.5	1605.8	16.0	98.3
Trial4_766	26	NA()	1.3	18.8517	5.9	0.2837	6.7	0.0388	3.2	0.48	245.4	3.0	5.4	5.5	253.6	11.7	12.6	12.6	330.8	115.4	116.5	116.8	245.4	5.4	NA
Trial4_767	132	NA()	1.3	13.7215	1.0	1.6583	3.2	0.1651	3.0	0.94	985.1	6.3	18.3	18.7	992.7	5.3	13.7	14.2	1010.6	9.4	16.8	18.5	1010.6	16.8	97.5
Trial4_768	82	NA()	1.2	9.1987	0.9	4.4200	3.0	0.2950	2.9	0.96	1666.5	10.0	29.3	30.0	1716.1	6.6	17.8	18.5	1777.9	7.7	14.7	16.3	1777.9	14.7	93.7
Trial4_769	31	NA()	1.5	18.1842	5.0	0.2853	5.9	0.0376	3.2	0.55	238.2	2.8	5.2	5.3	254.8	9.3	10.3	10.4	411.9	88.5	89.8	90.2	238.2	5.2	NA
Trial4_770	646	NA()	2.0	13.5890	0.8	1.7368	1.3	0.1712	1.0	0.79	1019.0	2.9	17.9	18.4	1022.3	2.4	13.1	13.6	1030.2	4.0	14.5	16.4	1030.2	14.5	98.9
Trial4_771	77	NA()	1.0	8.4257	0.8	5.4798	3.0	0.3350	2.9	0.96	1862.6	10.8	32.3	33.0	1897.4	6.9	18.5	19.2	1936.5	7.8	14.6	16.1	1936.5	14.6	96.2
Trial4_772	62	NA()	2.0	13.5941	1.5	1.7165	3.9	0.1693	3.6	0.92	1008.3	8.1	19.3	19.8	1014.7	7.2	14.7	15.2	1029.4	14.1	19.9	21.3	1029.4	19.9	98.0
Trial4_773	54	NA()	0.6	18.5051	3.4	0.3884	4.7	0.0521	3.2	0.68	327.7	3.4	6.9	7.0	333.2	5.6	8.0	8.1	372.7	37.4	40.5	41.4	327.7	6.9	NA
Trial4_774	47	NA()	1.3	11.3512	1.4	2.4094	3.9	0.1985	3.6	0.93	1167.0	9.8	22.3	22.8	1245.4	8.2	16.5	17.0	1384.6	12.9	18.5	19.9	1384.6	18.5	84.3
Trial4_775	50	NA()	0.8	12.0368	1.4	2.4415	3.8	0.2132	3.6	0.93	1246.0	10.1	23.6	24.1	1254.9	8.1	16.6	17.1	1271.0	13.5	19.0	20.4	1271.0	19.0	98.0
Trial4_776	67	NA()	0.9	12.7674	1.3	1.9744	3.7	0.1829	3.5	0.93	1082.8	9.0	20.8	21.3	1106.8	7.4	15.4	15.9	1155.1	12.1	18.3	19.8	1155.1	18.3	93.7
Trial4_777	24	NA()	2.0	13.3567	2.3	1.7123	4.6	0.1659	3.9	0.86	989.7	9.7	19.8	20.2	1013.1	10.3	16.4	16.8	1065.0	24.2	27.8	28.9	1065.0	27.8	92.9
Trial4_778	97	NA()	1.2	13.7573	1.2	1.7622	3.5	0.1759	3.3	0.94	1044.6	8.2	19.9	20.3	1031.6	6.6	14.5	15.0	1005.3	11.1	17.9	19.5	1005.3	17.9	103.9
Trial4_779	129	NA()	1.1	13.5094	1.1	1.5536	3.3	0.1523	3.2	0.94	913.8	6.9	17.4	17.8	951.9	5.9	13.7	14.2	1042.1	10.2	17.2	18.9	1042.1	17.2	87.7
Trial4_780	194	NA()	1.4	19.6949	2.5	0.2090	4.5	0.0299	3.8	0.84	189.8	1.5	3.8	3.9	192.7	2.8	4.5	4.6	230.6	32.2	35.9	37.0	189.8	3.8	NA
Trial4_781	75	NA()	1.6	5.4026	0.9	13.0345	2.6	0.5110	2.4	0.93	2660.7	11.6	42.6	43.7	2682.2	6.9	20.1	20.8	2699.1	8.2	14.0	15.3	2699.1	14.0	98.6
Trial4_782	73	NA()	1.4	13.0457	1.3	1.9297	3.7	0.1827	3.4	0.94	1081.5	8.8	20.7	21.1	1091.4	7.1	15.2	15.7	1112.2	11.8	18.1	19.7	1112.2	18.1	97.2
Trial4_783	32	NA()	0.9	5.1051	0.9	12.6802	3.4	0.4697	3.3	0.96	2482.2	13.5	41.0	42.0	2656.2	8.1	20.5	21.2	2792.2	9.2	14.5	15.8	2792.2	14.5	88.9
Trial4_784	110	NA()	1.6	10.3769	0.8	3.4808	2.9	0.2621	2.8	0.96	1500.5	9.7	27.0	27.6	1522.9	6.6	17.1	17.7	1554.9	7.7	15.1	16.7	1554.9	15.1	96.5
Trial4_785	180	NA()	1.5	11.8270	0.7	2.5595	2.5	0.2196	2.3	0.95	1280.0	7.4	23.0	23.6	1289.1	5.3	15.5	16.1	1305.3	6.6	14.9	16.6	1305.3	14.9	98.1
Trial4_786	207	NA()	1.6	12.6157	0.8	1.8345	2.7	0.1679	2.5	0.95	1000.7	5.9	18.4	18.8	1057.9	4.8	14.0	14.5	1178.8	7.0	15.3	17.1	1178.8	15.3	84.9
Trial4_787	150	NA()	1.6	15.7862	1.3	0.8595	3.6	0.0985	3.3	0.93	605.3	4.8	11.9	12.2	629.9	4.8	10.6	10.9	719.9	12.7	19.4	21.0	605.3	11.9	84.1

Trial4_788	112	NA()	1.6	9.5609	0.8	3.7821	2.9	0.2624	2.7	0.96	1502.0	10.0	27.1	27.7	1588.9	6.8	17.5	18.1	1707.1	7.4	14.6	16.2	1707.1	14.6	88.0
Trial4_789	105	NA()	1.0	18.4760	2.1	0.4878	4.3	0.0654	3.8	0.88	408.4	3.9	8.4	8.6	403.4	4.7	8.2	8.4	376.2	22.8	27.5	28.9	408.4	8.4	108.6
Trial4_790	131	NA()	3.1	13.5005	1.0	1.6967	3.2	0.1662	3.0	0.95	991.2	7.4	18.8	19.2	1007.3	6.0	14.1	14.6	1043.4	9.5	16.8	18.5	1043.4	16.8	95.0
Trial4_791	138	NA()	4.0	8.9359	0.9	5.2903	2.3	0.3430	2.1	0.92	1901.2	10.3	32.6	33.4	1867.3	6.2	18.2	18.9	1830.6	6.9	14.2	15.8	1830.6	14.2	103.9
Trial4_792	94	NA()	1.1	15.7340	2.1	1.0501	3.8	0.1199	3.2	0.83	729.9	5.9	13.0	13.4	728.9	5.8	11.8	12.1	726.9	15.1	26.3	27.5	729.9	22.0	100.4
Trial4_793	277	NA()	2.1	11.8876	0.9	2.4080	1.9	0.2077	1.6	0.88	1216.6	4.7	19.3	19.9	1245.0	3.6	14.6	15.2	1295.3	5.2	20.4	21.7	1295.3	17.0	93.9
Trial4_794	167	NA()	1.8	11.0615	1.0	3.0110	2.4	0.2417	2.1	0.90	1395.4	7.5	22.4	23.1	1410.4	5.2	15.9	16.5	1434.0	6.5	20.4	21.7	1434.0	19.3	97.3
Trial4_795	91	NA()	2.9	9.0249	1.1	4.7296	2.7	0.3097	2.5	0.92	1739.3	10.2	27.6	28.4	1772.5	6.6	17.8	18.4	1812.6	7.7	20.0	21.1	1812.6	20.0	96.0
Trial4_796	373	NA()	1.5	13.8164	0.9	1.6814	1.7	0.1686	1.5	0.86	1004.2	2.4	15.9	16.4	1001.5	2.3	12.7	13.3	996.6	5.3	21.3	22.6	996.6	18.1	100.8
Trial4_797	20	NA()	3.1	5.2645	1.4	13.5355	3.5	0.5170	3.2	0.91	2686.6	13.6	39.5	40.6	2717.8	8.5	20.5	21.2	2741.7	10.7	19.8	20.8	2741.7	23.7	98.0
Trial4_798	71	NA()	1.1	17.8171	3.2	0.5661	4.7	0.0732	3.5	0.74	455.3	4.3	8.5	8.8	455.5	5.6	9.1	9.4	457.4	25.8	34.2	35.3	455.3	15.3	99.6
Trial4_799	71	NA()	1.1	13.2184	2.1	1.4577	3.8	0.1398	3.2	0.84	843.6	7.0	15.1	15.5	913.1	6.7	13.6	14.1	1085.9	13.8	24.5	25.7	1085.9	41.3	77.7
Trial4_800	69	NA()	1.2	11.0633	1.5	3.1008	3.3	0.2489	2.9	0.89	1432.9	9.9	23.8	24.5	1432.9	7.1	16.7	17.3	1433.7	9.8	21.7	22.9	1433.7	28.4	99.9
Trial4_801	851	NA()	3.5	13.3676	0.8	1.7805	1.5	0.1727	1.2	0.82	1027.0	6.4	17.2	17.7	1038.4	4.6	13.6	14.1	1063.4	4.2	20.8	22.2	1063.4	16.6	96.6
Trial4_802	78	NA()	1.4	9.2565	1.2	4.5619	2.9	0.3064	2.6	0.91	1723.0	10.5	27.6	28.4	1742.3	6.9	17.8	18.4	1766.5	8.1	20.2	21.4	1766.5	21.7	97.5
Trial4_803	500	NA()	4.0	18.7693	1.5	0.3959	2.9	0.0539	2.5	0.85	338.5	0.4	5.6	5.8	338.7	1.4	5.8	6.1	340.7	10.9	25.4	26.8	338.5	8.1	NA
Trial4_804	121	NA()	1.9	12.8306	1.4	1.9987	3.1	0.1861	2.7	0.89	1100.1	7.4	18.6	19.1	1115.1	5.8	14.5	15.0	1145.3	8.8	22.0	23.3	1145.3	27.8	96.0
Trial4_805	122	NA()	1.6	5.6537	0.9	11.9323	1.9	0.4895	1.6	0.88	2568.5	9.8	37.0	38.2	2599.1	5.7	19.3	20.1	2623.8	6.6	18.1	19.2	2623.8	14.9	97.9
Trial4_806	41	NA()	1.1	13.3537	2.4	1.8804	4.1	0.1822	3.4	0.82	1079.0	9.5	19.3	19.8	1074.2	8.5	15.6	16.0	1065.5	16.9	26.5	27.6	1065.5	48.0	101.3
Trial4_807	121	NA()	2.4	13.6441	1.5	1.7451	3.2	0.1728	2.8	0.88	1027.3	7.6	17.7	18.2	1025.3	6.1	14.1	14.6	1022.0	10.2	22.9	24.2	1022.0	30.5	100.5
Trial4_808	108	NA()	2.5	13.5521	1.6	1.7306	3.3	0.1702	2.9	0.88	1013.1	7.7	17.6	18.1	1020.0	6.2	14.1	14.6	1035.7	10.4	23.0	24.3	1035.7	32.2	97.8
Trial4_809	56	NA()	0.6	5.4027	1.0	13.5908	2.6	0.5328	2.4	0.93	2753.2	12.2	39.7	40.9	2721.6	7.2	20.0	20.7	2699.0	8.8	18.9	19.9	2699.0	15.9	102.0
Trial4_810	88	NA()	1.4	21.7091	3.3	0.2981	4.9	0.0470	3.5	0.73	295.8	3.0	5.7	5.8	264.9	4.1	6.1	6.3	0.9	34.2	42.1	43.1	295.8	10.2	NA
Trial4_811	46	NA()	1.1	6.1258	1.1	10.9488	2.9	0.4867	2.7	0.93	2556.2	13.1	37.9	39.0	2518.8	7.7	19.9	20.6	2489.5	9.3	19.5	20.5	2489.5	18.5	102.7
Trial4_812	241	NA()	2.8	19.0516	2.4	0.3094	4.0	0.0428	3.2	0.80	270.0	1.8	4.8	4.9	273.7	2.5	5.3	5.5	306.8	18.2	29.4	30.7	270.0	8.5	NA
Trial4_813	86	NA()	1.7	15.6664	2.3	0.9924	4.0	0.1128	3.3	0.82	689.1	6.2	12.6	13.0	699.9	6.3	11.8	12.1	736.1	17.2	27.5	28.6	689.1	21.5	93.6
Trial4_814	151	NA()	0.8	17.5281	2.3	0.5670	3.9	0.0721	3.2	0.82	448.9	3.7	8.2	8.4	456.1	4.2	8.4	8.6	493.5	17.0	28.1	29.3	448.9	13.9	91.0
Trial4_815	45	NA()	4.3	9.2617	1.5	4.6963	3.4	0.3156	3.1	0.90	1768.2	12.5	28.9	29.7	1766.6	8.3	18.4	19.1	1765.4	10.4	21.2	22.4	1765.4	27.7	100.2
Trial4_816	220	NA()	2.3	12.7186	1.0	2.1194	2.3	0.1956	2.1	0.89	1151.6	6.5	18.9	19.5	1155.2	4.8	14.4	14.9	1162.7	6.5	21.1	22.5	1162.7	20.7	99.0
Trial4_817	293	NA()	1.8	10.0264	0.8	3.7364	1.5	0.2718	1.3	0.84	1550.1	5.9	24.0	24.7	1579.2	4.0	16.2	16.9	1619.2	4.5	19.4	20.7	1619.2	15.6	95.7
Trial4_818	20	NA()	0.4	15.2723	3.8	0.8009	5.0	0.0888	3.2	0.65	548.2	6.5	11.0	11.2	597.4	11.9	14.9	15.1	789.7	49.1	53.5	54.1	548.2	17.0	69.4
Trial4_819	295	NA()	1.6	18.1061	1.8	0.4595	3.4	0.0604	2.9	0.84	377.8	2.0	6.5	6.7	383.9	2.6	6.8	7.1	421.5	13.6	26.4	27.7	377.8	10.5	NA
Trial4_820	78	NA()	1.2	17.0245	2.6	0.8162	4.2	0.1008	3.4	0.79	619.3	5.9	11.6	11.9	605.9	6.1	10.9	11.2	557.4	19.4	29.4	30.5	619.3	19.9	111.1
Trial4_821	171	NA()	4.3	19.8290	3.4	0.2083	4.9	0.0300	3.5	0.71	190.3	1.6	3.6	3.7	192.1	2.7	4.4	4.5	214.9	29.0	37.3	38.3	190.3	6.5	NA
Trial4_822	269	NA()	2.9	5.4022	0.8	12.1255	1.4	0.4753	1.1	0.78	2506.7	7.1	35.7	36.9	2614.2	4.1	18.9	19.7	2699.2	4.5	17.3	18.4	2699.2	14.0	92.9
Trial4_823	65	NA()	1.7	10.3300	1.4	3.7420	3.2	0.2805	2.9	0.90	1593.8	11.6	26.5	27.2	1580.4	7.7	17.6	18.2	1563.4	9.6	21.3	22.5	1563.4	26.6	101.9
Trial4_824	85	NA()	1.0	8.9405	1.1	5.2110	2.8	0.3380	2.5	0.92	1877.3	12.6	30.2	31.1	1854.4	7.7	18.5	19.1	1829.7	8.6	20.3	21.4	1829.7	20.2	102.6
Trial4_825	49	NA()	0.6	18.2298	3.5	0.5164	4.8	0.0683	3.2	0.68	425.9	5.0	8.5	8.7	422.7	7.6	10.2	10.4	406.3	41.4	47.3	48.0	425.9	13.3	104.8
Trial4_826	83	NA()	1.6	12.2565	1.5	2.4184	3.3	0.2151	2.9	0.89	1255.8	10.5	21.9	22.5	1248.1	7.6	16.0	16.6	1235.7	10.2	22.3	23.6	1235.7	29.9	101.6
Trial4_827	335	NA()	1.6	13.1138	0.9	1.9469	1.8	0.1852	1.6	0.87	1095.6	4.7	17.6	18.2	1097.4	3.6	13.7	14.2	1101.8	5.4	21.0	22.3	1101.8	17.8	99.4
Trial4_828	154	NA()	1.0	13.5100	1.3	1.7334	2.9	0.1699	2.6	0.89	1011.7	7.8	17.6	18.1	1021.0	6.0	14.0	14.5	1042.0	8.6	22.2	23.5	1042.0	26.9	97.1
Trial4_829	92	NA()	2.6	9.2782	1.1	4.5945	2.8	0.3093	2.5	0.91	1737.3	12.1	28.4	29.1	1748.3	7.6	18.1	18.7	1762.2	8.2	20.3	21.5	1762.2	20.5	98.6
Trial4_830	92	NA()	2.1	13.3833	1.7	1.7279	3.5	0.1678	3.0	0.87	999.9	8.8	17.9	18.4	1019.0	7.1	14.5	15.0	1061.0	11.5	23.4	24.6	1061.0	34.0	94.2
Trial4_831	125	NA()	0.6	18.1556	2.5	0.5354	4.2	0.0705	3.3	0.80	439.4	4.2	8.3	8.5	435.4	4.8	8.4	8.7	415.4	20.2	30.4	31.5	439.4	14.1	105.8
Trial4_832	39	NA()	0.7	5.3821	1.1	12.8194	3.0	0.5006	2.8	0.93	2616.5	13.9	38.8	40.0	2666.5	8.3	20.3	21.1	2705.3	10.0	19.5	20.5	2705.3	18.5	96.7
Trial4_833	34	NA()	1.2	13.9080	2.7	1.7136	4.4	0.1729	3.5	0.79	1028.3	10.3	19.0	19.5	1013.7	9.6	15.8	16.3	983.2	21.0	29.5	30.5	983.2	55.5	104.6
Trial4_834	249	NA()	1.5	19.2101	2.9	0.2123	4.4	0.0296	3.3	0.75	188.0	1.4	3.4	3.5	195.5	2.6	4.3	4.5	287.9	27.7	36.1	37.2	188.0	6.2	NA
Trial4_835	77	NA()	0.6	5.3226	1.0	13.6591	2.3	0.5275	2.1	0.91	2731.0	12.3	39.5	40.7	2726.4	7.4	20.0	20.8	2723.7	9.0	19.0	20.0	2723.7	16.0	100.3
Trial4_836	188	NA()	1.2	19.2540	2.7	0.3005	4.3	0.0420	3.3	0.77	265.1	2.3	4.9	5.1	266.8	3.1	5.6	5.7	282.7	22.6	32.4	33.6	265.1	8.7	NA
Trial4_837	147	NA()	1.5	18.2916	2.6	0.4382	4.2	0.0582	3.3	0.79	364.4	3.4	6.9	7.0	369.0	4.1	7.3	7.5	398.7	20.3	30.5	31.6	364.4	11.8	NA
Trial4_838	55	NA()	0.8	10.7649	1.6	3.2485	3.5	0.2537	3.1	0.88	1457.7	12.0	25.0	25.7	1468.8	8.5	17.5	18.0	1485.7	11.1	22.2	23.3	1485.7	30.9	98.1
Trial4_839	105	NA()	2.2	17.2328	2.5	0.6631	4.2	0.0829	3.3	0.80	513.5	5.1	9.7	10.0	516.5	5.4	9.6	9.9	530.9	19.1	29.3	30.5	513.5	16.5	96.7
Trial4_840	119	NA()	1.2	5.5028	0.9	12.2320	1.9	0.4884	1.7	0.87	2563.7	11.4	37.5	38.6	2622.4	6.8	19.7	20.4	2668.7	7.9	18.6	19.6	2668.7	15.4	96.1
Trial4_841	23	NA()	1.4	5.3270	1.4	13.0520	3.4	0.5045	3.1	0.91	2633.1	14.9	39.4	40.5	2683.4	9.2	20.7	21.4	2722.3	11.3	20.2	21.1	2722.3	23.0	96.7
Trial4_842	15	NA()	1.2	11.6630	3.4	2.5568	4.6	0.2164	3.2	0.69	1262.7	12.8	23.2	23.7	1288.4	12.7	19.2	19.7	1332.3	26.0	32.6	33.4	1332.3	32.6	94.8
Trial4_843	115	NA()	2.0	9.9999	1.1	3.7370	2.7	0.2712	2.4	0.91	1546.6	10.8	25.6	26.3	1579.3	7.1	17.3	17.9	1624.1	7.7	20.4	21.6	1624.1	20.4	95.2
Trial4_844	9	NA()	1.6	13.3050	3.9	1.7833	5.1	0.1722	3.3	0.64	1024.0	12.0	19.9	20.4	1039.4	18.1	22.1	22.5	1072.8	49.7	53.7	54.2	1072.8	53.7	95.5
Trial4_845	17	NA()	1.7	12.5140	3.4	2.2248	4.7	0.2020	3.2	0.68	1186.2	12.1	21.9	22.4	1188.9	12.4	18.5	19.0	1194.8	26.9	33.5	34.4	1194.8	33.5	99.3
Trial4_846	63	NA()	1.1	13.7918	2.1	1.5826	3.9	0.1584	3.3	0.84	947.8	9.0	17.4	17.8	963.4	7.8	14.5	14.9	1000.2	14.5	25.2	26.4	1000.2	25.2	94.8
Trial4_847	144	NA()	1.0	13.6161	1.4	1.7580	3.0	0.1737	2.7	0.89	1032.4	8.0	17.9	18.4	1030.1	6.1	14.1	14.6	1026.2	9.0	22.4	23.7	1026.2	22.4	100.6
Trial4_848	515	NA()	13.9	13.1510	0.8	1.7791	1.3	0.1698	1.0	0.78	1010.9	1.3	15.8	16.4	1037.9	1.6	12.9	13.4	1096.1	4.2	20.7	22.1	1096.1	20.7	92.2
Trial4_849	135	NA()	2.7	12.7335	1.3	2.1401	2.9	0.1977	2.6	0.90	1163.1	8.8	20.0	20.5	1161.9	6.4	15.1	15.6	1160.4	8.6	21.9	23.1	1160.4	21.9	100.2
Trial4_850	53	NA()	1.2	12.0660	1.9	2.4046	3.7	0.2105	3.2	0.86	1231.6	11.0	21.9	22.4	1244.0	8.6	16.5	17.0	1266.3	13.3	23.8	25.0	1266.3	23.8	97.3
Trial4_851	181	NA()	2.5	12.7607	1.1	2.0685	2.6	0.1915	2.3	0.90	1129.6	7.7	19.1	19.6	1138.4	5.7	14.6	15.2	1156.1	7.4	21.5	22.8	1156.1	21.5	97.7
Trial4_852	79	NA()	1.4	13.2234	1.8	1.7921	3.6	0.1719	3.1	0.87	1022.8	9.2	18.4	18.9	1042.6	7.4	14.8	15.3	1085.1	12.1	23.6	24.8	1085.1	23.6	94.3
Trial4_853	32	NA()	2.9	13.7148	2.8	1.7304	4.5	0.1722	3.5	0.78	1024.2	10.4	19.0	19.5	1019.9	9.8	16.0	16.4	1011.5	21.2	29.5	30.5	1011.5	29.5	101.3
Trial4_854	12	NA()	2.8	10.4745	3.4	3.1412	4.7	0.2387	3.2	0.69	1380.1	13.9	25.1	25.7	1442.8	13.5	20.3	20.8	1537.3	25.5	31.8	32.6	1537.3	31.8	89.8
Trial4_855	168	NA()	1.0	17.6133	2.2	0.5327	3.9	0.0681	3.2	0.82	424.6	3.7	7.8	8.1	433.6	4.2	8.1	8.3	482.8	16.9	28.0	29.3	424.6	7.8	87.9
Trial4_856	182	NA()	1.6	12.4674	1.1	2.1730	2.6	0.1966	2.3	0.90	1156.9	7.8	19.5	20.0	1172.5	5.7	14.8	15.4	1202.1	7.3	21.3	22.6	1202.1	21.3	96.2
Trial4_857	52	NA()	0.6	8.9451	1.4	5.1688	3.3	0.3355	2.9	0.90	1864.9	13.5	30.4	31.3	1847.5	8.5	18.8	19.4	1828.7	10.1	21.0	22.1	1828.7	21.0	102.0
Trial4_858	16	NA()	1.5	12.2421	3.5	2.3793	4.7	0.2114	3.2	0.68	1236.0	12.5	22.7	23.3	1236.4	12.7	18.9	19.4	1238.0	27.0	33.6	34.4	1238.0	33.6	99.8
Trial4_859	107	NA()	1.8	17.3341	2.6	0.6402	4.2	0.0805	3.4	0.79	499.2	4.9	9.5	9.7	502.4	5.6	9.6	9.9	518.0	21.5	30.9	32.1	499.2	9.5	96.4
Trial4_860	14	NA()	2.7	12.2838	3.4	2.1157	4.7	0.1886	3.2	0.68	1113.6	11.9	21.0	21.4	1153.9	14.1	19.5	20.0	1231.3	32.9	38.4	39.2	1231.3	38.4	90.4
Trial4_861	89	NA()	0.9	10.6591	1.4	2.9642	3.2	0.2293	2.9	0.90	1330.6	10.6	22.9	23.5	1398.5	7.6	16.8	17.3	1504.4	9.0	21.2	22.4	1504.4	21.2	88.4
Trial4_862	70	NA()	0.6	5.1797	1.0	14.4028	2.4	0.5413	2.2	0.91	2788.9	12.8	40.2	41.5	2776.6	7.5	20.1	20.9	2768.4	9.1	18.9	20.0	2768.4	18.9	100.7
Trial4_863	66	NA()	2.3	13.7085	2.0	1.7207	3.8	0.1712	3.2	0.85	1018.5	9.4	18.5	18.9	1016.3	7.7	14.8	15.3	1012.5	13.5	24.6	25.8	1012.5	24.6	100.6
Trial4_864	49	NA()	0.5	5.3807	1.0	13.0802	2.8	0.5107	2.6	0.93	2659.5	13.8	39.2	40.4	2685.5	8.1	20.2	21.0	2705.8	9.5	19.2	20.2	2705.8	19.2	98.3
Trial4_865	80	NA()	1.5	8.1668	1.1	6.4445	2.7	0.3819	2.5	0.92	2085.1	13.0	32.7	33.7	2038.4	7.7	18.9	19.6	1992.2	8.7	20.0	21.1	1992.2	20.0	104.7
Trial4_866	41	NA()	1.2	12.4150	2.2	2.1944	4.0	0.1977	3.4	0.83	1162.9	11.0	21.1	21.6	1179.3	9.0	16.4	16.9	1210.4	15.2	25.1	26.2	1210.4	25.1	96.1
Trial4_867	65	NA()	0.6	3.2689	1.0	30.1187	2.2	0.7144	1.9	0.89	3475.0	10.3	46.4	47.9	3490.7	7.1	20.6	21.4	3500.4	9.5	18.3	19.2	3500.4	18.3	99.3
Trial4_868	598	NA()	2.4	12.2388	0.8	2.4073	1.3	0.2138	1.0	0.78	1248.9	0.9	19.2	19.8	1244.8	1.4	14.2	14.8	1238.5	3.3	20.2	21.5	1238.5	20.2	100.8
Trial4_869	350	NA()	2.1	9.0658	0.8	4.4309	1.4	0.2915	1.1	0.79	1648.9	5.5	25.1	25.9	1718.1	3.6	16.7	17.4	1804.4	3.9	18.8	20.1	1804.4	18.8	91.4
Trial4_870	82	NA()	0.8	13.8455	1.8	1.7231	3.6	0.1731	3.1	0.86	1029.2	9.2	18.5	18.9	1017.2	7.4	14.6	15.1	992.3	12.6	24.2	25.4	992.3	24.2	103.7
Trial4_871	89	NA()	1.4	10.9798	1.3	3.1625	3.1	0.2519	2.8	0.90	1448.5	11.1	24.5	25.2	1448.0	7.5	16.9	17.5	1448.2	9.0	21.3	22.5	1448.2	21.3	100.0
Trial4_872	156	NA()	2.5	13.7220	1.3	1.7249	2.9	0.1717	2.6	0.89	1021.7	7.7	17.7	18.2	1017.9	5.9	13.9	14.4	1010.5	8.6	22.3	23.6	1010.5	22.3	101.1
Trial4_873	67	NA()	1.9	9.1137	1.3	4.8467	3.1	0.3205	2.8	0.91	1792.2	12.8	29.3	30.1	1793.0	8.1	18.4	19.1	1794.8	9.2	20.6	21.8	1794.8	20.6	99.9
Trial4_874	153	NA()	1.7	13.1807	1.3	1.8140	2.9	0.1735	2.6	0.89	1031.3	7.8	17.8	18.3	1050.5	6.0	14.2	14.7	1091.6	8.4	21.9	23.2	1091.6	21.9	94.5
Trial4_875	39	NA()	1.2	8.7113	1.6	5.1477	3.5	0.3254	3.1	0.89	1815.9	13.8	30.0	30.8	1844.0	9.0	19.0	19.6	1876.6	10.9	21.3	22.3	1876.6	21.3	96.8
Trial4_876	32	NA()	1.2	5.7924	1.3	11.5631	3.2	0.4860	3.0	0.92	2553.3	14.6	38.4	39.5	2569.7	8.7	20.3	21.1	2583.4	10.4	19.9	20.9	2583.4	19.9	98.8
Trial4_877	56	NA()	0.8	13.1842	2.1	1.9327	3.9	0.1849	3.3	0.85	1093.6	10.1	19.7	20.2	1092.5	8.2	15.5	16.0	1091.1	13.8	24.5	25.7	1091.1	24.5	100.2
Trial4_878	14	NA()	1.5	13.3635	3.5	1.8872	4.8	0.1830	3.2	0.68	1083.3	11.8	20.5	21.0	1076.6	14.5	19.5	19.9	1064.0	37.1	42.3	43.0	1064.0	42.3	101.8
Trial4_879	163	NA()	1.1	19.3812	3.6	0.2047	5.0	0.0288	3.5	0.70	183.0	1.7	3.5	3.6	189.1	3.0	4.6	4.7	267.5	33.7	40.9	41.9	183.0	3.5	NA
Trial4_880	259	NA()	1.6	11.0620	0.8	3.0875	1.8	0.2478	1.6	0.88	1427.2	7.1	22.7	23.4	1429.6	4.8	15.8	16.4	1434.0	5.4	20.1	21.4	1434.0	20.1	99.5
Trial4_881	94	NA()	1.3	10.9743	1.3	3.0876	3.0	0.2459	2.7	0.90	1417.1	10.8	24.0	24.7	1429.6	7.5	16.8	17.4	1449.1	8.9	21.3	22.5	1449.1	21.3	97.8
Trial4_882	83	NA()	1.1	11.2628	1.4	3.1058	3.1	0.2538	2.8	0.90	1458.1	11.3	24.7	25.4	1434.1	7.7	17.0	17.5	1399.6	9.7	21.7	22.9	1399.6	21.7	104.2
Trial4_883	99	NA()	1.8	19.5627	3.5	0.2812	5.0	0.0399	3.6	0.72	252.3	2.7	5.0	5.1	251.6	4.7	6.4	6.5	246.1	41.1	47.3	48.1	252.3	5.0	NA
Trial4_884	38	NA()	1.6	13.9893	2.6	1.6506	4.4	0.1675	3.5	0.80	998.6	9.9	18.5	18.9	989.8	8.9	15.3	15.8	971.3	18.8	27.9	29.0	971.3	27.9	102.8
Trial4_885	57	NA()	3.6	12.9137	2.0	1.9794	3.8	0.1855	3.3	0.85	1096.8	10.1	19.8	20.3	1108.5	8.3	15.6	16.1	1132.5	13.9	24.5	25.7	1132.5	24.5	96.8
Trial4_886	294	NA()	0.8	16.6664	1.4	0.8243	2.8	0.0997	2.5	0.87	612.6	3.5	10.5	10.8	610.5	3.4	9.6	10.0	603.6	9.2	23.8	25.2	612.6	10.5	101.5
Trial4_887	182	NA()	2.5	17.8143	2.1	0.5408	3.8	0.0699	3.1	0.83	435.6	3.7	8.0	8.2	439.0	3.9	8.0	8.3	457.7	15.3	27.2	28.5	435.6	8.0	95.2
Trial4_888	55	NA()	2.3	9.4539	1.4	4.3891	3.3	0.3011	3.0	0.90	1696.7	12.9	28.3	29.0	1710.3	8.5	18.3	19.0	1727.8	10.1	21.2	22.3	1727.8	21.2	98.2
Trial4_889	37	NA()	0.9	13.6140	2.5	1.8556	4.3	0.1833	3.4	0.81	1085.0	10.6	19.9	20.4	1065.4	9.1	15.8	16.3	1026.5	17.8	27.1	28.2	1026.5	27.1	105.7
Trial4_890	58	NA()	1.3	7.6122	3.1	0.7980	4.5	0.0441	3.2	0.73	278.1	3.3	5.6	5.8	595.7	7.8	11.8	12.1	2116.3	21.6	28.0	28.8	278.1	5.6	NA
Trial4_891	276	NA()	3.1	9.3755	0.8	4.5331	1.4	0.3084	1.2	0.81	1732.7	7.1	26.6	27.4	1737.1	4.5	17.0	17.7	1743.1	4.9	19.2	20.4	1743.1	19.2	99.4
Trial4_892	42	NA()	2.1	13.8758	2.5	1.6599	4.2	0.1671	3.4	0.81	996.2	9.7	18.3	18.8	993.3	8.6	15.2	15.6	987.8	17.6	27.1	28.2	987.8	27.1	100.8
Trial4_893	116	NA()	1.9	19.2280	3.4	0.1927	4.6	0.0269	3.2	0.69	171.1	1.8	3.4	3.5	179.0	3.0	4.4	4.5	285.7	34.3	41.4	42.3	171.1	3.4	NA
Trial4_894	20	NA()	1.5	13.0665	3.3	1.8942	4.9	0.1796	3.6	0.73	1064.8	10.9	19.8	20.3	1079.1	11.6	17.5	17.9	1109.0	27.0	33.7	34.6	1109.0	33.7	96.0
Trial4_895	69	NA()	1.0	12.7610	1.8	2.0623	3.5	0.1910	3.1	0.87	1126.6	8.7	19.5	20.0	1136.4	7.0	15.1	15.7	1156.1	11.5	23.2	24.4	1156.1	23.2	97.4
Trial4_896	86	NA()	0.7	17.2174	2.6	0.7260	4.3	0.0907	3.4	0.79	559.7	4.9	10.3	10.5	554.2	5.5	10.0	10.4	532.8	19.9	29.8	31.0	559.7	10.3	105.0
Trial4_897	106	NA()	1.7	17.3094	2.5	0.6577	4.1	0.0826	3.3	0.80	511.7	4.3	9.3	9.6	513.2	5.0	9.4	9.7	521.1	19.5	29.5	30.7	511.7	9.3	98.2
Trial4_898	94	NA()	2.7	12.9802	1.6	1.9881	3.3	0.1873	2.9	0.88	1106.5	8.0	18.9	19.4	1111.5	6.4	14.8	15.3	1122.2	10.6	22.8	24.1	1122.2	22.8	98.6
Trial4_899	139	NA()	3.0	13.0523	1.4	1.8627	3.0	0.1764	2.6	0.89	1047.3	6.9	17.7	18.2	1067.9	5.5	14.1	14.6	1111.2	8.7	22.0	23.3	1111.2	22.0	94.3
Trial4_900	117	NA()	2.4	13.7172	1.5	1.7358	3.2	0.1728	2.8	0.88	1027.4	7.2	17.5	18.0	1021.9	5.8	13.9	14.4	1011.2	10.0	22.9	24.1	1011.2	22.9	101.6
Trial4_901	47	NA()	0.5	19.4806	4.6	0.1879	5.6	0.0266	3.2	0.57	169.0	1.9	3.4	3.4	174.9	6.0	6.7	6.8	255.8	81.4	84.7	85.2	169.0	3.4	NA
Trial4_902	299	NA()	2.8	13.0584	0.9	1.9632	1.9	0.1860	1.7	0.88	1099.8	3.1	17.3	17.9	1103.0	2.8	13.5	14.1	1110.2	5.5	21.0	22.3	1110.2	21.0	99.1
Trial4_903	236	NA()	1.4	12.9253	1.1	1.8720	2.3	0.1756	2.0	0.89	1042.7	4.2	16.8	17.3	1071.3	3.6	13.5	14.0	1130.7	6.3	21.2	22.5	1130.7	21.2	92.2
Trial4_904	150	NA()	1.7	5.2866	0.9	13.6116	1.5	0.5221	1.2	0.81	2708.2	7.7	38.1	39.3	2723.1	4.6	19.2	20.0	2734.8	5.6	17.6	18.7	2734.8	17.6	99.0
Trial4_905	56	NA()	1.7	4.5850	0.9	18.1181	2.5	0.6028	2.3	0.93	3040.9	9.9	42.1	43.4	2996.0	6.2	19.9	20.7	2966.7	7.9	18.2	19.2	2966.7	18.2	102.5
Trial4_906	40	NA()	0.7	3.7644	1.0	23.8912	2.7	0.6526	2.5	0.94	3238.2	10.0	44.1	45.5	3264.0	6.4	20.2	21.0	3280.5	8.2	17.9	18.9	3280.5	17.9	98.7
Trial4_907	49	NA()	1.3	11.1928	1.7	3.0196	3.6	0.2452	3.1	0.87	1413.9	9.5	23.4	24.1	1412.6	7.2	16.7	17.3	1411.5	11.2	22.4	23.6	1411.5	22.4	100.2
Trial4_908	63	NA()	0.6	15.8876	2.8	0.8862	4.4	0.1022	3.4	0.77	627.1	5.2	11.3	11.6	644.3	6.5	11.4	11.7	706.3	22.4	31.1	32.1	627.1	11.3	88.8
Trial4_909	70	NA()	0.6	20.6321	4.4	0.1743	5.5	0.0261	3.2	0.58	166.1	1.7	3.3	3.3	163.2	5.2	6.0	6.0	122.2	77.2	80.8	81.3	166.1	3.3	NA
Trial4_910	412	NA()	2.0	13.5266	0.8	1.7513	1.6	0.1719	1.3	0.85	1022.5	1.6	16.0	16.6	1027.7	1.9	12.9	13.4	1039.5	4.8	21.1	22.4	1039.5	21.1	98.4
Trial4_911	135	NA()	0.9	19.9292	3.3	0.2851	4.8	0.0412	3.4	0.72	260.4	2.1	4.8	4.9	254.7	3.5	5.6	5.8	203.2	30.3	38.3	39.3	260.4	4.8	NA
Trial4_912	52	NA()	1.1	13.8369	2.2	1.6926	4.0	0.1699	3.3	0.83	1011.8	8.2	17.8	18.3	1005.7	7.4	14.6	15.0	993.6	15.3	25.6	26.8	993.6	25.6	101.8
Trial4_913	32	NA()	0.5	5.5288	1.2	12.8102	3.2	0.5139	2.9	0.93	2673.3	12.5	39.0	40.1	2665.8	7.5	20.0	20.7	2660.9	9.3	19.2	20.2	2660.9	19.2	100.5
Trial4_914	326	NA()	15.8	13.2627	0.9	1.8542	1.9	0.1784	1.6	0.87	1058.4	3.1	16.7	17.3	1064.9	2.8	13.3	13.8	1079.2	5.4	21.0	22.4	1079.2	21.0	98.1
Trial4_915	79	NA()	1.6	19.0204	3.3	0.5003	4.8	0.0690	3.5	0.72	430.4	3.9	8.0	8.2	411.9	5.5	8.6	8.9	310.5	30.1	38.0	38.9	430.4	8.0	138.6
Trial4_916	69	NA()	2.6	13.0526	1.8	2.0334	3.6	0.1926	3.1	0.86	1135.3	8.3	19.4	20.0	1126.7	6.8	15.0	15.5	1111.1	12.0	23.5	24.7	1111.1	23.5	102.2
Trial4_917	304	NA()	2.4	13.4928	0.9	1.8284	2.0	0.1790	1.7	0.87	1061.6	3.4	16.9	17.4	1055.7	3.0	13.2	13.8	1044.5	5.7	21.2	22.6	1044.5	21.2	101.6
Trial4_918	58	NA()	1.6	12.8332	1.9	2.0717	3.7	0.1929	3.2	0.85	1137.1	8.6	19.6	20.1	1139.5	7.2	15.3	15.8	1144.9	13.0	23.9	25.1	1144.9	23.9	99.3
Trial4_919	241	NA()	1.6	20.8015	2.5	0.2849	4.0	0.0430	3.2	0.78	271.4	1.5	4.7	4.9	254.6	2.4	5.0	5.2	102.9	21.0	31.9	33.1	271.4	4.7	NA
Trial4_920	77	NA()	1.4	18.0664	2.9	0.6565	4.5	0.0861	3.4	0.76	532.2	4.6	9.8	10.0	512.5	5.7	9.7	10.0	426.4	24.3	33.2	34.3	532.2	9.8	124.8
Trial4_921	28	NA()	1.8	11.9993	2.5	2.5176	4.2	0.2192	3.4	0.81	1277.6	10.5	22.2	22.8	1277.1	9.4	17.1	17.7	1277.1	18.1	26.8	27.8	1277.1	26.8	100.0
Trial4_922	168	NA()	2.5	5.8940	0.9	11.4566	1.5	0.4900	1.2	0.81	2570.5	8.4	36.7	37.9	2561.0	4.9	19.0	19.8	2554.3	5.7	17.9	19.0	2554.3	17.9	100.6
Trial4_923	62	NA()	1.0	16.0294	3.0	0.7282	4.6	0.0847	3.5	0.75	524.1	4.8	9.8	10.0	555.5	6.1	10.4	10.7	687.4	22.6	31.3	32.3	524.1	9.8	76.2
Trial4_924	48	NA()	2.0	13.7590	2.3	1.6661	4.1	0.1663	3.3	0.82	991.9	8.3	17.6	18.0	995.7	7.6	14.6	15.1	1005.0	16.0	26.1	27.2	1005.0	26.1	98.7
Trial4_925	81	NA()	2.4	10.9337	1.4	3.2158	3.1	0.2551	2.8	0.90	1464.9	9.6	24.1	24.8	1461.0	6.7	16.7	17.3	1456.2	9.0	21.3	22.5	1456.2	21.3	100.6

Trial4_926	198	NA()	0.7	20.2556	3.3	0.1899	4.8	0.0279	3.4	0.72	177.5	1.4	3.3	3.4	176.6	2.4	4.0	4.1	165.4	29.7	38.0	39.0	177.5	3.3	NA
Trial4_927	10	NA()	2.3	15.0200	5.5	1.2138	6.4	0.1323	3.3	0.51	800.9	9.5	15.9	16.2	806.9	26.6	28.8	29.0	824.6	96.2	98.5	98.9	800.9	15.9	97.1
Trial4_928	285	NA()	1.9	9.1148	0.8	4.8401	1.4	0.3201	1.1	0.79	1790.2	6.1	27.1	27.9	1791.9	3.9	17.0	17.7	1794.6	4.3	19.0	20.2	1794.6	19.0	99.8
Trial4_929	75	NA()	0.8	18.1559	3.3	0.5398	4.8	0.0711	3.5	0.73	442.9	4.4	8.4	8.6	438.3	5.8	9.1	9.3	415.4	28.2	36.2	37.2	442.9	8.4	106.6
Trial4_930	29	NA()	0.3	5.4036	1.3	12.5921	3.3	0.4937	3.0	0.92	2586.7	13.7	38.4	39.6	2649.6	8.2	20.3	21.0	2698.8	9.9	19.4	20.4	2698.8	19.4	95.8
Trial4_931	52	NA()	0.7	5.5887	1.0	12.6576	2.7	0.5133	2.5	0.93	2670.6	12.4	38.9	40.1	2654.5	7.4	19.9	20.7	2643.0	9.0	19.1	20.1	2643.0	19.1	101.0
Trial4_932	68	NA()	2.2	13.4318	1.9	1.8287	3.7	0.1782	3.2	0.86	1057.3	9.5	19.0	19.5	1055.8	7.7	15.0	15.5	1053.7	13.0	24.2	25.4	1053.7	24.2	100.3
Trial4_933	202	NA()	1.5	13.7042	1.2	1.8008	2.6	0.1791	2.3	0.89	1061.8	6.7	17.8	18.3	1045.7	5.1	13.8	14.3	1013.1	7.5	21.9	23.2	1013.1	21.9	104.8
Trial4_934	124	NA()	2.2	19.9099	3.6	0.2586	5.0	0.0374	3.5	0.70	236.4	2.4	4.6	4.7	233.5	3.7	5.5	5.7	205.5	34.2	41.5	42.5	236.4	4.6	NA
Trial4_935	73	NA()	2.1	11.6596	1.7	2.2214	3.5	0.1879	3.1	0.88	1110.2	9.5	19.7	20.2	1187.8	7.6	15.8	16.3	1332.9	11.0	22.5	23.7	1332.9	22.5	83.3
Trial4_936	69	NA()	1.1	13.0655	1.8	2.0206	3.6	0.1916	3.1	0.87	1129.8	9.7	20.0	20.5	1122.5	7.6	15.4	15.9	1109.1	12.2	23.7	24.9	1109.1	23.7	101.9
Trial4_937	46	NA()	0.9	18.4216	3.6	0.5046	4.8	0.0675	3.2	0.67	420.8	4.8	8.4	8.6	414.8	7.9	10.4	10.6	382.8	45.2	50.6	51.3	420.8	8.4	109.9
Trial4_938	60	NA()	1.8	12.2306	1.8	2.4089	3.6	0.2138	3.1	0.87	1248.9	10.4	21.8	22.3	1245.3	7.9	16.2	16.7	1239.8	12.0	23.2	24.4	1239.8	23.2	100.7
Trial4_939	26	NA()	0.8	13.3166	3.0	1.8398	4.6	0.1778	3.5	0.77	1054.8	10.3	19.4	19.9	1059.8	9.9	16.3	16.7	1071.0	21.2	29.4	30.4	1071.0	29.4	98.5
Trial4_940	45	NA()	1.8	17.4726	3.5	0.5043	4.7	0.0639	3.2	0.68	399.5	4.5	8.0	8.1	414.6	7.1	9.8	10.0	500.5	38.1	44.2	45.0	399.5	8.0	NA
Trial4_941	26	NA()	3.4	13.0552	3.0	1.8689	4.6	0.1770	3.5	0.77	1050.8	10.3	19.3	19.8	1070.1	10.0	16.4	16.9	1110.7	21.7	29.7	30.6	1110.7	29.7	94.6
Trial4_942	114	NA()	2.9	11.4998	1.3	2.7036	2.9	0.2256	2.6	0.90	1311.4	9.4	22.1	22.7	1329.4	6.7	16.0	16.6	1359.5	8.2	21.2	22.4	1359.5	21.2	96.5
Trial4_943	66	NA()	1.2	11.5120	1.6	2.6913	3.4	0.2248	3.0	0.88	1307.2	10.7	22.6	23.2	1326.1	7.9	16.6	17.1	1357.5	11.1	22.5	23.7	1357.5	22.5	96.3
Trial4_944	111	NA()	1.4	11.3639	1.2	2.8752	2.9	0.2371	2.6	0.90	1371.5	9.8	23.0	23.7	1375.4	6.8	16.3	16.9	1382.4	8.4	21.2	22.4	1382.4	21.2	99.2
Trial4_945	167	NA()	0.3	13.4802	1.2	1.8453	2.7	0.1805	2.5	0.89	1069.6	7.4	18.2	18.7	1061.7	5.6	14.1	14.6	1046.5	8.1	22.0	23.3	1046.5	22.0	102.2
Trial4_946	58	NA()	1.7	13.2840	2.1	1.6891	3.9	0.1628	3.3	0.84	972.4	9.0	17.7	18.1	1004.4	7.8	14.8	15.3	1076.0	14.5	25.0	26.2	1076.0	25.0	90.4
Trial4_947	626	NA()	1.6	20.7982	1.6	0.2746	3.0	0.0414	2.5	0.84	261.8	0.5	4.4	4.5	246.4	1.2	4.5	4.7	103.2	12.5	27.0	28.5	261.8	4.4	NA
Trial4_948	14	NA()	1.6	4.4674	1.5	16.8816	3.7	0.5472	3.3	0.91	2813.6	15.4	41.4	42.6	2928.1	9.5	21.1	21.8	3008.5	11.6	20.0	20.9	3008.5	20.0	93.5
Trial4_949	124	NA()	1.9	13.5833	1.5	1.8108	3.1	0.1785	2.8	0.89	1058.6	8.1	18.4	18.8	1049.4	6.3	14.3	14.8	1031.1	9.6	22.7	23.9	1031.1	22.7	102.7
Trial4_950	158	NA()	2.1	9.0287	0.9	4.2249	2.2	0.2768	2.0	0.92	1575.1	9.4	25.3	26.1	1678.9	6.2	17.3	17.9	1811.9	6.3	19.5	20.7	1811.9	19.5	86.9
Trial4_951	147	NA()	2.2	4.2964	0.9	18.4842	1.5	0.5762	1.2	0.78	2933.4	9.5	40.8	42.1	3015.3	5.8	19.8	20.6	3071.1	7.2	17.7	18.7	3071.1	17.7	95.5
Trial4_952	46	NA()	0.5	15.6997	3.5	0.7155	5.0	0.0815	3.6	0.71	505.1	5.5	9.9	10.1	548.0	7.5	11.2	11.5	731.6	28.6	35.8	36.7	505.1	9.9	69.0
Trial4_953	133	NA()	1.1	13.1498	1.4	1.7903	3.1	0.1708	2.7	0.89	1016.6	7.7	17.6	18.1	1041.9	6.1	14.2	14.7	1096.3	9.0	22.2	23.5	1096.3	22.2	92.7
Trial4_954	136	NA()	2.4	19.6922	3.1	0.3054	4.6	0.0436	3.4	0.74	275.3	2.6	5.2	5.4	270.6	3.5	5.8	6.0	230.9	25.5	34.6	35.7	275.3	5.2	NA
Trial4_955	31	NA()	1.4	13.2813	2.7	1.8095	4.4	0.1744	3.5	0.79	1036.2	10.1	19.0	19.5	1048.9	9.3	15.9	16.3	1076.4	19.1	27.9	29.0	1076.4	27.9	96.3
Trial4_956	41	NA()	1.3	5.4560	1.1	12.0699	3.0	0.4778	2.8	0.93	2517.8	13.5	37.6	38.8	2609.8	8.1	20.2	20.9	2682.8	9.5	19.3	20.3	2682.8	19.3	93.8
Trial4_957	61	NA()	1.1	13.5216	2.1	1.6016	3.9	0.1571	3.3	0.84	940.8	8.7	17.1	17.6	970.8	7.8	14.6	15.0	1040.2	15.4	25.6	26.7	1040.2	25.6	90.4
Trial4_958	16	NA()	1.5	13.5131	3.5	1.8342	4.7	0.1798	3.2	0.68	1066.1	11.1	19.9	20.4	1057.8	13.6	18.8	19.2	1041.5	35.1	40.7	41.4	1041.5	40.7	102.4
Trial4_959	82	NA()	1.9	13.7407	1.8	1.7151	3.6	0.1710	3.1	0.87	1017.6	8.8	18.1	18.6	1014.2	7.0	14.4	14.9	1007.7	11.7	23.7	24.9	1007.7	23.7	101.0
Trial4_960	46	NA()	0.9	12.7802	2.2	2.0407	4.0	0.1892	3.3	0.84	1117.2	10.2	20.1	20.6	1129.2	8.6	15.9	16.4	1153.1	15.4	25.3	26.5	1153.1	25.3	96.9
Trial4_961	89	NA()	1.3	10.9225	1.3	3.0652	3.1	0.2429	2.8	0.90	1401.9	10.5	23.7	24.3	1424.0	7.3	16.8	17.3	1458.1	9.0	21.3	22.5	1458.1	21.3	96.1
Trial4_962	117	NA()	1.5	15.3779	2.5	0.5941	4.2	0.0663	3.4	0.80	413.8	4.0	7.8	8.0	473.5	5.1	9.0	9.3	775.2	19.1	28.6	29.7	413.8	7.8	53.4
Trial4_963	43	NA()	1.3	17.2384	3.3	0.6542	4.6	0.0818	3.2	0.69	507.0	5.7	10.0	10.2	511.1	7.5	10.9	11.2	530.2	32.3	39.2	40.1	507.0	10.0	95.6
Trial4_964	28	NA()	2.4	13.5814	2.9	1.7590	4.6	0.1733	3.5	0.77	1030.5	10.3	19.1	19.5	1030.5	10.0	16.2	16.6	1031.3	22.2	30.2	31.2	1031.3	30.2	99.9
Trial4_965	61	NA()	1.7	17.8544	3.7	0.2868	4.9	0.0372	3.2	0.66	235.2	2.8	4.8	4.9	256.1	5.6	7.2	7.3	452.7	48.3	53.3	54.0	235.2	4.8	NA
Trial4_966	26	NA()	1.1	12.9328	2.9	1.9477	4.6	0.1828	3.5	0.77	1082.1	10.8	19.9	20.4	1097.6	10.7	17.0	17.5	1129.5	23.5	31.0	31.9	1129.5	31.0	95.8
Trial4_967	27	NA()	0.8	5.4710	1.3	13.5121	3.3	0.5364	3.0	0.92	2768.3	14.0	40.4	41.6	2716.1	8.7	20.5	21.3	2678.3	11.2	20.2	21.1	2678.3	20.2	103.4
Trial4_968	279	NA()	1.0	18.1412	1.8	0.4872	3.4	0.0641	2.9	0.85	400.7	2.5	7.0	7.2	403.0	2.8	7.1	7.4	417.2	12.6	25.9	27.3	400.7	7.0	96.0
Trial4_969	100	NA()	1.2	5.0191	1.0	15.0024	2.0	0.5464	1.7	0.87	2810.0	11.3	40.0	41.3	2815.4	6.8	19.9	20.7	2820.0	8.5	18.6	19.6	2820.0	18.6	99.6
Trial4_970	82	NA()	1.3	13.1679	1.7	1.9373	3.5	0.1851	3.0	0.87	1094.8	9.3	19.4	19.9	1094.1	7.2	15.0	15.5	1093.6	11.1	23.1	24.4	1093.6	23.1	100.1
Trial4_971	242	NA()	5.1	12.8703	1.0	1.9076	2.3	0.1781	2.0	0.89	1056.8	6.0	17.5	18.0	1083.7	4.6	13.9	14.4	1139.1	6.5	21.2	22.5	1139.1	21.2	92.8
Trial4_972	178	NA()	0.8	18.1090	2.2	0.4889	3.9	0.0642	3.2	0.82	401.4	3.3	7.3	7.5	404.2	3.7	7.5	7.8	421.2	16.0	27.7	29.0	401.4	7.3	95.3
Trial4_973	225	NA()	3.2	13.4391	1.2	1.5218	2.6	0.1484	2.3	0.89	892.0	5.6	15.1	15.6	939.2	4.7	12.9	13.4	1052.6	7.4	21.7	23.1	1052.6	21.7	84.7
Trial4_974	383	NA()	6.5	12.3940	0.8	1.9564	1.6	0.1759	1.4	0.86	1044.7	3.4	16.6	17.1	1100.6	2.9	13.5	14.1	1213.7	4.7	20.5	21.8	1213.7	20.5	86.1
Trial4_975	196	NA()	1.8	16.7170	1.7	0.8007	3.3	0.0971	2.9	0.86	597.5	4.4	10.6	10.9	597.2	4.3	9.9	10.2	597.1	12.0	25.0	26.4	597.5	10.6	100.1
Trial4_976	39	NA()	1.4	5.8052	1.2	10.9865	3.1	0.4628	2.9	0.93	2451.8	13.9	37.1	38.2	2522.0	8.3	20.1	20.8	2579.7	9.7	19.5	20.5	2579.7	19.5	95.0
Trial4_977	373	NA()	1.7	17.2315	1.5	0.5889	2.9	0.0736	2.5	0.86	458.0	1.9	7.7	7.9	470.2	2.3	7.8	8.1	531.0	10.0	24.4	25.8	458.0	7.7	86.3
Trial4_978	175	NA()	1.0	5.8771	0.9	10.6410	1.6	0.4538	1.3	0.82	2412.0	10.0	35.4	36.5	2492.3	5.8	19.1	19.9	2559.1	6.3	18.1	19.2	2559.1	18.1	94.3
Trial4_979	42	NA()	1.8	9.1843	1.6	4.8797	3.5	0.3252	3.1	0.89	1815.0	12.5	29.5	30.3	1798.8	8.3	18.5	19.2	1780.8	10.6	21.3	22.4	1780.8	21.3	101.9
Trial4_980	26	NA()	2.1	13.1023	3.0	1.9140	4.6	0.1820	3.5	0.77	1077.7	9.9	19.4	19.9	1086.0	9.8	16.4	16.8	1103.6	21.4	29.5	30.4	1103.6	29.5	97.7
Trial4_981	21	NA()	1.0	13.8456	3.5	1.6934	4.8	0.1701	3.2	0.67	1012.8	9.7	18.6	19.0	1006.0	11.6	17.1	17.5	992.3	30.3	36.7	37.5	992.3	36.7	102.1
Trial4_982	77	NA()	1.2	9.9949	1.3	3.8807	3.1	0.2814	2.8	0.90	1598.6	10.7	26.1	26.9	1609.7	7.2	17.4	18.0	1625.0	8.7	20.8	21.9	1625.0	20.8	98.4
Trial4_983	129	NA()	1.9	13.1078	1.4	1.9683	3.0	0.1872	2.7	0.89	1106.2	7.6	18.7	19.3	1104.7	5.9	14.5	15.0	1102.7	9.0	22.2	23.5	1102.7	22.2	100.3
Trial4_984	302	NA()	1.2	19.0456	2.2	0.3160	3.7	0.0437	3.0	0.82	275.5	1.4	4.7	4.9	278.8	2.1	5.2	5.4	307.5	16.2	28.2	29.5	275.5	4.7	NA
Trial4_985	97	NA()	1.8	19.7684	3.8	0.3073	5.2	0.0441	3.5	0.68	278.1	2.7	5.3	5.4	272.1	4.6	6.6	6.7	222.0	38.2	44.8	45.6	278.1	5.3	NA
Trial4_986	119	NA()	2.8	13.6853	1.5	1.7878	3.2	0.1775	2.8	0.88	1053.5	6.8	17.7	18.2	1041.0	5.6	14.0	14.5	1015.9	9.9	22.8	24.1	1015.9	22.8	103.7
Trial4_987	34	NA()	2.0	9.8558	1.8	4.0351	3.7	0.2886	3.2	0.87	1634.4	11.2	26.8	27.5	1641.3	8.3	18.0	18.7	1651.0	12.3	22.5	23.6	1651.0	22.5	99.0
Trial4_988	25	NA()	1.4	12.9871	3.1	1.8919	4.7	0.1783	3.5	0.75	1057.6	9.1	18.8	19.3	1078.3	10.2	16.6	17.0	1121.2	24.4	31.7	32.6	1121.2	31.7	94.3
Trial4_989	73	NA()	2.1	16.9882	2.9	0.6894	4.5	0.0850	3.4	0.76	525.8	4.5	9.6	9.9	532.5	5.9	10.1	10.4	562.1	24.4	32.9	34.0	525.8	9.6	93.5
Trial4_990	96	NA()	1.3	13.1493	1.6	1.9107	3.3	0.1823	2.9	0.88	1079.5	7.4	18.3	18.8	1084.8	6.1	14.5	15.0	1096.4	10.6	22.9	24.2	1096.4	22.9	98.5
Trial4_991	331	NA()	3.5	13.5842	1.0	1.6792	2.0	0.1655	1.7	0.87	987.4	2.9	15.7	16.2	1000.7	2.7	12.8	13.3	1030.9	5.7	21.3	22.6	1030.9	21.3	95.8
Trial4_992	46	NA()	0.6	5.2202	1.0	14.0774	2.8	0.5332	2.6	0.93	2755.0	11.6	39.5	40.8	2754.9	6.9	19.9	20.7	2755.6	8.5	18.7	19.7	2755.6	18.7	100.0
Trial4_993	70	NA()	1.0	5.6255	0.9	11.9856	2.5	0.4892	2.3	0.93	2567.4	11.0	37.4	38.5	2603.3	6.5	19.5	20.3	2632.1	7.6	18.5	19.5	2632.1	18.5	97.5
Trial4_994	477	NA()	2.2	12.6753	0.8	2.0285	1.3	0.1866	1.0	0.78	1102.7	0.8	17.1	17.7	1125.1	1.5	13.5	14.0	1169.4	4.1	20.5	21.8	1169.4	20.5	94.3
Trial4_995	191	NA()	1.3	17.8735	2.1	0.5275	3.7	0.0684	3.1	0.82	426.6	2.7	7.5	7.7	430.1	3.4	7.7	7.9	450.3	15.6	27.4	28.7	426.6	7.5	94.7
Trial4_996	13	NA()	1.7	13.8712	3.5	1.7005	4.8	0.1711	3.2	0.67	1018.4	10.0	18.8	19.2	1008.7	13.8	18.7	19.1	988.5	38.3	43.5	44.2	988.5	43.5	103.0
Trial4_997	27	NA()	0.6	13.5287	3.1	1.7089	4.7	0.1678	3.5	0.75	999.7	8.8	17.9	18.4	1011.9	9.6	15.8	16.3	1039.2	23.4	31.1	32.0	1039.2	31.1	96.2
Trial4_998	86	NA()	1.1	18.7672	3.7	0.3675	5.1	0.0500	3.5	0.69	314.8	2.9	5.9	6.1	317.8	5.0	7.4	7.5	341.0	36.0	42.7	43.6	314.8	5.9	NA
Trial4_999	82	NA()	0.9	12.3585	1.6	2.3779	3.3	0.2132	2.9	0.88	1246.0	8.4	20.9	21.4	1236.0	6.5	15.5	16.0	1219.4	10.3	22.4	23.7	1219.4	22.4	102.2
Trial4_1000	94	NA()	1.8	13.2736	1.6	1.9329	3.3	0.1862	2.9	0.88	1100.5	7.5	18.6	19.1	1092.5	6.1	14.5	15.0	1077.5	10.5	22.9	24.2	1077.5	22.9	102.1
Trial4_1001	49	NA()	1.5	13.0550	2.1	1.9742	3.9	0.1870	3.3	0.84	1105.2	8.5	19.1	19.6	1106.8	7.5	15.2	15.7	1110.7	14.7	25.0	26.1	1110.7	25.0	99.5
Trial4_1002	141	NA()	1.8	8.9917	0.9	5.3573	2.1	0.3495	1.9	0.91	1932.3	8.7	29.4	30.4	1878.1	5.3	17.7	18.3	1819.3	6.2	19.4	20.6	1819.3	19.4	106.2
Trial4_1003	48	NA()	1.1	12.0979	2.1	2.1067	3.9	0.1849	3.3	0.84	1093.9	8.1	18.8	19.3	1151.0	7.4	15.4	15.9	1261.2	13.7	24.0	25.2	1261.2	24.0	86.7
Trial4_1004	121	NA()	0.9	13.3606	1.5	1.8941	3.1	0.1836	2.7	0.88	1086.7	6.1	17.9	18.5	1079.0	5.1	14.0	14.5	1064.4	9.2	22.4	23.7	1064.4	22.4	102.1
Trial4_1005	67	NA()	1.6	11.2130	1.6	2.8095	3.4	0.2286	3.0	0.88	1327.1	8.2	21.8	22.5	1358.1	6.4	16.1	16.7	1408.1	9.9	21.8	23.0	1408.1	21.8	94.2
Trial4_1006	79	NA()	0.4	9.9239	1.3	4.0465	3.0	0.2914	2.7	0.91	1648.4	8.8	26.1	26.8	1643.6	6.1	17.1	17.8	1638.3	8.0	20.5	21.7	1638.3	20.5	100.6
Trial4_1007	53	NA()	1.3	9.0989	1.4	4.6886	3.2	0.3096	2.9	0.90	1738.5	9.8	27.5	28.3	1765.2	6.8	17.8	18.5	1797.8	8.9	20.5	21.6	1797.8	20.5	96.7
Trial4_1008	146	NA()	0.6	7.5849	0.9	7.1384	1.9	0.3929	1.7	0.89	2136.1	7.3	31.5	32.5	2128.8	4.4	18.1	18.8	2122.6	5.2	18.5	19.7	2122.6	18.5	100.6
Trial4_1009	837	NA()	1.1	14.8739	0.8	1.0119	1.5	0.1092	1.3	0.84	668.1	5.4	12.0	12.3	709.8	4.5	11.0	11.4	845.0	5.0	21.7	23.1	668.1	12.0	79.1
Trial4_1010	368	NA()	1.8	13.3041	0.8	1.9763	1.6	0.1908	1.3	0.85	1125.6	1.5	17.5	18.1	1107.5	1.9	13.4	14.0	1072.9	4.8	20.9	22.3	1072.9	20.9	104.9
Trial4_1011	42	NA()	2.3	9.4454	1.6	4.5068	3.5	0.3089	3.1	0.89	1735.2	10.2	27.6	28.4	1732.2	7.2	17.9	18.5	1729.5	10.2	21.2	22.3	1729.5	21.2	100.3
Trial4_1012	40	NA()	2.0	14.1353	2.9	1.2474	4.5	0.1279	3.5	0.77	776.1	6.2	13.8	14.2	822.2	7.4	13.3	13.7	950.0	20.5	29.1	30.2	776.1	13.8	81.7
Trial4_1013	63	NA()	1.7	9.7139	1.4	4.1166	3.2	0.2902	2.9	0.90	1642.3	9.2	26.1	26.9	1657.6	6.5	17.4	18.0	1677.9	8.9	20.8	21.9	1677.9	20.8	97.9
Trial4_1014	695	NA()	18.7	13.6381	0.8	1.5761	1.4	0.1560	1.1	0.81	934.3	4.8	15.4	15.9	960.8	3.7	12.8	13.3	1022.9	4.4	21.0	22.4	1022.9	21.0	91.3
Trial4_1015	16	NA()	2.1	8.0528	2.3	5.9021	4.1	0.3449	3.4	0.83	1910.0	12.1	30.4	31.3	1961.5	9.7	19.7	20.3	2017.1	15.1	23.4	24.4	2017.1	23.4	94.7
Trial4_1016	289	NA()	1.1	8.9185	0.8	4.8231	1.3	0.3121	1.0	0.78	1751.1	3.7	26.1	27.0	1788.9	2.6	16.8	17.5	1834.1	3.6	18.7	19.9	1834.1	18.7	95.5
Trial4_1017	112	NA()	0.8	8.7634	0.9	5.3234	2.4	0.3385	2.2	0.92	1879.4	8.3	28.7	29.6	1872.6	5.3	17.6	18.3	1865.9	6.3	19.4	20.5	1865.9	19.4	100.7
Trial4_1018	41	NA()	1.0	17.7035	3.5	0.5319	4.7	0.0683	3.1	0.67	426.1	3.9	7.9	8.2	433.1	7.3	10.1	10.3	471.5	41.0	46.7	47.4	426.1	7.9	90.4
Trial4_1019	73	NA()	1.8	12.3008	1.7	2.3572	3.4	0.2104	3.0	0.87	1230.9	7.7	20.4	21.0	1229.7	6.2	15.3	15.9	1228.6	10.7	22.6	23.8	1228.6	22.6	100.2
Trial4_1020	129	NA()	2.0	18.2885	2.5	0.5428	4.1	0.0720	3.2	0.79	448.4	3.0	7.9	8.1	440.3	3.9	8.0	8.3	399.1	18.8	29.5	30.7	448.4	7.9	112.4
Trial4_1021	68	NA()	1.7	13.2993	1.9	1.8043	3.7	0.1741	3.1	0.85	1034.8	6.4	17.4	17.9	1047.0	6.2	14.3	14.8	1073.7	13.4	24.4	25.5	1073.7	24.4	96.4
Trial4_1022	129	NA()	1.5	15.8841	2.7	0.4760	4.3	0.0549	3.3	0.78	344.3	2.1	6.0	6.2	395.3	3.6	7.4	7.6	706.8	19.1	28.8	29.9	344.3	6.0	NA
Trial4_1023	33	NA()	1.2	13.4695	2.8	1.7435	4.4	0.1704	3.4	0.78	1014.3	7.3	17.4	17.9	1024.8	7.8	14.9	15.3	1048.1	18.6	27.6	28.7	1048.1	27.6	96.8
Trial4_1024	92	NA()	0.7	9.7902	1.2	4.1715	2.8	0.2963	2.5	0.91	1673.1	7.7	26.0	26.8	1668.4	5.4	17.0	17.6	1663.4	7.3	20.1	21.3	1663.4	20.1	100.6
Trial4_1025	75	NA()	1.5	11.1049	1.5	3.0046	3.2	0.2421	2.9	0.89	1397.6	7.5	22.5	23.1	1408.8	5.8	16.1	16.7	1426.1	9.1	21.4	22.6	1426.1	21.4	98.0
Trial4_1026	100	NA()	1.0	11.1722	1.3	3.1001	2.9	0.2513	2.6	0.90	1445.2	7.0	22.9	23.6	1432.7	5.2	16.0	16.6	1415.0	7.7	20.9	22.1	1415.0	20.9	102.1
Trial4_1027	115	NA()	2.7	13.6941	1.5	1.7253	3.2	0.1714	2.8	0.88	1020.0	5.9	17.0	17.5	1018.0	5.2	13.7	14.2	1014.6	10.2	22.9	24.2	1014.6	22.9	100.5
Trial4_1028	469	NA()	1.1	17.2009	1.5	0.4381	2.9	0.0547	2.5	0.86	343.2	0.5	5.7	5.8	368.9	1.5	6.3	6.5	534.9	10.3	24.5	25.9	343.2	5.7	NA
Trial4_1029	111	NA()	1.7	13.5488	1.5	1.8118	3.2	0.1781	2.8	0.88	1056.7	6.2	17.5	18.1	1049.7	5.2	13.9	14.4	1036.2	9.8	22.7	24.0	1036.2	22.7	102.0
Trial4_1030	70	NA()	6.0	13.3802	1.9	1.6620	3.7	0.1614	3.2	0.85	964.3	6.9	16.6	17.1	994.2	6.4	14.0	14.5	1061.5	13.1	24.3	25.5	1061.5	24.3	90.9
Trial4_1031	185	NA()	0.7	16.1772	1.6	0.9241	3.2	0.1085	2.8	0.86	663.9	3.6	11.2	11.6	664.5	3.8	10.3	10.7	667.7	11.0	24.4	25.7	663.9	11.2	99.4
Trial4_1032	95	NA()	1.1	5.3513	0.9	13.2286	2.1	0.5137	1.9	0.90	2672.9	9.3	38.0	39.2	2696.1	5.5	19.4	20.1	2714.8	6.6	18.0	19.0	2714.8	18.0	98.4
Trial4_1033	84	NA()	1.0	19.2568	3.4	0.2798	4.6	0.0391	3.1	0.68	247.2	2.2	4.6	4.7	250.5	4.0	5.9	6.1	282.3	35.8	42.6	43.5	247.2	4.6	NA
Trial4_1034	201	NA()	1.8	9.7922	0.8	4.0702	1.8	0.2892	1.6	0.89	1637.5	6.1	25.1	25.9	1648.4	4.1	16.6	17.2	1663.0	5.0	19.4	20.7	1663.0	19.4	98.5
Trial4_1035	204	NA()	2.3	13.5065	1.1	1.8076	2.5	0.1771	2.2	0.89	1051.4	4.7	17.0	17.6	1048.2	3.9	13.4	14.0	1042.5	7.1	21.7	23.0	1042.5	21.7	100.9
Trial4_1036	97	NA()	1.1	10.3039	1.2	3.4949	2.9	0.2613	2.6	0.91	1496.5	9.2	24.3	25.0	1526.1	6.4	16.8	17.4	1568.2	7.9	20.6	21.8	1568.2	20.6	95.4
Trial4_1037	150	NA()	1.2	11.1543	1.1	3.0228	2.5	0.2447	2.2	0.90	1410.8	6.9	22.5	23.1	1413.4	4.9	15.8	16.4	1418.1	6.6	20.5	21.7	1418.1	20.5	99.5
Trial4_1038	51	NA()	1.4	13.7067	2.2	1.7243	4.0	0.1715	3.3	0.83	1020.3	7.7	17.7	18.2	1017.6	7.2	14.6	15.0	1012.7	15.8	25.9	27.0	1012.7	25.9	100.7
Trial4_1039	151	NA()	1.5	9.8335	2.2	0.5815	4.0	0.0415	3.4	0.83	262.1	1.9	4.7	4.8	465.4	3.9	8.3	8.6	1655.2	14.3	23.6	24.6	262.1	4.7	NA
Trial4_1040	142	NA()	2.4	8.6948	0.9	5.1629	2.1	0.3257	2.0	0.91	1817.6	8.3	28.0	28.8	1846.5	5.3	17.5	18.2	1880.1	6.0	19.2	20.4	1880.1	19.2	96.7
Trial4_1041	16	NA()	1.3	13.9403	3.5	1.6854	4.7	0.1705	3.2	0.68	1014.7	10.2	18.8	19.3	1003.0	13.1	18.1	18.5	978.4	35.4	41.0	41.7	978.4	41.0	103.7
Trial4_1042	38	NA()	1.6	7.6814	1.4	6.7288	3.3	0.3750	3.0	0.91	2053.0	12.4	32.1	33.1	2076.4	8.0	19.1	19.8	2100.4	9.8	20.3	21.4	2100.4	20.3	97.7
Trial4_1043	129	NA()	2.6	13.0267	1.4	1.9323	3.0	0.1826	2.7	0.89	1081.4	7.1	18.2	18.8	1092.3	5.7	14.3	14.8	1115.1	8.9	22.1	23.4	1115.1	22.1	97.0
Trial4_1044	279	NA()	4.1	13.5905	1.0	1.7113	2.1	0.1688	1.9	0.88	1005.2	3.6	16.1	16.6	1012.8	3.1	13.0	13.5	1030.0	6.1	21.4	22.7	1030.0	21.4	97.6
Trial4_1045	125	NA()	2.8	13.2585	1.4	1.8502	3.0	0.1780	2.7	0.89	1056.0	6.5	17.7	18.2	1063.5	5.3	14.0	14.5	1079.8	8.9	22.2	23.5	1079.8	22.2	97.8
Trial4_1046	223	NA()	0.6	13.5975	1.1	1.7944	2.4	0.1770	2.1	0.89	1050.8	4.7	17.0	17.5	1043.4	3.8	13.4	13.9	1028.9	6.8	21.6	23.0	1028.9	21.6	102.1
Trial4_1047	151	NA()	1.8	19.1659	2.9	0.3233	4.4	0.0450	3.3	0.75	283.5	2.0	5.1	5.2	284.4	3.1	5.8	5.9	293.2	23.3	32.9	34.0	283.5	5.1	NA
Trial4_1048	82	NA()	0.9	5.6860	0.9	11.8086	2.3	0.4872	2.1	0.92	2558.5	9.7	36.9	38.1	2589.3	5.8	19.3	20.1	2614.3	6.8	18.2	19.3	2614.3	18.2	97.9
Trial4_1049	36	NA()	1.7	8.6776	1.6	5.4196	3.5	0.3412	3.1	0.89	1892.6	11.3	29.9	30.7	1888.0	7.7	18.5	19.2	1883.6	10.4	21.0	22.1	1883.6	21.0	100.5
Trial4_1050	34	NA()	1.2	13.1633	2.6	1.8655	4.3	0.1782	3.4	0.79	1057.0	8.4	18.4	18.9	1068.9	8.5	15.5	16.0	1094.3	19.0	27.8	28.9	1094.3	27.8	96.6
Trial4_1051	248	NA()	1.2	15.5019	1.3	1.1383	2.7	0.1280	2.3	0.87	776.6	3.2	12.7	13.1	771.7	3.2	11.1	11.5	758.3	8.3	22.9	24.3	776.6	12.7	102.4
Trial4_1052	37	NA()	2.3	13.7196	2.6	1.7835	4.3	0.1775	3.4	0.79	1053.6	8.3	18.4	18.9	1039.5	8.2	15.2	15.7	1010.8	18.8	27.9	28.9	1010.8	27.9	104.2
Trial4_1053	220	NA()	4.5	12.4583	1.1	1.9107	2.4	0.1727	2.2	0.89	1027.1	4.3	16.6	17.1	1084.8	3.7	13.6	14.2	1203.6	6.5	21.0	22.3	1203.6	21.0	85.3
Trial4_1054	158	NA()	2.1	13.8375	1.3	1.7056	2.9	0.1712	2.6	0.88	1019.0	5.4	16.8	17.3	1010.6	4.5	13.4	13.9	993.5	8.4	22.3	23.6	993.5	22.3	102.6
Trial4_1055	27	NA()	1.2	12.5494	2.9	1.8882	4.5	0.1719	3.5	0.77	1022.8	8.5	18.1	18.5	1077.0	9.0	15.9	16.3	1189.2	20.2	28.4	29.4	1189.2	28.4	86.0
Trial4_1056	86	NA()	0.8	19.2590	3.5	0.2683	4.7	0.0375	3.1	0.67	237.3	2.1	4.5	4.6	241.3	4.2	6.0	6.1	282.1	40.1	46.4	47.2	237.3	4.5	NA
Trial4_1057	253	NA()	2.6	13.6198	1.1	1.7065	2.3	0.1686	2.0	0.88	1004.7	3.8	16.1	16.7	1011.0	3.4	13.0	13.6	1025.6	6.6	21.6	22.9	1025.6	21.6	98.0
Trial4_1058	49	NA()	1.3	12.9421	2.3	1.7630	4.0	0.1656	3.3	0.82	987.6	7.7	17.2	17.7	1031.9	7.5	14.8	15.3	1128.1	16.1	25.8	26.9	1128.1	25.8	87.5
Trial4_1059	62	NA()	1.2	10.7999	1.5	3.2275	3.3	0.2529	3.0	0.89	1453.5	9.3	23.8	24.5	1463.8	6.9	16.7	17.3	1479.6	9.9	21.6	22.8	1479.6	21.6	98.2
Trial4_1060	25	NA()	1.1	13.2340	3.0	1.8846	4.6	0.1810	3.5	0.76	1072.3	9.0	18.9	19.4	1075.7	9.5	16.1	16.6	1083.5	22.1	30.0	31.0	1083.5	30.0	99.0
Trial4_1061	46	NA()	1.2	13.7831	2.4	1.6508	4.1	0.1651	3.4	0.81	985.0	8.0	17.4	17.8	989.8	8.0	14.8	15.3	1001.5	18.7	27.8	28.8	1001.5	27.8	98.4
Trial4_1062	100	NA()	1.6	9.5528	1.1	4.4315	2.7	0.3072	2.4	0.91	1726.8	9.9	27.4	28.2	1718.2	6.4	17.5	18.1	1708.7	7.5	20.1	21.3	1708.7	20.1	101.1
Trial4_1063	192	NA()	2.4	13.3652	1.2	1.8703	2.6	0.1814	2.3	0.89	1074.5	6.1	17.8	18.3	1070.6	4.7	13.9	14.4	1063.7	7.4	21.7	23.0	1063.7	21.7	101.0

Trial4_1064	38	NA()	2.3	12.4925	2.3	2.1840	4.1	0.1980	3.4	0.82	1164.4	9.8	20.5	21.0	1176.0	8.6	16.2	16.7	1198.2	16.4	25.8	26.9	1198.2	25.8	97.2
Trial4_1065	26	NA()	0.6	10.0726	2.2	3.4994	4.0	0.2558	3.4	0.84	1468.1	11.7	25.0	25.7	1527.1	9.3	18.1	18.7	1610.6	14.3	23.7	24.8	1610.6	23.7	91.2
Trial4_1066	54	NA()	1.8	10.0789	1.5	3.8516	3.4	0.2817	3.0	0.89	1599.8	11.1	26.4	27.1	1603.6	7.7	17.6	18.3	1609.4	10.2	21.5	22.6	1609.4	21.5	99.4
Trial4_1067	115	NA()	1.1	13.2351	1.5	1.9506	3.1	0.1873	2.8	0.89	1106.8	7.7	18.8	19.3	1098.6	6.0	14.5	15.0	1083.3	9.6	22.5	23.7	1083.3	22.5	102.2

Element2 (LA-SC-ICP-MS)																									
						Isotope ratios						Apparent ages (Ma)													
Analysis	U (ppm)	206Pb 204Pb	U/Th	206Pb* 207Pb*	± (%)	207Pb* 235U*	± (%)	206Pb* 238U	± (%)	error corr.	206Pb* 238U*	±1s(a) (Ma)	±1s(b) (Ma)	±1s(c) (Ma)	207Pb* 235U	±1s(a) (Ma)	±1s(b) (Ma)	±1s(c) (Ma)	206Pb* 207Pb*	±1s(a) (Ma)	±1s(b) (Ma)	±1s(c) (Ma)	Best age (Ma)	± (Ma)	Conc (%)
49127	138	NA()	0.9	20.5018	3.5	0.1443	4.7	0.0215	3.2	0.67	136.9	1.3	2.9	2.9	136.8	2.6	3.7	3.8	137.1	42.2	47.1	47.9	136.9	2.9	NA
49127	140	NA()	0.9	20.6480	3.4	0.1423	4.7	0.0213	3.2	0.68	136.0	1.3	2.9	2.9	135.1	2.4	3.6	3.7	120.4	39.1	44.3	45.2	136.0	2.9	NA
49127	127	NA()	0.8	20.0314	3.6	0.1552	4.8	0.0226	3.2	0.66	143.8	1.4	3.0	3.1	146.5	3.1	4.2	4.3	191.4	46.9	51.2	52.0	143.8	3.0	NA
49127	163	NA()	1.2	20.9590	3.5	0.1410	5.2	0.0214	3.9	0.75	136.8	1.2	2.8	2.9	133.9	2.5	3.6	3.7	85.1	41.2	46.2	47.2	136.8	2.8	NA
49127	123	NA()	1.7	19.7900	3.6	0.1501	4.8	0.0215	3.2	0.66	137.4	1.4	2.9	3.0	142.0	2.9	4.0	4.1	219.5	44.7	49.2	50.0	137.4	2.9	NA
49127	60	NA()	1.7	20.8484	4.4	0.1421	5.5	0.0215	3.2	0.59	137.1	1.6	3.0	3.1	134.9	4.4	5.1	5.1	97.5	76.7	79.5	80.0	137.1	3.0	NA
49127	481	NA()	1.4	20.4117	2.0	0.1488	3.9	0.0220	3.4	0.86	140.5	0.3	2.6	2.7	140.8	1.1	2.9	3.0	147.4	18.2	27.6	29.0	140.5	2.6	NA
49127	554	NA()	1.7	19.7383	1.9	0.1507	3.8	0.0216	3.3	0.87	137.6	0.3	2.6	2.6	142.5	0.9	2.9	3.0	225.5	15.8	25.9	27.4	137.6	2.6	NA
49127	148	NA()	0.8	17.6287	4.7	0.1660	6.8	0.0212	5.0	0.73	135.4	4.3	5.0	5.0	155.9	7.0	7.6	7.6	480.9	80.1	82.4	82.9	135.4	5.0	NA
49127	151	NA()	1.0	20.3057	3.6	0.1448	5.3	0.0213	3.9	0.74	136.1	1.2	2.8	2.9	137.3	2.8	3.8	3.9	159.6	45.8	50.3	51.1	136.1	2.8	NA
49127	241	NA()	1.4	17.5222	2.6	0.1710	4.6	0.0217	3.7	0.82	138.7	0.9	2.7	2.8	160.3	1.9	3.6	3.7	494.3	23.6	30.6	31.8	138.7	2.7	NA
49127	239	NA()	1.0	20.6754	3.0	0.1367	4.8	0.0205	3.8	0.79	130.9	0.9	2.6	2.7	130.1	1.7	3.0	3.1	117.2	28.3	35.2	36.3	130.9	2.6	NA
49127	863	NA()	1.5	20.5710	1.6	0.1497	3.4	0.0223	3.1	0.89	142.5	1.2	2.9	3.0	141.7	1.3	3.0	3.1	129.2	12.9	24.5	26.1	142.5	2.9	NA
49127	161	NA()	1.0	19.9964	3.4	0.1441	5.1	0.0209	3.9	0.76	133.4	1.2	2.7	2.8	136.7	2.3	3.5	3.6	195.4	35.3	40.9	41.8	133.4	2.7	NA
91500	48	NA()	2.3	13.3834	1.9	1.8719	4.1	0.1818	3.6	0.89	1076.7	9.4	20.9	21.3	1071.2	8.0	15.9	16.4	1061.0	15.1	23.4	24.6	1061.0	23.4	101.5
91500	46	NA()	2.5	13.0259	1.9	1.9074	4.1	0.1803	3.7	0.89	1068.5	9.5	20.8	21.2	1083.7	8.1	16.0	16.5	1115.2	14.6	23.0	24.2	1115.2	23.0	95.8
91500	48	NA()	2.4	13.1718	1.9	1.8653	4.1	0.1783	3.6	0.89	1057.5	9.4	20.6	21.0	1068.9	8.1	15.9	16.4	1092.9	14.8	23.1	24.4	1092.9	23.1	96.8
91500	49	NA()	2.5	13.2158	1.9	1.8799	4.1	0.1803	3.6	0.89	1068.5	9.5	20.8	21.2	1074.0	8.0	15.9	16.4	1086.3	14.3	22.8	24.1	1086.3	22.8	98.4
91500	47	NA()	2.5	13.2587	1.9	1.8211	4.1	0.1752	3.7	0.89	1040.7	9.4	20.4	20.8	1053.1	8.1	15.8	16.3	1079.7	14.8	23.1	24.4	1079.7	23.1	96.4
91500	53	NA()	2.4	13.3786	1.8	1.8683	4.0	0.1814	3.6	0.89	1074.4	9.6	20.9	21.4	1069.9	7.9	15.8	16.3	1061.7	13.9	22.7	23.9	1061.7	22.7	101.2
91500	53	NA()	2.4	13.3828	1.8	1.8103	4.0	0.1758	3.6	0.89	1043.9	9.4	20.4	20.8	1049.2	7.8	15.7	16.1	1061.1	14.1	22.7	24.0	1061.1	22.7	98.4
91500	54	NA()	2.4	13.0903	1.8	1.9064	4.0	0.1811	3.6	0.89	1072.8	9.6	20.9	21.3	1083.3	7.9	15.9	16.4	1105.4	13.8	22.4	23.7	1105.4	22.4	97.1
91500	52	NA()	2.4	13.2783	1.9	1.7905	4.1	0.1725	3.6	0.89	1025.9	9.2	20.0	20.5	1042.0	7.9	15.6	16.1	1076.8	14.5	22.9	24.2	1076.8	22.9	95.3
91500	54	NA()	2.5	13.7656	1.9	1.7964	4.0	0.1794	3.6	0.89	1063.9	9.5	20.7	21.2	1044.2	7.9	15.7	16.1	1004.1	14.8	23.3	24.6	1004.1	23.3	106.0
91500	51	NA()	2.2	13.6172	1.9	1.7874	4.0	0.1766	3.6	0.89	1048.4	9.4	20.4	20.9	1040.9	7.8	15.6	16.1	1026.0	14.4	23.0	24.3	1026.0	23.0	102.2
91500	51	NA()	2.4	13.4806	1.9	1.7888	4.1	0.1750	3.6	0.89	1039.4	9.4	20.3	20.8	1041.4	7.9	15.6	16.1	1046.4	14.2	22.8	24.1	1046.4	22.8	99.3
91500	55	NA()	2.5	13.3399	1.8	1.8754	4.0	0.1815	3.6	0.89	1075.3	9.7	21.0	21.4	1072.4	8.0	15.9	16.4	1067.5	14.3	22.9	24.1	1067.5	22.9	100.7
91500	49	NA()	2.3	13.1234	1.9	1.8170	4.1	0.1730	3.6	0.89	1028.7	9.4	20.2	20.6	1051.6	8.1	15.8	16.3	1100.3	15.0	23.2	24.5	1100.3	23.2	93.5
91500	48	NA()	2.3	13.2611	1.9	1.8651	4.1	0.1795	3.7	0.89	1064.0	9.7	20.8	21.2	1068.8	8.2	16.0	16.4	1079.4	15.0	23.2	24.5	1079.4	23.2	98.6
94-35	38	NA()	2.1	18.6696	11.2	0.0627	11.8	0.0085	3.5	0.30	54.5	0.9	1.4	1.4	61.7	6.6	6.7	6.7	352.7	245.1	245.9	246.1	54.5	1.4	NA
94-35	48	NA()	1.3	20.7923	9.9	0.0589	10.5	0.0089	3.4	0.32	57.0	0.9	1.4	1.4	58.1	5.4	5.5	5.5	103.9	223.1	224.1	224.3	57.0	1.4	NA
94-35	45	NA()	1.3	17.6558	8.8	0.0676	9.4	0.0087	3.4	0.36	55.6	0.9	1.4	1.4	66.4	5.4	5.6	5.6	477.5	182.8	183.8	184.0	55.6	1.4	NA
94-35	54	NA()	1.3	20.0940	9.2	0.0596	9.8	0.0087	3.4	0.35	55.7	0.9	1.4	1.4	58.8	5.1	5.2	5.2	184.1	203.1	204.2	204.3	55.7	1.4	NA
94-35	45	NA()	1.6	20.1604	10.7	0.0605	11.2	0.0088	3.4	0.30	56.8	0.9	1.4	1.4	59.6	6.0	6.1	6.2	176.4	240.6	241.5	241.7	56.8	1.4	NA
94-35	58	NA()	1.3	19.9592	10.1	0.0602	10.6	0.0087	3.4	0.32	55.9	0.9	1.4	1.4	59.3	5.6	5.7	5.7	199.8	223.5	224.5	224.7	55.9	1.4	NA
94-35	39	NA()	1.5	18.8998	10.8	0.0609	11.4	0.0084	3.5	0.31	53.6	1.0	1.4	1.4	60.0	6.1	6.3	6.3	324.9	236.3	237.2	237.3	53.6	1.4	NA
94-35	55	NA()	1.2	18.6946	7.3	0.0658	8.0	0.0089	3.4	0.42	57.3	0.9	1.4	1.4	64.7	4.3	4.5	4.5	349.7	150.6	151.9	152.2	57.3	1.4	NA
94-35	54	NA()	1.2	21.1958	7.5	0.0576	8.2	0.0089	3.4	0.41	56.8	0.9	1.4	1.4	56.9	3.9	4.1	4.1	58.3	163.9	165.3	165.5	56.8	1.4	NA
94-35	60	NA()	1.4	20.6232	7.6	0.0573	8.3	0.0086	3.3	0.40	55.1	0.8	1.3	1.3	56.6	3.9	4.1	4.1	123.2	163.7	165.0	165.3	55.1	1.3	NA
94-35	45	NA()	2.0	18.3441	8.9	0.0637	9.6	0.0085	3.5	0.36	54.4	0.9	1.4	1.4	62.7	5.2	5.4	5.4	392.3	188.8	189.8	190.0	54.4	1.4	NA
94-35	54	NA()	1.2	21.6445	10.1	0.0571	10.7	0.0090	3.3	0.31	57.6	0.8	1.4	1.4	56.4	5.4	5.5	5.5	8.1	233.0	234.0	234.2	57.6	1.4	NA
94-35	34	NA()	1.7	20.8054	16.1	0.0563	16.5	0.0085	3.5	0.21	54.6	1.0	1.4	1.5	55.6	8.6	8.7	8.7	102.5	376.7	377.4	377.5	54.6	1.4	NA
94-35	40	NA()	2.1	21.2546	10.2	0.0573	10.8	0.0088	3.4	0.32	56.7	0.9	1.4	1.4	56.6	5.5	5.6	5.6	51.7	234.4	235.4	235.5	56.7	1.4	NA
94-35	47	NA()	1.6	20.2241	11.5	0.0578	12.0	0.0085	3.4	0.28	54.4	0.9	1.3	1.4	57.0	6.2	6.3	6.3	169.0	259.3	260.2	260.3	54.4	1.3	NA
94-35	64	NA()	2.1	20.1658	7.0	0.0599	7.7	0.0088	3.3	0.43	56.3	0.8	1.3	1.4	59.1	3.7	3.9	3.9	175.7	147.3	148.8	149.1	56.3	1.3	NA
94-35	45	NA()	2.3	19.9166	12.3	0.0580	12.8	0.0084	3.4	0.27	53.8	0.9	1.3	1.4	57.3	6.7	6.8	6.8	204.7	278.5	279.2	279.4	53.8	1.3	NA
PLES	608	NA()	7.2	18.6472	1.1	0.3969	2.6	0.0537	2.4	0.90	337.2	0.8	6.2	6.4	339.4	1.3	6.1	6.4	355.5	8.6	21.8	23.4	337.2	6.2	NA
PLES	609	NA()	7.2	18.5029	1.1	0.4050	2.6	0.0544	2.3	0.90	341.3	0.8	6.3	6.5	345.3	1.3	6.2	6.4	372.9	8.5	21.7	23.3	341.3	6.3	NA
PLES	385	NA()	9.4	18.5215	1.4	0.3939	3.2	0.0529	2.9	0.90	332.6	1.0	6.2	6.3	337.2	1.6	6.2	6.4	370.7	10.8	22.7	24.3	332.6	6.2	NA
PLES	927	NA()	6.1	18.4207	0.9	0.4084	2.2	0.0546	2.0	0.90	342.6	3.1	7.0	7.1	347.7	2.9	6.8	7.0	382.9	7.0	21.1	22.8	342.6	7.0	NA
PLES	706	NA()	5.5	18.6849	1.1	0.3954	2.5	0.0536	2.2	0.90	336.6	1.4	6.3	6.5	338.3	1.6	6.2	6.4	350.9	8.1	21.6	23.3	336.6	6.3	NA
PLES	873	NA()	5.7	18.4730	1.0	0.3982	2.2	0.0534	2.0	0.90	335.2	2.6	6.7	6.8	340.3	2.5	6.5	6.7	376.6	7.3	21.2	22.9	335.2	6.7	NA
PLES	705	NA()	5.3	18.5856	1.1	0.3984	2.5	0.0537	2.2	0.90	337.3	1.4	6.3	6.5	340.5	1.6	6.2	6.4	362.9	8.0	21.5	23.2	337.3	6.3	NA
PLES	663	NA()	5.1	19.0126	1.1	0.3940	2.5	0.0544	2.2	0.90	341.2	1.2	6.3	6.5	337.3	1.5	6.1	6.4	311.4	8.4	21.9	23.5	341.2	6.3	NA
PLES	661	NA()	5.2	18.7005	1.1	0.4098	2.5	0.0556	2.2	0.90	348.8	1.2	6.5	6.6	348.7	1.5	6.3	6.5	349.0	8.1	21.6	23.3	348.8	6.5	NA
PLES	642	NA()	7.2	18.5949	1.1	0.4096	2.6	0.0553	2.3	0.90	346.8	1.0	6.4	6.6	348.6	1.4	6.3	6.5	361.8	8.4	21.7	23.4	346.8	6.4	NA
FC	258	NA()	1.5	13.1996	0.8	1.8845	2.2	0.1805	2.0	0.92	1069.6	5.0	19.2	19.6	1075.6	3.9	14.3	14.8	1088.7	5.9	18.7	20.2	1088.7	18.7	98.2
FC	224	NA()	1.2	13.1319	0.9	1.9224	2.4	0.1832	2.2	0.92	1084.3	5.7	19.6	20.1	1088.9	4.4	14.5	15.0	1099.0	6.4	18.9	20.4	1099.0	18.9	98.7
FC	217	NA()	1.2	13.1100	0.9	1.8548	2.4	0.1764	2.3	0.92	1047.5	5.7	19.0	19.5	1065.1	4.4	14.4	14.9	1102.4	6.5	18.9	20.4	1102.4	18.9	95.0
FC	212	NA()	1.2	13.4011	0.9	1.8885	2.4	0.1836	2.2	0.92	1086.8	6.0	19.7	20.2	1077.1	4.6	14.5	15.0	1058.3	6.7	19.0	20.5	1058.3	19.0	102.7
FC	186	NA()	1.1	13.2129	1.0	1.8938	2.6	0.1816	2.4	0.93	1075.5	6.5	19.7	20.2	1078.9	5.0	14.7	15.2	1086.7	7.2	19.2	20.6	1086.7	19.2	99.0
FC	188	NA()	1.2	13.2978	1.0	1.9395	2.6	0.1871	2.4	0.93	1105.9	6.6	20.2	20.7	1094.8	4.9	14.8	15.3	1073.9	7.0	19.1	20.6	1073.9	19.1	103.0
FC	114	NA()	1.9	13.1949	1.2	1.9554	3.2	0.1872	3.0	0.92	1106.2	8.0	20.7	21.2	1100.3	6.1	15.2	15.7	1089.5	9.1	19.9	21.4	1089.5	19.9	101.5
FC	160	NA()	1.3	13.0265	1.1	1.9636	2.8	0.1856	2.6	0.93	1097.5	7.0	20.2	20.7	1103.1	5.4	15.0	15.5	1115.1	7.7	19.3	20.7	1115.1	19.3	98.4
FC	741	NA()	1.2	13.1190	0.8	1.9727	1.4	0.1878	1.1	0.80	1109.4	4.3	19.6	20.1	1106.2	3.1	14.3	14.8	1101.0	3.7	18.1	19.7	1101.0	18.1	100.8
FC	957	NA()	0.9	13.1312	0.8	1.9503	1.5	0.1858	1.3	0.84	1098.7	8.4	20.7	21.2	1098.5	5.8	15.1	15.6	1099.1	4.7	18.4	19.9	1099.1	18.4	100.0
FC	972	NA()	1.0	13.2448	0.8	2.0364	1.6	0.1957	1.3	0.85	1152.2	9.2	21.8	22.3	1127.8	6.2	15.4	15.9	1081.9	5.0	18.5	20.0	1081.9	18.5	106.5
FC	210	NA()	0.4	13.0963	0.9	1.9512	2.5	0.1854	2.3	0.92	1096.5	6.1	19.9	20.4	1098.9	4.6	14.7	15.2	1104.5	6.7	18.9	20.4	1104.5	18.9	99.3
FC	180	NA()	0.3	13.2466	1.0	1.9500	2.7	0.1874	2.5	0.93	1107.4	6.8	20.3	20.7	1098.4	5.1	14.8	15.3	1081.6	7.2	19.2	20.7	1081.6	19.2	102.4
FC	1157	NA()	0.9	13.2798	0.9	1.9501	1.8	0.1879	1.6	0.88	1110.0	12.5	22.8	23.3	1098.5	8.5	16.3	16.8	1076.6	6.4	18.9	20.4	1076.6	18.9	103.1
FC	137	NA()	1.4	13.1074	1.1	1.9484	3.0	0.1853	2.8	0.93	1095.9	7.6	20.4	20.8	1097.9	5.7	15.1	15.6	1102.8	8.3	19.6	21.0	1102.8	19.6	99.4
R33	80	NA()	1.7	17.8501	2.7	0.5075	4.7	0.0657	3.8	0.81	410.4	4.1	8.5	8.7	416.8	5.4	8.9	9.1	453.2	26.4	33.0	34.0	410.4	8.5	90.5
R33	76	NA()	1.2	18.1222	2.8	0.5179	4.7	0.0681	3.8	0.81	424.7	4.2	8.8	9.0	423.7	5.3	8.9	9.2	419.5	25.4	32.2	33.3	424.7	8.8	101.2
R33	624	NA()	1.1	18.0138	1.0	0.5126	2.4	0.0670	2.1	0.90	418.0	1.1	7.7	7.9	420.2	1.5	7.3	7.6	432.9	7.7	21.2	22.9	418.0	7.7	96.6
R33	665	NA()	1.2	18.0212	1.0	0.5202	2.2	0.0680	1.9	0.89	424.3	1.4	7.8	8.0	425.3	1.7	7.4	7.7	432.0	7.2	21.0	22.7	424.3	7.8	98.2
R33	471	NA()	1.6	18.0097	1.2	0.5180	2.7	0.0677	2.4	0.90	422.3	0.5	7.7	7.9	423.8	1.4	7.3	7.6	433.4	8.7	21.6	23.2	422.3	7.7	97.4
R33	446	NA()	1.2	17.5457	1.2	0.5149	2.8	0.0656	2.5	0.90	409.3	0.7	7.5	7.7	421.7	1.5	7.3	7.6	491.3	8.9	21.5	23.1	409.3	7.5	83.3
R33	78	NA()	1.5	18.0657	2.7	0.5202	4.7	0.0682	3.8	0.81	425.3	4.0	8.7	8.9	425.3	5.2	8.9	9.1	426.5	25.5	32.3	33.4	425.3	8.7	99.7
R33	154	NA()	1.2	18.2137	2.0	0.5064	4.0	0.0669	3.5	0.87	417.6	3.4	8.3	8.5	416.0	3.9	8.1	8.3	408.3	17.5	26.4	27.8	417.6	8.3	102.3
R33	340	NA()	1.2	17.8764	1.4	0.5042	3.2	0.0654	2.9	0.90	408.4	1.6	7.6	7.8	414.6	2.2	7.4	7.6	450.0	10.7	22.4	24.0	408.4	7.6	90.8
R33	133	NA()	1.3	17.8473	2.1	0.5194	4.1	0.0673	3.6	0.86	419.6	3.6	8.4	8.6	424.7	4.0	8.2	8.5	453.6	16.7	25.8	27.2	419.6	8.4	92.5
R33	110	NA()	1.2	17.9106	2.3	0.5109	4.3	0.0664	3.7	0.85	414.4	3.7	8.4	8.6	419.1	4.3	8.3	8.6	445.7	18.7	27.2	28.5	414.4	8.4	93.0
R33	358	NA()	1.1	17.8330	1.3	0.5274	3.1	0.0682	2.8	0.90	425.5	1.5	7.9	8.1	430.1	2.0	7.5	7.8	455.4	10.0	22.1	23.7	425.5	7.9	93.4
R33	291	NA()	1.6	17.5898	1.4	0.5409	3.3	0.0690	3.0	0.90	430.3	2.1	8.1	8.3	439.0	2.6	7.8	8.1	485.8	11.1	22.5	24.0	430.3	8.1	88.6
R33	40	NA()	1.8	18.2748	3.7	0.5037	4.9	0.0668	3.2	0.66	416.8	4.5	8.8	9.0	414.2	8.2	10.8	11.0	400.8	47.8	51.7	52.4	416.8	8.8	104.0
R33	993	NA()	0.8	18.2669	0.8	0.5620	1.8	0.0745	1.6	0.90	463.1	4.7	9.6	9.8	452.8	4.0	8.6	8.8	401.8	6.0	20.7	22.4	463.1	9.6	115.3
FISH CANYON	145	NA()	2.5	21.3737	7.9	0.0279	8.5	0.0043	3.2	0.38	27.9	0.3	0.6	0.6	28.0	2.0	2.1	2.1	38.3	174.6	175.8	176.1	27.9	0.6	NA
FISH CANYON	125	NA()	1.1	20.2355	8.4	0.0297	9.0	0.0044	3.3	0.36	28.0	0.4	0.6	0.6	29.7	2.3	2.4	2.4	167.7	183.6	184.8	185.0	28.0	0.6	NA
FISH CANYON	492	NA()	0.4	20.6385	3.7	0.0301	4.8	0.0045	3.0	0.64	29.0	0.1	0.6	0.6	30.1	0.6	0.9	0.9	121.4	49.5	53.7	54.5	29.0	0.6	NA
FISH CANYON	134	NA()	0.8	22.6684	7.7	0.0281	8.4	0.0046	3.2	0.39	29.8	0.4	0.7	0.7	28.2	2.0	2.1	2.1	-104.3	174.9	176.3	176.5	29.8	0.7	NA
FISH CANYON	366	NA()	1.6	21.4713	4.2	0.0295	5.2	0.0046	3.1	0.59	29.5	0.2	0.6	0.6	29.5	0.9	1.1	1.1	27.5	70.5	73.7	74.2	29.5	0.6	NA
FISH CANYON	330	NA()	2.0	21.2609	4.4	0.0288	5.3	0.0044	3.1	0.58	28.6	0.2	0.6	0.6	28.8	0.9	1.1	1.1	51.0	75.3	78.2	78.8	28.6	0.6	NA
FISH CANYON	220	NA()	2.2	20.8896	5.3	0.0296	6.1	0.0045	3.1	0.51	28.9	0.3	0.6	0.6	29.6	1.3	1.4	1.4	92.9	102.3	104.4	104.8	28.9	0.6	NA
FISH CANYON	177	NA()	1.6	20.2272	6.0	0.0321	6.8	0.0047	3.2	0.47	30.3	0.3	0.7	0.7	32.1	1.7	1.8	1.8	168.7	121.7	123.5	123.8	30.3	0.7	NA
FISH CANYON	182	NA()	1.0	21.9276	5.3	0.0279	6.2	0.0044	3.2	0.51	28.6	0.3	0.6	0.6	28.0	1.2	1.4	1.4	-23.2	106.0	108.2	108.6	28.6	0.6	NA
FISH CANYON	153	NA()	1.0	20.0404	6.1	0.0298	6.9	0.0043	3.2	0.47	27.9	0.3	0.6	0.6	29.9	1.6	1.7	1.7	190.3	123.9	125.6	125.9	27.9	0.6	NA
FISH CANYON	320	NA()	2.4	21.1022	4.6	0.0279	5.5	0.0043	3.1	0.56	27.4	0.2	0.6	0.6	27.9	1.0	1.1	1.1	68.9	82.1	84.8	85.3	27.4	0.6	NA
FISH CANYON	161	NA()	0.6	20.2153	7.1	0.0302	7.8	0.0044	3.2	0.41	28.5	0.3	0.6	0.6	30.2	1.9	2.0	2.0	170.0	150.0	151.4	151.6	28.5	0.6	NA
FISH CANYON	438	NA()	1.8	20.4478	3.9	0.0287	5.0	0.0043	3.0	0.61	27.4	0.1	0.5	0.5	28.7	0.7	0.9	0.9	143.3	59.1	62.6	63.3	27.4	0.5	NA
FISH CANYON	372	NA()	1.4	21.3879	4.2	0.0294	5.2	0.0046	3.1	0.59	29.3	0.2	0.6	0.6	29.4	0.9	1.1	1.1	36.7	71.4	74.5	75.1	29.3	0.6	NA
ORACLE	125	NA()	1.6	11.1205	1.0	3.1767	2.7	0.2563	2.5	0.93	1471.0	9.3	26.4	27.0	1451.5	6.2	17.2	17.8	1423.9	7.5	18.5	19.9	1423.9	18.5	103.3
ORACLE	147	NA()	1.4	11.2123	0.9	2.9688	2.6	0.2415	2.4	0.93	1394.6	8.3	25.0	25.6	1399.7	5.7	16.8	17.3	1408.2	6.9	18.3	19.7	1408.2	18.3	99.0
ORACLE	222	NA()	0.5	11.0851	0.9	3.0788	2.0	0.2476	1.8	0.90	1426.3	6.7	24.9	25.6	1427.4	4.6	16.5	17.1	1430.0	5.5	17.8	19.2	1430.0	17.8	99.7
ORACLE	118	NA()	1.6	11.2243	1.0	3.0632	2.8	0.2495	2.6	0.93	1435.8	9.2	25.8	26.4	1423.5	6.3	17.1	17.6	1406.1	7.7	18.6	20.0	1406.1	18.6	102.1
ORACLE	149	NA()	1.3	11.0891	0.9	3.0293	2.5	0.2437	2.3	0.93	1406.1	8.2	25.1	25.7	1415.0	5.7	16.8	17.4	1429.3	6.8	18.2	19.6	1429.3	18.2	98.4
ORACLE	104	NA()	2.1	10.9429	1.0	3.2637	2.9	0.2591	2.7	0.94	1485.4	9.5	26.7	27.3	1472.4	6.5	17.4	18.0	1454.6	7.9	18.6	20.0	1454.6	18.6	102.1
ORACLE	111	NA()	2.0	11.1001	1.0	3.1637	2.9	0.2548	2.7	0.93	1463.2	9.2	26.2	26.9	1448.4	6.3	17.2	17.8	1427.4	7.8	18.6	20.0	1427.4	18.6	102.5
ORACLE	127	NA()	1.1	11.2074	1.0	3.1189	2.7	0.2536	2.5	0.93	1457.1	8.8	26.0	26.6	1437.4	6.0	17.0	17.6	1409.0	7.4	18.5	19.9	1409.0	18.5	103.4
ORACLE	93	NA()	1.7	11.2899	1.1	2.9932	3.1	0.2452	2.9	0.93	1413.7	9.7	25.7	26.3	1405.9	6.7	17.2	17.7	1395.0	8.5	19.0	20.4	1395.0	19.0	101.3
ORACLE	177	NA()	1.7	11.1930	0.8	3.0678	2.2	0.2492	2.1	0.93	1434.1	7.7	25.3	26.0	1424.7	5.2	16.7	17.3	1411.5	6.2	18.1	19.5	1411.5	18.1	101.6
ECSTALL	327	NA()	1.4	21.0225	3.2	0.0932	4.9	0.0142	3.7	0.76	91.0	0.5	1.8	1.8	90.5	1.4	2.3	2.3	77.8	36.4	42.1	43.0	91.0	1.8	NA
ECSTALL	218	NA()	2.2	20.8818	3.5	0.0951	4.6	0.0144	3.1	0.67	92.2	0.7	1.9	1.9	92.3	1.7	2.5	2.5	93.8	40.6	45.7	46.6	92.2	1.9	NA
ECSTALL	192	NA()	1.7	20.6281	3.5	0.0994	4.7	0.0149	3.1	0.66	95.2	0.8	1.9	2.0	96.2	1.9	2.7	2.7	122.6	44.5	49.2	50.0	95.2	1.9	NA
ECSTALL	140	NA()	2.0	20.7047	3.8	0.0951	4.9	0.0143	3.2	0.64	91.5	0.9	1.9	2.0	92.3	2.2	2.9	2.9	113.9	54.0	57.9	58.6	91.5	1.9	NA
ECSTALL	132	NA()	1.8	20.0395	4.0	0.1015	5.1	0.0148	3.2	0.62	94.5	0.9	2.0	2.0	98.2	2.7	3.3	3.4	190.4	63.2	66.5	67.1	94.5	2.0	NA
ECSTALL	263	NA()	1.6	20.3529	3.4	0.0974	5.1	0.0144	3.8	0.75	92.1	0.6	1.8	1.9	94.4	1.5	2.4	2.5	154.2	36.8	42.2	43.2	92.1	1.8	NA
ECSTALL	228	NA()	1.9	20.1937	3.5	0.1011	5.2	0.0148	3.9	0.75	94.8	0.7	1.9	1.9	97.8	1.7	2.6	2.7	172.5	40.2	45.2	46.1	94.8	1.9	NA

ECSTALL	228	NA()	1.6	20.9147	3.4	0.0974	5.1	0.0148	3.9	0.75	94.6	0.7	1.9	1.9	94.4	1.6	2.4	2.5	90.1	37.7	43.1	44.1	94.6	1.9	NA
ECSTALL	260	NA()	1.4	20.0349	3.4	0.0967	5.1	0.0141	3.8	0.75	90.0	0.6	1.8	1.8	93.8	1.5	2.4	2.5	191.0	36.6	42.0	42.9	90.0	1.8	NA
ECSTALL	324	NA()	1.2	20.6164	3.1	0.0941	4.9	0.0141	3.8	0.77	90.1	0.4	1.7	1.8	91.3	1.3	2.2	2.3	123.9	32.5	38.6	39.7	90.1	1.7	NA
ECSTALL	258	NA()	1.6	20.4463	3.3	0.0977	5.1	0.0145	3.8	0.75	92.8	0.6	1.8	1.9	94.7	1.5	2.4	2.4	143.4	34.4	40.2	41.2	92.8	1.8	NA
ECSTALL	283	NA()	1.7	21.0219	3.3	0.0984	5.0	0.0150	3.8	0.76	96.1	0.6	1.9	1.9	95.3	1.4	2.4	2.4	77.9	34.9	40.8	41.8	96.1	1.9	NA
ECSTALL	385	NA()	1.5	20.6704	2.8	0.0977	4.6	0.0147	3.7	0.79	93.8	0.3	1.8	1.8	94.6	1.1	2.2	2.2	117.8	27.4	34.5	35.7	93.8	1.8	NA
TEM	175	NA()	1.7	17.9442	1.9	0.4810	3.9	0.0626	3.4	0.87	391.6	2.7	7.6	7.8	398.8	3.3	7.6	7.8	441.5	15.5	25.1	26.5	391.6	7.6	NA
TEM	137	NA()	1.6	17.9875	2.1	0.5052	4.1	0.0659	3.6	0.86	411.7	3.2	8.1	8.3	415.2	3.9	8.1	8.3	436.2	17.9	26.7	28.0	411.7	8.1	94.4
TEM	183	NA()	1.6	18.4102	1.8	0.4954	3.8	0.0662	3.4	0.88	413.1	2.8	8.0	8.2	408.6	3.2	7.7	7.9	384.2	14.5	24.6	26.1	413.1	8.0	107.5
TEM	137	NA()	1.5	18.0910	2.1	0.5222	4.1	0.0685	3.5	0.86	427.4	3.3	8.4	8.6	426.6	4.0	8.3	8.5	423.4	18.1	26.8	28.1	427.4	8.4	100.9
TEM	77	NA()	2.2	17.9099	2.7	0.5110	4.7	0.0664	3.8	0.81	414.5	3.9	8.5	8.7	419.1	5.1	8.8	9.0	445.8	25.0	31.8	32.9	414.5	8.5	93.0
TEM	79	NA()	2.3	17.9659	2.8	0.5035	4.7	0.0656	3.8	0.81	409.8	3.8	8.4	8.6	414.1	5.1	8.7	9.0	438.8	25.9	32.5	33.6	409.8	8.4	93.4
TEM	92	NA()	2.4	17.9445	2.5	0.5064	4.5	0.0659	3.8	0.83	411.6	3.7	8.3	8.5	416.0	4.6	8.5	8.7	441.5	22.2	29.7	30.9	411.6	8.3	93.2
TEM	139	NA()	1.9	18.3181	2.1	0.4877	4.2	0.0648	3.6	0.86	404.9	3.2	8.0	8.2	403.3	3.9	7.9	8.2	395.5	18.7	27.3	28.6	404.9	8.0	102.4
TEM	167	NA()	1.5	18.0550	1.9	0.5299	3.9	0.0694	3.4	0.88	432.7	3.1	8.4	8.6	431.8	3.5	8.1	8.4	427.8	15.2	24.9	26.4	432.7	8.4	101.1
TEM	124	NA()	1.8	17.8627	2.2	0.5048	4.3	0.0654	3.6	0.85	408.5	3.3	8.1	8.3	414.9	4.2	8.2	8.4	451.7	19.6	27.8	29.1	408.5	8.1	90.4
TEM	123	NA()	1.7	17.5202	2.2	0.5311	4.2	0.0675	3.6	0.86	421.2	3.4	8.4	8.6	432.5	4.1	8.4	8.6	494.5	18.0	26.6	27.9	421.2	8.4	85.2
TEM	105	NA()	1.7	17.5025	2.3	0.5316	4.4	0.0675	3.7	0.85	421.1	3.6	8.4	8.6	432.9	4.4	8.5	8.8	496.7	19.6	27.7	28.9	421.1	8.4	84.8
TEM	76	NA()	2.5	17.9299	2.8	0.5178	4.7	0.0674	3.8	0.81	420.2	3.9	8.6	8.7	423.7	5.2	8.9	9.1	443.3	26.2	32.8	33.8	420.2	8.6	94.8
TEM	106	NA()	1.9	18.3662	2.4	0.4861	4.4	0.0648	3.7	0.83	404.6	3.4	8.1	8.3	402.3	4.4	8.2	8.4	389.6	22.2	29.8	31.0	404.6	8.1	103.9
TEM	384	NA()	1.5	18.2182	1.3	0.5077	3.0	0.0671	2.7	0.90	418.7	1.1	7.7	7.9	416.9	1.8	7.3	7.6	407.7	10.0	22.2	23.8	418.7	7.7	102.7
PEIXE	67	NA()	1.6	17.3811	2.5	0.7413	4.5	0.0935	3.8	0.83	576.1	5.3	11.6	11.9	563.2	6.1	10.8	11.1	512.0	22.4	29.6	30.8	576.1	11.6	112.5
PEIXE	62	NA()	1.6	16.7311	2.5	0.7630	4.6	0.0926	3.8	0.83	571.0	5.4	11.6	11.8	575.7	6.2	11.0	11.3	595.2	22.2	29.3	30.5	571.0	11.6	95.9
PEIXE	60	NA()	1.5	16.7299	2.5	0.7624	4.6	0.0925	3.8	0.83	570.6	5.4	11.6	11.8	575.4	6.2	11.0	11.3	595.4	21.9	29.1	30.3	570.6	11.6	95.8
PEIXE	61	NA()	1.6	16.4642	2.6	0.7362	4.6	0.0880	3.8	0.83	543.4	5.2	11.1	11.3	560.2	6.3	10.9	11.2	629.9	23.3	30.1	31.2	543.4	11.1	86.3
PEIXE	87	NA()	1.3	17.2549	2.2	0.7414	4.2	0.0928	3.6	0.86	572.2	5.0	11.4	11.7	563.2	5.4	10.5	10.8	528.1	19.0	27.2	28.4	572.2	11.4	108.4
PEIXE	77	NA()	1.5	17.2977	2.3	0.7329	4.4	0.0920	3.7	0.85	567.3	5.1	11.4	11.6	558.2	5.5	10.5	10.8	522.6	19.2	27.3	28.6	567.3	11.4	108.5
PEIXE	83	NA()	1.3	16.8884	2.2	0.7639	4.3	0.0936	3.7	0.86	576.9	5.1	11.5	11.8	576.3	5.4	10.6	10.9	574.9	17.5	26.0	27.3	576.9	11.5	100.3
PEIXE	105	NA()	1.3	17.0362	2.0	0.7527	4.1	0.0930	3.6	0.87	573.5	4.8	11.4	11.6	569.8	5.0	10.3	10.7	555.9	16.2	25.2	26.5	573.5	11.4	103.2
PEIXE	99	NA()	5.3	16.8080	2.0	0.7354	4.1	0.0897	3.6	0.87	553.7	4.8	11.0	11.3	559.7	5.1	10.3	10.6	585.3	16.9	25.6	26.9	553.7	11.0	94.6
PEIXE	112	NA()	6.6	16.8499	1.9	0.7629	4.0	0.0933	3.5	0.88	574.9	4.8	11.4	11.6	575.7	5.0	10.4	10.7	579.8	16.0	25.0	26.4	574.9	11.4	99.1
PEIXE	80	NA()	10.6	17.0184	2.4	0.7155	4.4	0.0884	3.7	0.85	545.8	5.2	11.1	11.3	548.0	5.7	10.5	10.8	558.2	20.0	27.8	29.0	545.8	11.1	97.8
PEIXE	199	NA()	3.6	17.0336	1.5	0.7200	3.4	0.0890	3.1	0.90	549.5	3.8	10.6	10.8	550.7	3.8	9.6	9.9	556.3	11.5	22.5	24.0	549.5	10.6	98.8
PEIXE	189	NA()	3.8	16.9542	1.5	0.7429	3.5	0.0914	3.1	0.90	563.8	4.0	10.9	11.1	564.1	3.9	9.8	10.1	566.5	11.4	22.4	23.9	563.8	10.9	99.5
PEIXE	171	NA()	3.6	16.8233	1.5	0.7685	3.5	0.0938	3.2	0.90	578.0	4.3	11.2	11.5	578.9	4.2	10.1	10.4	583.3	11.8	22.5	24.0	578.0	11.2	99.1
PEIXE	186	NA()	3.6	16.9041	1.5	0.7447	3.5	0.0913	3.2	0.90	563.5	4.0	10.9	11.2	565.2	4.0	9.8	10.2	572.9	11.4	22.4	23.9	563.5	10.9	98.4
PEIXE	31	NA()	3.2	16.4276	3.4	0.7420	4.7	0.0884	3.2	0.69	546.3	6.1	11.5	11.8	563.6	8.4	12.3	12.5	634.8	33.3	38.4	39.3	546.3	11.5	86.1

Nu Plasma HR (LA-MC-ICP-MS)							Isotope ratios					Apparent ages (Ma)													
Analysis	U (ppm)	206Pb 204Pb	U/Th	206Pb* 207Pb*	± (%)	207Pb* 235U*	± (%)	206Pb* 238U	± (%)	error corr.	206Pb* 238U*	±1s(a) (Ma)	±1s(b) (Ma)	±1s(c) (Ma)	207Pb* 235U	±1s(a) (Ma)	±1s(b) (Ma)	±1s(c) (Ma)	206Pb* 207Pb*	±1s(a) (Ma)	±1s(b) (Ma)	±1s(c) (Ma)	Best age (Ma)	± (Ma)	Conc (%)
49127	138	NA()	0.9	20.5018	3.5	0.1443	4.7	0.0215	3.2	0.67	136.9	1.3	2.9	2.9	136.8	2.6	3.7	3.8	137.1	42.2	47.1	47.9	136.9	2.9	NA
49127	140	NA()	0.9	20.6480	3.4	0.1423	4.7	0.0213	3.2	0.68	136.0	1.3	2.9	2.9	135.1	2.4	3.6	3.7	120.4	39.1	44.3	45.2	136.0	2.9	NA
49127	127	NA()	0.8	20.0314	3.6	0.1552	4.8	0.0226	3.2	0.66	143.8	1.4	3.0	3.1	146.5	3.1	4.2	4.3	191.4	46.9	51.2	52.0	143.8	3.0	NA
49127	163	NA()	1.2	20.9590	3.5	0.1410	5.2	0.0214	3.9	0.75	136.8	1.2	2.8	2.9	133.9	2.5	3.6	3.7	85.1	41.2	46.2	47.2	136.8	2.8	NA
49127	123	NA()	1.7	19.7900	3.6	0.1501	4.8	0.0215	3.2	0.66	137.4	1.4	2.9	3.0	142.0	2.9	4.0	4.1	219.5	44.7	49.2	50.0	137.4	2.9	NA
49127	60	NA()	1.7	20.8484	4.4	0.1421	5.5	0.0215	3.2	0.59	137.1	1.6	3.0	3.1	134.9	4.4	5.1	5.1	97.5	76.7	79.5	80.0	137.1	3.0	NA
49127	481	NA()	1.4	20.4117	2.0	0.1488	3.9	0.0220	3.4	0.86	140.5	0.3	2.6	2.7	140.8	1.1	2.9	3.0	147.4	18.2	27.6	29.0	140.5	2.6	NA
49127	554	NA()	1.7	19.7383	1.9	0.1507	3.8	0.0216	3.3	0.87	137.6	0.3	2.6	2.6	142.5	0.9	2.9	3.0	225.5	15.8	25.9	27.4	137.6	2.6	NA
49127	148	NA()	0.8	17.6287	4.7	0.1660	6.8	0.0212	5.0	0.73	135.4	4.3	5.0	5.0	155.9	7.0	7.6	7.6	480.9	80.1	82.4	82.9	135.4	5.0	NA
49127	151	NA()	1.0	20.3057	3.6	0.1448	5.3	0.0213	3.9	0.74	136.1	1.2	2.8	2.9	137.3	2.8	3.8	3.9	159.6	45.8	50.3	51.1	136.1	2.8	NA
49127	241	NA()	1.4	17.5222	2.6	0.1710	4.6	0.0217	3.7	0.82	138.7	0.9	2.7	2.8	160.3	1.9	3.6	3.7	494.3	23.6	30.6	31.8	138.7	2.7	NA
49127	239	NA()	1.0	20.6754	3.0	0.1367	4.8	0.0205	3.8	0.79	130.9	0.9	2.6	2.7	130.1	1.7	3.0	3.1	117.2	28.3	35.2	36.3	130.9	2.6	NA
49127	863	NA()	1.5	20.5710	1.6	0.1497	3.4	0.0223	3.1	0.89	142.5	1.2	2.9	3.0	141.7	1.3	3.0	3.1	129.2	12.9	24.5	26.1	142.5	2.9	NA
49127	161	NA()	1.0	19.9964	3.4	0.1441	5.1	0.0209	3.9	0.76	133.4	1.2	2.7	2.8	136.7	2.3	3.5	3.6	195.4	35.3	40.9	41.8	133.4	2.7	NA
91500	48	NA()	2.3	13.3834	1.9	1.8719	4.1	0.1818	3.6	0.89	1076.7	9.4	20.9	21.3	1071.2	8.0	15.9	16.4	1061.0	15.1	23.4	24.6	1061.0	23.4	101.5
91500	46	NA()	2.5	13.0259	1.9	1.9074	4.1	0.1803	3.7	0.89	1068.5	9.5	20.8	21.2	1083.7	8.1	16.0	16.5	1115.2	14.6	23.0	24.2	1115.2	23.0	95.8
91500	48	NA()	2.4	13.1718	1.9	1.8653	4.1	0.1783	3.6	0.89	1057.5	9.4	20.6	21.0	1068.9	8.1	15.9	16.4	1092.9	14.8	23.1	24.4	1092.9	23.1	96.8
91500	49	NA()	2.5	13.2158	1.9	1.8799	4.1	0.1803	3.6	0.89	1068.5	9.5	20.8	21.2	1074.0	8.0	15.9	16.4	1086.3	14.3	22.8	24.1	1086.3	22.8	98.4
91500	47	NA()	2.5	13.2587	1.9	1.8211	4.1	0.1752	3.7	0.89	1040.7	9.4	20.4	20.8	1053.1	8.1	15.8	16.3	1079.7	14.8	23.1	24.4	1079.7	23.1	96.4
91500	53	NA()	2.4	13.3786	1.8	1.8683	4.0	0.1814	3.6	0.89	1074.4	9.6	20.9	21.4	1069.9	7.9	15.8	16.3	1061.7	13.9	22.7	23.9	1061.7	22.7	101.2
91500	53	NA()	2.4	13.3828	1.8	1.8103	4.0	0.1758	3.6	0.89	1043.9	9.4	20.4	20.8	1049.2	7.8	15.7	16.1	1061.1	14.1	22.7	24.0	1061.1	22.7	98.4
91500	54	NA()	2.4	13.0903	1.8	1.9064	4.0	0.1811	3.6	0.89	1072.8	9.6	20.9	21.3	1083.3	7.9	15.9	16.4	1105.4	13.8	22.4	23.7	1105.4	22.4	97.1
91500	52	NA()	2.4	13.2783	1.9	1.7905	4.1	0.1725	3.6	0.89	1025.9	9.2	20.0	20.5	1042.0	7.9	15.6	16.1	1076.8	14.5	22.9	24.2	1076.8	22.9	95.3
91500	54	NA()	2.5	13.7656	1.9	1.7964	4.0	0.1794	3.6	0.89	1063.9	9.5	20.7	21.2	1044.2	7.9	15.7	16.1	1004.1	14.8	23.3	24.6	1004.1	23.3	106.0
91500	51	NA()	2.2	13.6172	1.9	1.7874	4.0	0.1766	3.6	0.89	1048.4	9.4	20.4	20.9	1040.9	7.8	15.6	16.1	1026.0	14.4	23.0	24.3	1026.0	23.0	102.2
91500	51	NA()	2.4	13.4806	1.9	1.7888	4.1	0.1750	3.6	0.89	1039.4	9.4	20.3	20.8	1041.4	7.9	15.6	16.1	1046.4	14.2	22.8	24.1	1046.4	22.8	99.3
91500	55	NA()	2.5	13.3399	1.8	1.8754	4.0	0.1815	3.6	0.89	1075.3	9.7	21.0	21.4	1072.4	8.0	15.9	16.4	1067.5	14.3	22.9	24.1	1067.5	22.9	100.7
91500	49	NA()	2.3	13.1234	1.9	1.8170	4.1	0.1730	3.6	0.89	1028.7	9.4	20.2	20.6	1051.6	8.1	15.8	16.3	1100.3	15.0	23.2	24.5	1100.3	23.2	93.5
91500	48	NA()	2.3	13.2611	1.9	1.8651	4.1	0.1795	3.7	0.89	1064.0	9.7	20.8	21.2	1068.8	8.2	16.0	16.4	1079.4	15.0	23.2	24.5	1079.4	23.2	98.6
94-35	38	NA()	2.1	18.6696	11.2	0.0627	11.8	0.0085	3.5	0.30	54.5	0.9	1.4	1.4	61.7	6.6	6.7	6.7	352.7	245.1	245.9	246.1	54.5	1.4	NA
94-35	48	NA()	1.3	20.7923	9.9	0.0589	10.5	0.0089	3.4	0.32	57.0	0.9	1.4	1.4	58.1	5.4	5.5	5.5	103.9	223.1	224.1	224.3	57.0	1.4	NA
94-35	45	NA()	1.3	17.6558	8.8	0.0676	9.4	0.0087	3.4	0.36	55.6	0.9	1.4	1.4	66.4	5.4	5.6	5.6	477.5	182.8	183.8	184.0	55.6	1.4	NA
94-35	54	NA()	1.3	20.0940	9.2	0.0596	9.8	0.0087	3.4	0.35	55.7	0.9	1.4	1.4	58.8	5.1	5.2	5.2	184.1	203.1	204.2	204.3	55.7	1.4	NA
94-35	45	NA()	1.6	20.1604	10.7	0.0605	11.2	0.0088	3.4	0.30	56.8	0.9	1.4	1.4	59.6	6.0	6.1	6.2	176.4	240.6	241.5	241.7	56.8	1.4	NA
94-35	58	NA()	1.3	19.9592	10.1	0.0602	10.6	0.0087	3.4	0.32	55.9	0.9	1.4	1.4	59.3	5.6	5.7	5.7	199.8	223.5	224.5	224.7	55.9	1.4	NA
94-35	39	NA()	1.5	18.8998	10.8	0.0609	11.4	0.0084	3.5	0.31	53.6	1.0	1.4	1.4	60.0	6.1	6.3	6.3	324.9	236.3	237.2	237.3	53.6	1.4	NA
94-35	55	NA()	1.2	18.6946	7.3	0.0658	8.0	0.0089	3.4	0.42	57.3	0.9	1.4	1.4	64.7	4.3	4.5	4.5	349.7	150.6	151.9	152.2	57.3	1.4	NA
94-35	54	NA()	1.2	21.1958	7.5	0.0576	8.2	0.0089	3.4	0.41	56.8	0.9	1.4	1.4	56.9	3.9	4.1	4.1	58.3	163.9	165.3	165.5	56.8	1.4	NA
94-35	60	NA()	1.4	20.6232	7.6	0.0573	8.3	0.0086	3.3	0.40	55.1	0.8	1.3	1.3	56.6	3.9	4.1	4.1	123.2	163.7	165.0	165.3	55.1	1.3	NA
94-35	45	NA()	2.0	18.3441	8.9	0.0637	9.6	0.0085	3.5	0.36	54.4	0.9	1.4	1.4	62.7	5.2	5.4	5.4	392.3	188.8	189.8	190.0	54.4	1.4	NA
94-35	54	NA()	1.2	21.6445	10.1	0.0571	10.7	0.0090	3.3	0.31	57.6	0.8	1.4	1.4	56.4	5.4	5.5	5.5	8.1	233.0	234.0	234.2	57.6	1.4	NA
94-35	34	NA()	1.7	20.8054	16.1	0.0563	16.5	0.0085	3.5	0.21	54.6	1.0	1.4	1.5	55.6	8.6	8.7	8.7	102.5	376.7	377.4	377.5	54.6	1.4	NA
94-35	40	NA()	2.1	21.2546	10.2	0.0573	10.8	0.0088	3.4	0.32	56.7	0.9	1.4	1.4	56.6	5.5	5.6	5.6	51.7	234.4	235.4	235.5	56.7	1.4	NA
94-35	47	NA()	1.6	20.2241	11.5	0.0578	12.0	0.0085	3.4	0.28	54.4	0.9	1.3	1.4	57.0	6.2	6.3	6.3	169.0	259.3	260.2	260.3	54.4	1.3	NA
94-35	64	NA()	2.1	20.1658	7.0	0.0599	7.7	0.0088	3.3	0.43	56.3	0.8	1.3	1.4	59.1	3.7	3.9	3.9	175.7	147.3	148.8	149.1	56.3	1.3	NA
94-35	45	NA()	2.3	19.9166	12.3	0.0580	12.8	0.0084	3.4	0.27	53.8	0.9	1.3	1.4	57.3	6.7	6.8	6.8	204.7	278.5	279.2	279.4	53.8	1.3	NA
PLES	608	NA()	7.2	18.6472	1.1	0.3969	2.6	0.0537	2.4	0.90	337.2	0.8	6.2	6.4	339.4	1.3	6.1	6.4	355.5	8.6	21.8	23.4	337.2	6.2	NA
PLES	609	NA()	7.2	18.5029	1.1	0.4050	2.6	0.0544	2.3	0.90	341.3	0.8	6.3	6.5	345.3	1.3	6.2	6.4	372.9	8.5	21.7	23.3	341.3	6.3	NA
PLES	385	NA()	9.4	18.5215	1.4	0.3939	3.2	0.0529	2.9	0.90	332.6	1.0	6.2	6.3	337.2	1.6	6.2	6.4	370.7	10.8	22.7	24.3	332.6	6.2	NA
PLES	927	NA()	6.1	18.4207	0.9	0.4084	2.2	0.0546	2.0	0.90	342.6	3.1	7.0	7.1	347.7	2.9	6.8	7.0	382.9	7.0	21.1	22.8	342.6	7.0	NA
PLES	706	NA()	5.5	18.6849	1.1	0.3954	2.5	0.0536	2.2	0.90	336.6	1.4	6.3	6.5	338.3	1.6	6.2	6.4	350.9	8.1	21.6	23.3	336.6	6.3	NA
PLES	873	NA()	5.7	18.4730	1.0	0.3982	2.2	0.0534	2.0	0.90	335.2	2.6	6.7	6.8	340.3	2.5	6.5	6.7	376.6	7.3	21.2	22.9	335.2	6.7	NA
PLES	705	NA()	5.3	18.5856	1.1	0.3984	2.5	0.0537	2.2	0.90	337.3	1.4	6.3	6.5	340.5	1.6	6.2	6.4	362.9	8.0	21.5	23.2	337.3	6.3	NA
PLES	663	NA()	5.1	19.0126	1.1	0.3940	~~2.5~~	0.0544	2.2	0.90	341.2	1.2	6.3	6.5	337.3	1.5	6.1	6.4	311.4	8.4	21.9	23.5	341.2	6.3	NA
PLES	661	NA()	5.2	18.7005	1.1	0.4098	2.5	0.0556	2.2	0.90	348.8	1.2	6.5	6.6	348.7	1.5	6.3	6.5	349.0	8.1	21.6	23.3	348.8	6.5	NA
PLES	642	NA()	7.2	18.5949	1.1	0.4096	2.6	0.0553	2.3	0.90	346.8	1.0	6.4	6.6	348.6	1.4	6.3	6.5	361.8	8.4	21.7	23.4	346.8	6.4	NA
FC	258	NA()	1.5	13.1996	0.8	1.8845	2.2	0.1805	2.0	0.92	1069.6	5.0	19.2	19.6	1075.6	3.9	14.3	14.8	1088.7	5.9	18.7	20.2	1088.7	18.7	98.2
FC	224	NA()	1.2	13.1319	0.9	1.9224	2.4	0.1832	2.2	0.92	1084.3	5.7	19.6	20.1	1088.9	4.4	14.5	15.0	1099.0	6.4	18.9	20.4	1099.0	18.9	98.7
FC	217	NA()	1.2	13.1100	0.9	1.8548	2.4	0.1764	2.3	0.92	1047.5	5.7	19.0	19.5	1065.1	4.4	14.4	14.9	1102.4	6.5	18.9	20.4	1102.4	18.9	95.0
FC	212	NA()	1.2	13.4011	0.9	1.8885	2.4	0.1836	2.2	0.92	1086.8	6.0	19.7	20.2	1077.1	4.6	14.5	15.0	1058.3	6.7	19.0	20.5	1058.3	19.0	102.7
FC	186	NA()	1.1	13.2129	1.0	1.8938	2.6	0.1816	2.4	0.93	1075.5	6.5	19.7	20.2	1078.9	5.0	14.7	15.2	1086.7	7.2	19.2	20.6	1086.7	19.2	99.0
FC	188	NA()	1.2	13.2978	1.0	1.9395	2.6	0.1871	2.4	0.93	1105.9	6.6	20.2	20.7	1094.8	4.9	14.8	15.3	1073.9	7.0	19.1	20.6	1073.9	19.1	103.0
FC	114	NA()	1.9	13.1949	1.2	1.9554	3.2	0.1872	3.0	0.92	1106.2	8.0	20.7	21.2	1100.3	6.1	15.2	15.7	1089.5	9.1	19.9	21.4	1089.5	19.9	101.5
FC	160	NA()	1.3	13.0265	1.1	1.9636	2.8	0.1856	2.6	0.93	1097.5	7.0	20.2	20.7	1103.1	5.4	15.0	15.5	1115.1	7.7	19.3	20.7	1115.1	19.3	98.4
FC	741	NA()	1.2	13.1190	0.8	1.9727	1.4	0.1878	1.1	0.80	1109.4	4.3	19.6	20.1	1106.2	3.1	14.3	14.8	1101.0	3.7	18.1	19.7	1101.0	18.1	100.8
FC	957	NA()	0.9	13.1312	0.8	1.9503	1.5	0.1858	1.3	0.84	1098.7	8.4	20.7	21.2	1098.5	5.8	15.1	15.6	1099.1	4.7	18.4	19.9	1099.1	18.4	100.0
FC	972	NA()	1.0	13.2448	0.8	2.0364	1.6	0.1957	1.3	0.85	1152.2	9.2	21.8	22.3	1127.8	6.2	15.4	15.9	1081.9	5.0	18.5	20.0	1081.9	18.5	106.5
FC	210	NA()	0.4	13.0963	0.9	1.9512	2.5	0.1854	2.3	0.92	1096.5	6.1	19.9	20.4	1098.9	4.6	14.7	15.2	1104.5	6.7	18.9	20.4	1104.5	18.9	99.3
FC	180	NA()	0.3	13.2466	1.0	1.9500	2.7	0.1874	2.5	0.93	1107.4	6.8	20.3	20.7	1098.4	5.1	14.8	15.3	1081.6	7.2	19.2	20.7	1081.6	19.2	102.4
FC	1157	NA()	0.9	13.2798	0.9	1.9501	1.8	0.1879	1.6	0.88	1110.0	12.5	22.8	23.3	1098.5	8.5	16.3	16.8	1076.6	6.4	18.9	20.4	1076.6	18.9	103.1
FC	137	NA()	1.4	13.1074	1.1	1.9484	3.0	0.1853	2.8	0.93	1095.9	7.6	20.4	20.8	1097.9	5.7	15.1	15.6	1102.8	8.3	19.6	21.0	1102.8	19.6	99.4
R33	80	NA()	1.7	17.8501	2.7	0.5075	4.7	0.0657	3.8	0.81	410.4	4.1	8.5	8.7	416.8	5.4	8.9	9.1	453.2	26.4	33.0	34.0	410.4	8.5	90.5
R33	76	NA()	1.2	18.1222	2.8	0.5179	4.7	0.0681	3.8	0.81	424.7	4.2	8.8	9.0	423.7	5.3	8.9	9.2	419.5	25.4	32.2	33.3	424.7	8.8	101.2
R33	624	NA()	1.1	18.0138	1.0	0.5126	2.4	0.0670	2.1	0.90	418.0	1.1	7.7	7.9	420.2	1.5	7.3	7.6	432.9	7.7	21.2	22.9	418.0	7.7	96.6
R33	665	NA()	1.2	18.0212	1.0	0.5202	2.2	0.0680	1.9	0.89	424.3	1.4	7.8	8.0	425.3	1.7	7.4	7.7	432.0	7.2	21.0	22.7	424.3	7.8	98.2
R33	471	NA()	1.6	18.0097	1.2	0.5180	2.7	0.0677	2.4	0.90	422.3	0.5	7.7	7.9	423.8	1.4	7.3	7.6	433.4	8.7	21.6	23.2	422.3	7.7	97.4
R33	446	NA()	1.2	17.5457	1.2	0.5149	2.8	0.0656	2.5	0.90	409.3	0.7	7.5	7.7	421.7	1.5	7.3	7.6	491.3	8.9	21.5	23.1	409.3	7.5	83.3
R33	78	NA()	1.5	18.0657	2.7	0.5202	4.7	0.0682	3.8	0.81	425.3	4.0	8.7	8.9	425.3	5.2	8.9	9.1	426.5	25.5	32.3	33.4	425.3	8.7	99.7
R33	154	NA()	1.2	18.2137	2.0	0.5064	4.0	0.0669	3.5	0.87	417.6	3.4	8.3	8.5	416.0	3.9	8.1	8.3	408.3	17.5	26.4	27.8	417.6	8.3	102.3
R33	340	NA()	1.2	17.8764	1.4	0.5042	3.2	0.0654	2.9	0.90	408.4	1.6	7.6	7.8	414.6	2.2	7.4	7.6	450.0	10.7	22.4	24.0	408.4	7.6	90.8
R33	133	NA()	1.3	17.8473	2.1	0.5194	4.1	0.0673	3.6	0.86	419.6	3.6	8.4	8.6	424.7	4.0	8.2	8.5	453.6	16.7	25.8	27.2	419.6	8.4	92.5
R33	110	NA()	1.2	17.9106	2.3	0.5109	4.3	0.0664	3.7	0.85	414.4	3.7	8.4	8.6	419.1	4.3	8.3	8.6	445.7	18.7	27.2	28.5	414.4	8.4	93.0
R33	358	NA()	1.1	17.8330	1.3	0.5274	3.1	0.0682	2.8	0.90	425.5	1.5	7.9	8.1	430.1	2.0	7.5	7.8	455.4	10.0	22.1	23.7	425.5	7.9	93.4
R33	291	NA()	1.6	17.5898	1.4	0.5409	3.3	0.0690	3.0	0.90	430.3	2.1	8.1	8.3	439.0	2.6	7.8	8.1	485.8	11.1	22.5	24.0	430.3	8.1	88.6
R33	40	NA()	1.8	18.2748	3.7	0.5037	4.9	0.0668	3.2	0.66	416.8	4.5	8.8	9.0	414.2	8.2	10.8	11.0	400.8	47.8	51.7	52.4	416.8	8.8	104.0
R33	993	NA()	0.8	18.2669	0.8	0.5620	1.8	0.0745	1.6	0.90	463.1	4.7	9.6	9.8	452.8	4.0	8.6	8.8	401.8	6.0	20.7	22.4	463.1	9.6	115.3

FISH CANYON	145	NA()	2.5	21.3737	7.9	0.0279	8.5	0.0043	3.2	0.38	27.9	0.3	0.6	0.6	28.0	2.0	2.1	2.1	38.3	174.6	175.8	176.1	27.9	0.6	NA
FISH CANYON	125	NA()	1.1	20.2355	8.4	0.0297	9.0	0.0044	3.3	0.36	28.0	0.4	0.6	0.6	29.7	2.3	2.4	2.4	167.7	183.6	184.8	185.0	28.0	0.6	NA
FISH CANYON	492	NA()	0.4	20.6385	3.7	0.0301	4.8	0.0045	3.0	0.64	29.0	0.1	0.6	0.6	30.1	0.6	0.9	0.9	121.4	49.5	53.7	54.5	29.0	0.6	NA
FISH CANYON	134	NA()	0.8	22.6684	7.7	0.0281	8.4	0.0046	3.2	0.39	29.8	0.4	0.7	0.7	28.2	2.0	2.1	2.1	-104.3	174.9	176.3	176.5	29.8	0.7	NA
FISH CANYON	366	NA()	1.6	21.4713	4.2	0.0295	5.2	0.0046	3.1	0.59	29.5	0.2	0.6	0.6	29.5	0.9	1.1	1.1	27.5	70.5	73.7	74.2	29.5	0.6	NA
FISH CANYON	330	NA()	2.0	21.2609	4.4	0.0288	5.3	0.0044	3.1	0.58	28.6	0.2	0.6	0.6	28.8	0.9	1.1	1.1	51.0	75.3	78.2	78.8	28.6	0.6	NA
FISH CANYON	220	NA()	2.2	20.8896	5.3	0.0296	6.1	0.0045	3.1	0.51	28.9	0.3	0.6	0.6	29.6	1.3	1.4	1.4	92.9	102.3	104.4	104.8	28.9	0.6	NA
FISH CANYON	177	NA()	1.6	20.2272	6.0	0.0321	6.8	0.0047	3.2	0.47	30.3	0.3	0.7	0.7	32.1	1.7	1.8	1.8	168.7	121.7	123.5	123.8	30.3	0.7	NA
FISH CANYON	182	NA()	1.0	21.9276	5.3	0.0279	6.2	0.0044	3.2	0.51	28.6	0.3	0.6	0.6	28.0	1.2	1.4	1.4	-23.2	106.0	108.2	108.6	28.6	0.6	NA
FISH CANYON	153	NA()	1.0	20.0404	6.1	0.0298	6.9	0.0043	3.2	0.47	27.9	0.3	0.6	0.6	29.9	1.6	1.7	1.7	190.3	123.9	125.6	125.9	27.9	0.6	NA
FISH CANYON	320	NA()	2.4	21.1022	4.6	0.0279	5.5	0.0043	3.1	0.56	27.4	0.2	0.6	0.6	27.9	1.0	1.1	1.1	68.9	82.1	84.8	85.3	27.4	0.6	NA
FISH CANYON	161	NA()	0.6	20.2153	7.1	0.0302	7.8	0.0044	3.2	0.41	28.5	0.3	0.6	0.6	30.2	1.9	2.0	2.0	170.0	150.0	151.4	151.6	28.5	0.6	NA
FISH CANYON	438	NA()	1.8	20.4478	3.9	0.0287	5.0	0.0043	3.0	0.61	27.4	0.1	0.5	0.5	28.7	0.7	0.9	0.9	143.3	59.1	62.6	63.3	27.4	0.5	NA
FISH CANYON	372	NA()	1.4	21.3879	4.2	0.0294	5.2	0.0046	3.1	0.59	29.3	0.2	0.6	0.6	29.4	0.9	1.1	1.1	36.7	71.4	74.5	75.1	29.3	0.6	NA
ORACLE	125	NA()	1.6	11.1205	1.0	3.1767	2.7	0.2563	2.5	0.93	1471.0	9.3	26.4	27.0	1451.5	6.2	17.2	17.8	1423.9	7.5	18.5	19.9	1423.9	18.5	103.3
ORACLE	147	NA()	1.4	11.2123	0.9	2.9688	2.6	0.2415	2.4	0.93	1394.6	8.3	25.0	25.6	1399.7	5.7	16.8	17.3	1408.2	6.9	18.3	19.7	1408.2	18.3	99.0
ORACLE	222	NA()	0.5	11.0851	0.9	3.0788	2.0	0.2476	1.8	0.90	1426.3	6.7	24.9	25.6	1427.4	4.6	16.5	17.1	1430.0	5.5	17.8	19.2	1430.0	17.8	99.7
ORACLE	118	NA()	1.6	11.2243	1.0	3.0632	2.8	0.2495	2.6	0.93	1435.8	9.2	25.8	26.4	1423.5	6.3	17.1	17.6	1406.1	7.7	18.6	20.0	1406.1	18.6	102.1
ORACLE	149	NA()	1.3	11.0891	0.9	3.0293	2.5	0.2437	2.3	0.93	1406.1	8.2	25.1	25.7	1415.0	5.7	16.8	17.4	1429.3	6.8	18.2	19.6	1429.3	18.2	98.4
ORACLE	104	NA()	2.1	10.9429	1.0	3.2637	2.9	0.2591	2.7	0.94	1485.4	9.5	26.7	27.3	1472.4	6.5	17.4	18.0	1454.6	7.9	18.6	20.0	1454.6	18.6	102.1
ORACLE	111	NA()	2.0	11.1001	1.0	3.1637	2.9	0.2548	2.7	0.93	1463.2	9.2	26.2	26.9	1448.4	6.3	17.2	17.8	1427.4	7.8	18.6	20.0	1427.4	18.6	102.5
ORACLE	127	NA()	1.1	11.2074	1.0	3.1189	2.7	0.2536	2.5	0.93	1457.1	8.8	26.0	26.6	1437.4	6.0	17.0	17.6	1409.0	7.4	18.5	19.9	1409.0	18.5	103.4
ORACLE	93	NA()	1.7	11.2899	1.1	2.9932	3.1	0.2452	2.9	0.93	1413.7	9.7	25.7	26.3	1405.9	6.7	17.2	17.7	1395.0	8.5	19.0	20.4	1395.0	19.0	101.3
ORACLE	177	NA()	1.7	11.1930	0.8	3.0678	2.2	0.2492	2.1	0.93	1434.1	7.7	25.3	26.0	1424.7	5.2	16.7	17.3	1411.5	6.2	18.1	19.5	1411.5	18.1	101.6
ECSTALL	327	NA()	1.4	21.0225	3.2	0.0932	4.9	0.0142	3.7	0.76	91.0	0.5	1.8	1.8	90.5	1.4	2.3	2.3	77.8	36.4	42.1	43.0	91.0	1.8	NA
ECSTALL	218	NA()	2.2	20.8818	3.5	0.0951	4.6	0.0144	3.1	0.67	92.2	0.7	1.9	1.9	92.3	1.7	2.5	2.5	93.8	40.6	45.7	46.6	92.2	1.9	NA
ECSTALL	192	NA()	1.7	20.6281	3.5	0.0994	4.7	0.0149	3.1	0.66	95.2	0.8	1.9	2.0	96.2	1.9	2.7	2.7	122.6	44.5	49.2	50.0	95.2	1.9	NA
ECSTALL	140	NA()	2.0	20.7047	3.8	0.0951	4.9	0.0143	3.2	0.64	91.5	0.9	1.9	2.0	92.3	2.2	2.9	2.9	113.9	54.0	57.9	58.6	91.5	1.9	NA
ECSTALL	132	NA()	1.8	20.0395	4.0	0.1015	5.1	0.0148	3.2	0.62	94.5	0.9	2.0	2.0	98.2	2.7	3.3	3.4	190.4	63.2	66.5	67.1	94.5	2.0	NA
ECSTALL	263	NA()	1.6	20.3529	3.4	0.0974	5.1	0.0144	3.8	0.75	92.1	0.6	1.8	1.9	94.4	1.5	2.4	2.5	154.2	36.8	42.2	43.2	92.1	1.8	NA
ECSTALL	228	NA()	1.9	20.1937	3.5	0.1011	5.2	0.0148	3.9	0.75	94.8	0.7	1.9	1.9	97.8	1.7	2.6	2.7	172.5	40.2	45.2	46.1	94.8	1.9	NA
ECSTALL	228	NA()	1.6	20.9147	3.4	0.0974	5.1	0.0148	3.9	0.75	94.6	0.7	1.9	1.9	94.4	1.6	2.4	2.5	90.1	37.7	43.1	44.1	94.6	1.9	NA
ECSTALL	260	NA()	1.4	20.0349	3.4	0.0967	5.1	0.0141	3.8	0.75	90.0	0.6	1.8	1.8	93.8	1.5	2.4	2.5	191.0	36.6	42.0	42.9	90.0	1.8	NA
ECSTALL	324	NA()	1.2	20.6164	3.1	0.0941	4.9	0.0141	3.8	0.77	90.1	0.4	1.7	1.8	91.3	1.3	2.2	2.3	123.9	32.5	38.6	39.7	90.1	1.7	NA
ECSTALL	258	NA()	1.6	20.4463	3.3	0.0977	5.1	0.0145	3.8	0.75	92.8	0.6	1.8	1.9	94.7	1.5	2.4	2.4	143.4	34.4	40.2	41.2	92.8	1.8	NA
ECSTALL	283	NA()	1.7	21.0219	3.3	0.0984	5.0	0.0150	3.8	0.76	96.1	0.6	1.9	1.9	95.3	1.4	2.4	2.4	77.9	34.9	40.8	41.8	96.1	1.9	NA
ECSTALL	385	NA()	1.5	20.6704	2.8	0.0977	4.6	0.0147	3.7	0.79	93.8	0.3	1.8	1.8	94.6	1.1	2.2	2.2	117.8	27.4	34.5	35.7	93.8	1.8	NA
TEM	175	NA()	1.7	17.9442	1.9	0.4810	3.9	0.0626	3.4	0.87	391.6	2.7	7.6	7.8	398.8	3.3	7.6	7.8	441.5	15.5	25.1	26.5	391.6	7.6	NA
TEM	137	NA()	1.6	17.9875	2.1	0.5052	4.1	0.0659	3.6	0.86	411.7	3.2	8.1	8.3	415.2	3.9	8.1	8.3	436.2	17.9	26.7	28.0	411.7	8.1	94.4
TEM	183	NA()	1.6	18.4102	1.8	0.4954	3.8	0.0662	3.4	0.88	413.1	2.8	8.0	8.2	408.6	3.2	7.7	7.9	384.2	14.5	24.6	26.1	413.1	8.0	107.5
TEM	137	NA()	1.5	18.0910	2.1	0.5222	4.1	0.0685	3.5	0.86	427.4	3.3	8.4	8.6	426.6	4.0	8.3	8.5	423.4	18.1	26.8	28.1	427.4	8.4	100.9
TEM	77	NA()	2.2	17.9099	2.7	0.5110	4.7	0.0664	3.8	0.81	414.5	3.9	8.5	8.7	419.1	5.1	8.8	9.0	445.8	25.0	31.8	32.9	414.5	8.5	93.0
TEM	79	NA()	2.3	17.9659	2.8	0.5035	4.7	0.0656	3.8	0.81	409.8	3.8	8.4	8.6	414.1	5.1	8.7	9.0	438.8	25.9	32.5	33.6	409.8	8.4	93.4
TEM	92	NA()	2.4	17.9445	2.5	0.5064	4.5	0.0659	3.8	0.83	411.6	3.7	8.3	8.5	416.0	4.6	8.5	8.7	441.5	22.2	29.7	30.9	411.6	8.3	93.2
TEM	139	NA()	1.9	18.3181	2.1	0.4877	4.2	0.0648	3.6	0.86	404.9	3.2	8.0	8.2	403.3	3.9	7.9	8.2	395.5	18.7	27.3	28.6	404.9	8.0	102.4
TEM	167	NA()	1.5	18.0550	1.9	0.5299	3.9	0.0694	3.4	0.88	432.7	3.1	8.4	8.6	431.8	3.5	8.1	8.4	427.8	15.2	24.9	26.4	432.7	8.4	101.1
TEM	124	NA()	1.8	17.8627	2.2	0.5048	4.3	0.0654	3.6	0.85	408.5	3.3	8.1	8.3	414.9	4.2	8.2	8.4	451.7	19.6	27.8	29.1	408.5	8.1	90.4
TEM	123	NA()	1.7	17.5202	2.2	0.5311	4.2	0.0675	3.6	0.86	421.2	3.4	8.4	8.6	432.5	4.1	8.4	8.6	494.5	18.0	26.6	27.9	421.2	8.4	85.2
TEM	105	NA()	1.7	17.5025	2.3	0.5316	4.4	0.0675	3.7	0.85	421.1	3.6	8.4	8.6	432.9	4.4	8.5	8.8	496.7	19.6	27.7	28.9	421.1	8.4	84.8
TEM	76	NA()	2.5	17.9299	2.8	0.5178	4.7	0.0674	3.8	0.81	420.2	3.9	8.6	8.7	423.7	5.2	8.9	9.1	443.3	26.2	32.8	33.8	420.2	8.6	94.8
TEM	106	NA()	1.9	18.3662	2.4	0.4861	4.4	0.0648	3.7	0.83	404.6	3.4	8.1	8.3	402.3	4.4	8.2	8.4	389.6	22.2	29.8	31.0	404.6	8.1	103.9
TEM	384	NA()	1.5	18.2182	1.3	0.5077	3.0	0.0671	2.7	0.90	418.7	1.1	7.7	7.9	416.9	1.8	7.3	7.6	407.7	10.0	22.2	23.8	418.7	7.7	102.7
PEIXE	67	NA()	1.6	17.3811	2.5	0.7413	4.5	0.0935	3.8	0.83	576.1	5.3	11.6	11.9	563.2	6.1	10.8	11.1	512.0	22.4	29.6	30.8	576.1	11.6	112.5
PEIXE	62	NA()	1.6	16.7311	2.5	0.7630	4.6	0.0926	3.8	0.83	571.0	5.4	11.6	11.8	575.7	6.2	11.0	11.3	595.2	22.2	29.3	30.5	571.0	11.6	95.9
PEIXE	60	NA()	1.5	16.7299	2.5	0.7624	4.6	0.0925	3.8	0.83	570.6	5.4	11.6	11.8	575.4	6.2	11.0	11.3	595.4	21.9	29.1	30.3	570.6	11.6	95.8
PEIXE	61	NA()	1.6	16.4642	2.6	0.7362	4.6	0.0880	3.8	0.83	543.4	5.2	11.1	11.3	560.2	6.3	10.9	11.2	629.9	23.3	30.1	31.2	543.4	11.1	86.3
PEIXE	87	NA()	1.3	17.2549	2.2	0.7414	4.2	0.0928	3.6	0.86	572.2	5.0	11.4	11.7	563.2	5.4	10.5	10.8	528.1	19.0	27.2	28.4	572.2	11.4	108.4
PEIXE	77	NA()	1.5	17.2977	2.3	0.7329	4.4	0.0920	3.7	0.85	567.3	5.1	11.4	11.6	558.2	5.5	10.5	10.8	522.6	19.2	27.3	28.6	567.3	11.4	108.5
PEIXE	83	NA()	1.3	16.8884	2.2	0.7639	4.3	0.0936	3.7	0.86	576.9	5.1	11.5	11.8	576.3	5.4	10.6	10.9	574.9	17.5	26.0	27.3	576.9	11.5	100.3
PEIXE	105	NA()	1.3	17.0362	2.0	0.7527	4.1	0.0930	3.6	0.87	573.5	4.8	11.4	11.6	569.8	5.0	10.3	10.7	555.9	16.2	25.2	26.5	573.5	11.4	103.2
PEIXE	99	NA()	5.3	16.8080	2.0	0.7354	4.1	0.0897	3.6	0.87	553.7	4.8	11.0	11.3	559.7	5.1	10.3	10.6	585.3	16.9	25.6	26.9	553.7	11.0	94.6
PEIXE	112	NA()	6.6	16.8499	1.9	0.7629	4.0	0.0933	3.5	0.88	574.9	4.8	11.4	11.6	575.7	5.0	10.4	10.7	579.8	16.0	25.0	26.4	574.9	11.4	99.1
PEIXE	80	NA()	10.6	17.0184	2.4	0.7155	4.4	0.0884	3.7	0.85	545.8	5.2	11.1	11.3	548.0	5.7	10.5	10.8	558.2	20.0	27.8	29.0	545.8	11.1	97.8
PEIXE	199	NA()	3.6	17.0336	1.5	0.7200	3.4	0.0890	3.1	0.90	549.5	3.8	10.6	10.8	550.7	3.8	9.6	9.9	556.3	11.5	22.5	24.0	549.5	10.6	98.8
PEIXE	189	NA()	3.8	16.9542	1.5	0.7429	3.5	0.0914	3.1	0.90	563.8	4.0	10.9	11.1	564.1	3.9	9.8	10.1	566.5	11.4	22.4	23.9	563.8	10.9	99.5
PEIXE	171	NA()	3.6	16.8233	1.5	0.7685	3.5	0.0938	3.2	0.90	578.0	4.3	11.2	11.5	578.9	4.2	10.1	10.4	583.3	11.8	22.5	24.0	578.0	11.2	99.1
PEIXE	186	NA()	3.6	16.9041	1.5	0.7447	3.5	0.0913	3.2	0.90	563.5	4.0	10.9	11.2	565.2	4.0	9.8	10.2	572.9	11.4	22.4	23.9	563.5	10.9	98.4
PEIXE	31	NA()	3.2	16.4276	3.4	0.7420	4.7	0.0884	3.2	0.69	546.3	6.1	11.5	11.8	563.6	8.4	12.3	12.5	634.8	33.3	38.4	39.3	546.3	11.5	86.1