Supplementary Materials

Equation	Model	Descriptors	R^2	Q^2	R_{pred}^2	$\overline{r_{m(Overall)}^2}$	$\Delta r_{m(Overall)}^2$	$^{c}R_{p}^{2}$
1	2D- QSAR	S_aasN Atype_N_69 D/Dtr05 Atype_C_43 S_sCl	0.73	0.70	0.675	0.59	0.197	0.712
2	G-QSAR	R3-SsClE-index R2-chi5chain R2-SsNH2count R1-k1alpha R1-chi3Cluster	0.72	0.69	0.695	0.57	0.19	0.70
3	QAAR	Jurs-FPSA-2 R2-chi3Cluster R3- Electronegativity Count	0.86	0.79	-	-	-	0.78

Table S1. Summary of all 3 QSAR models along with their descriptors and important validation parameters.

Table S2. 2D-QSAR results consisting of compound names, values of important descriptors, observed, calculated and LOO predicted activity values and Glide docking scores.

Training set compounds												
Name ^a	S_aasN	Atype_N_69	D/Dtr05	Atype_C_43	S_sCl	pIC ₅₀ ^b	pIC ₅₀ ^c	pIC _d 50	gScore ^e			
Bla	1.277	1	105.723	1	5.380	6.409	6.678	6.712	-6.128			
B1b	1.292	1	97.819	1	0.000	7.319	6.531	6.433	-7.175			
B1c	1.296	1	130.127	1	0.000	6.201	6.590	6.629	-6.751			
B2d	1.376	0	236.690	1	5.374	6.335	6.439	6.441	-6.084			
B2f	1.413	0	147.737	1	0.000	6.241	6.161	6.159	-5.325			
B2g	1.417	0	180.045	1	0.000	6.049	6.220	6.223	-5.679			
B2h	1.416	0	194.538	1	5.491	6.171	6.388	6.391	-5.021			
B2j	1.392	0	118.089	1	0.000	6.223	6.099	6.092	-5.533			
B21	1.391	0	134.548	1	0.000	6.361	6.127	6.117	-5.909			
B4a	1.426	0	186.348	1	11.351	6.250	6.532	6.541	-6.765			
B4d	1.440	0	194.902	1	5.875	6.371	6.411	6.411	-6.739			
B4e	1.539	0	163.609	1	5.723	6.182	6.402	6.410	-6.123			
B4f	1.467	0	170.625	1	0.000	6.403	6.229	6.226	-6.472			

B4h	1.466	0	187.084	1	0.000	6.312	6.257	6.256	-7.099
B4i	1.431	0	148.882	1	5.430	6.382	6.314	6.313	-6.226
B4j	1.446	0	140.978	1	0.000	6.334	6.166	6.161	-6.364
B41	1.446	0	157.437	1	0.000	6.193	6.195	6.195	-6.070
B5b	1.405	0	194.452	1	5.882	6.420	6.392	6.392	-7.497
B5c	1.409	0	226.760	1	5.917	6.538	6.452	6.451	-6.548
B5d	1.404	0	210.910	1	5.889	6.252	6.421	6.423	-7.674
B5e	1.412	0	164.195	1	0.000	6.171	6.190	6.190	-6.444
B5f	1.431	0	186.634	1	0.000	6.231	6.238	6.239	-6.864
B5g	1.435	0	218.942	1	0.000	6.439	6.297	6.294	-6.381
B5i	1.482	0	150.656	1	5.677	6.158	6.350	6.355	-6.091
B5j	1.410	0	156.987	1	0.000	6.331	6.176	6.172	-6.816
B5k	1.414	0	189.295	1	0.000	6.223	6.235	6.235	-5.611
B51	1.410	0	173.445	1	0.000	6.127	6.205	6.207	-6.508
B6a	1.377	1	202.356	1	11.368	7.357	7.053	7.031	-9.224
B6b	1.392	1	194.452	1	5.877	7.143	6.904	6.892	-9.233
B6c	1.396	1	226.760	1	5.912	7.469	6.963	6.931	-9.267
B6d	1.391	1	210.910	1	5.885	7.194	6.932	6.918	-8.469
B6e	1.403	1	194.538	1	5.488	6.609	6.899	6.915	-8.208
B6f	1.418	1	186.634	1	0.000	6.706	6.750	6.752	-6.877
B6g	1.422	1	218.942	1	0.000	6.572	6.808	6.824	-8.189
B6i	1.382	1	164.891	1	5.437	6.697	6.836	6.843	-6.716
B6j	1.397	1	156.987	1	0.000	6.618	6.688	6.691	-6.825
B6k	1.401	1	189.295	1	0.000	6.730	6.746	6.747	-6.291
B61	1.397	1	173.445	1	0.000	6.025	6.716	6.753	-6.521
B7a	1.351	0	227.433	1	11.396	6.333	6.567	6.577	-8.918
B7b	1.525	0	184.593	1	0.000	6.193	6.282	6.284	-6.572
B7e	1.467	0	191.888	1	0.000	6.194	6.266	6.267	-7.704
B7f	1.370	0	256.203	1	5.928	6.272	6.484	6.491	-6.581
B7g	1.427	0	226.676	1	5.910	6.315	6.461	6.463	-5.731
B7h	1.369	0	272.662	1	5.935	6.572	6.513	6.510	-7.979
B7i	1.336	0	235.954	1	16.771	6.650	6.715	6.721	-7.709
B7k	1.355	0	260.358	1	11.327	6.469	6.625	6.632	-8.248
B71	1.493	1	168.716	1	0.000	6.745	6.756	6.757	-6.996
B7m	1.377	0	219.615	1	5.500	6.449	6.412	6.412	-8.466
B7n	1.392	0	211.711	1	0.000	6.266	6.263	6.263	-5.997
B7p	1.391	0	228.169	1	0.000	6.326	6.291	6.290	-7.246
B7r	1.396	0	248.385	1	0.000	6.376	6.329	6.327	-6.717
B7t	1.396	0	264.844	1	0.000	6.365	6.357	6.357	-8.004
B7x	1.386	0	172.013	1	5.840	6.335	6.343	6.343	-9.060

B7y	1.356	0	189.968	1	5.449	6.451	6.349	6.346	-6.184
B7za	1.375	0	214.372	1	0.000	6.192	6.259	6.261	-7.396
B7zb	1.371	0	198.522	1	0.000	6.334	6.229	6.226	-6.101
B7zd	1.375	0	218.738	1	0.000	6.361	6.266	6.264	-5.695
B7ze	1.379	0	251.046	1	0.000	6.384	6.325	6.322	-8.010
B7zf	1.375	0	235.197	1	0.000	6.322	6.295	6.294	-6.842
B7zh	1.356	0	190.584	1	5.318	6.322	6.346	6.347	-6.477
B7zj	1.356	0	207.043	1	5.324	6.578	6.375	6.371	-6.799
B8a	1.401	0	255.703	1	11.472	7.377	6.644	6.616	-6.335
B8b	1.416	0	247.798	1	5.935	7.523	6.493	6.473	-7.230
B8d	1.416	0	264.257	1	5.943	6.623	6.522	6.519	-7.649
B8e	1.517	0	218.453	1	0.000	6.331	6.337	6.337	-6.613
B8f	1.442	0	239.981	1	0.000	6.314	6.337	6.338	-6.862
B8g	1.447	0	272.289	1	0.000	6.305	6.396	6.400	-5.784
B8h	1.442	0	256.439	1	0.000	6.279	6.366	6.369	-7.475
B8i	1.407	0	218.238	1	5.483	6.271	6.425	6.427	-5.118
B8j	1.422	0	210.333	1	0.000	6.334	6.275	6.274	-6.889
B8k	1.427	0	247.885	1	5.534	6.331	6.488	6.491	-7.283
B81	1.421	0	226.792	1	0.000	6.197	6.304	6.306	-7.360
B9a	2.715	1	329.160	2	11.454	5.487	5.487	5.488	-5.400
B9b	2.737	1	317.248	2	5.926	5.126	5.333	5.349	-7.383
B9e	2.751	1	317.693	2	5.525	5.156	5.331	5.343	-6.185
B9f	2.773	1	305.782	2	0.000	5.185	5.177	5.176	-6.978
B9h	2.768	1	330.417	2	0.000	5.320	5.217	5.209	-7.175
B9i	2.726	1	273.203	2	5.474	5.334	5.239	5.231	-6.752
B9j	2.748	1	261.292	2	0.000	5.280	5.086	5.069	-6.928
B9k	2.754	1	311.567	2	0.000	5.198	5.177	5.176	-6.128
C10a	2.827	1	315.216	2	11.772	5.544	5.528	5.526	-7.088
C10b	2.853	1	306.943	2	5.994	5.195	5.376	5.388	-5.271
C10c	2.861	1	343.889	2	6.033	5.484	5.445	5.442	-6.264
C10e	2.864	1	304.265	2	5.774	5.209	5.370	5.382	-6.781
C10f	2.890	1	295.992	2	0.000	5.191	5.218	5.220	-4.994
C10g	2.897	1	332.937	2	0.000	5.338	5.286	5.282	-6.072
C10h	2.889	1	313.789	2	0.000	5.302	5.249	5.244	-5.365
C10j	2.865	1	253.563	2	0.000	5.289	5.132	5.119	-5.679
C10k	2.872	1	290.508	2	0.000	5.212	5.200	5.199	-6.736
C2b	1.382	1	91.335	1	0.000	6.413	6.566	6.579	-7.483
C3a	1.458	0	150.204	1	11.597	6.465	6.492	6.493	-7.397
C3b	1.477	0	144.832	1	5.900	6.337	6.343	6.343	-4.906
C3d	1.480	0	156.555	1	5.909	6.340	6.365	6.365	-4.094

C3f C3g C3h C3i C3j C3k C5c C5e C5h C5i C5j C5k C6a C6b C6c C6c C6c C6c C6c C6c C6c C6c	1.503 1.508 1.506 1.464 1.482	0 0 0	137.529 160.961	1 1	0.000 0.000	6.243 6.050	6.189 6.233	6.187 6.227	-7.232
C3g C3h C3i C3j C3k C5c C5e C5h C5i C5j C5k C6a C6b C6c C6d C6e C6g C6h C6i	1.508 1.506 1.464 1.482	0 0	160.961	1	0.000	6.050	6 223	6 227	7 650
C3h C3i C3j C3k C5c C5e C5h C5i C5j C5k C6a C6b C6c C6c C6c C6c C6c C6c C6c C6c	1.506 1.464 1.482	0	1 40 0 50			0.050	0.255	0.237	-7.039
C3i C3j C3k C5c C5e C5h C5i C5j C5k C6a C6b C6c C6d C6c C6d C6e C6g C6h C6i	1.464 1.482		149.252	1	0.000	6.171	6.211	6.212	-5.488
C3j C3k C5c C5e C5h C5i C5j C5k C6a C6b C6c C6d C6e C6g C6h C6i	1 482	0	115.315	1	5.636	6.162	6.278	6.283	-8.071
C3k C5c C5e C5h C5i C5j C5k C6a C6b C6c C6d C6c C6d C6e C6g C6h C6i	1.102	0	109.943	1	0.000	6.234	6.130	6.125	-6.245
C5c C5e C5h C5i C5j C5k C6a C6b C6c C6d C6c C6g C6h C6i	1.487	0	133.375	1	0.000	6.209	6.174	6.172	-7.195
C5e C5h C5i C5j C5k C6a C6b C6c C6d C6e C6g C6h C6i	1.536	0	188.972	1	5.974	6.259	6.452	6.457	-5.734
C5h C5i C5j C5k C6a C6b C6c C6d C6c C6d C6e C6g C6h C6i	1.452	0	178.530	1	5.480	6.182	6.378	6.380	-6.667
C5i C5j C5k C6a C6b C6c C6d C6e C6g C6h C6i	1.561	0	169.960	1	0.000	5.312	6.275	6.313	-6.436
C5j C5k C6a C6b C6c C6d C6e C6g C6h C6i	1.469	1	150.656	1	5.674	7.215	6.861	6.839	-6.456
C5k C6a C6b C6c C6d C6e C6g C6h C6i	1.536	0	130.651	1	0.000	6.339	6.194	6.187	-6.868
C6a C6b C6c C6d C6e C6g C6h C6i	1.542	0	154.083	1	0.000	6.249	6.237	6.237	-6.626
C6b C6c C6d C6e C6g C6h C6i	1.477	0	185.545	1	11.689	6.354	6.565	6.573	-6.787
C6c C6d C6e C6g C6h C6i	1.500	0	145.284	1	0.000	6.410	6.201	6.195	-6.378
C6d C6e C6g C6h	1.500	0	203.605	1	5.989	6.533	6.459	6.458	-6.606
C6e C6g C6h C6i	1.498	0	191.896	1	5.958	6.267	6.437	6.440	-7.027
C6g C6h C6i	1.503	0	178.242	1	5.736	6.167	6.410	6.415	-6.359
C6h C6i	1.526	0	196.302	1	0.000	6.452	6.304	6.299	-6.473
C6i	1.366	0	219.529	1	5.893	6.193	6.417	6.422	-6.468
001	1.395	0	164.891	1	5.441	6.158	6.324	6.328	-5.507
C6j	1.495	0	180.173	1	5.949	6.410	6.415	6.415	-6.187
C6k	1.506	0	168.716	1	0.000	6.246	6.245	6.245	-6.386
C7a	1.464	1	185.545	1	11.681	7.409	7.076	7.046	-5.440
C7b	1.482	1	180.173	1	5.945	7.194	6.926	6.909	-4.697
C7f	1.508	1	172.870	1	0.000	6.717	6.771	6.775	-4.767
C7g	1.513	1	196.302	1	0.000	6.578	6.815	6.833	-6.440
C7h	1.512	1	184.593	1	0.000	6.342	6.794	6.827	-7.027
C7i	1.518	0	136.023	1	5.665	7.215	6.342	6.304	-6.715
C7k	1.350	0	244.508	1	11.278	6.745	6.593	6.587	-7.446
C8aa	1.355	0	264.107	1	11.459	6.194	6.635	6.657	-6.820
C8ab	1.465	0	180.179	1	0.000	6.437	6.244	6.241	-5.626
C8ac	1.447	0	208.171	1	5.718	6.440	6.434	6.433	-6.162
C8af	1.469	0	214.522	1	0.000	6.322	6.306	6.306	-6.529
C8ag	1.428	0	181.745	1	11.059	6.345	6.517	6.523	-5.921
C8ah	1.446	0	176.373	1	5.369	6.324	6.369	6.369	-5.862
C8ai	1.452	0	199.805	1	5.400	6.446	6.413	6.413	-5.285
C8aj	1.450	0	188.096	1	5.376	6.577	6.391	6.389	-5.515
C8b		-	000 046						
C8d	1.456	0	203.346	1	5.961	6.193	6.436	6.438	-6.092

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B2c

B2e

B2i

B2k

B4b

B4c

1.390

1.398

1.377

1.396

1.441

1.445

0

0

0

0

0

0

187.863

155.641

125.994

150.398

178.443

210.751

1

1

1

1

1

1

C8e	1.442	0	243.060	1	11.776	6.210	6.650	6.666	-8.004	
C8f	1.460	0	237.688	1	5.995	6.280	6.499	6.502	-6.702	
C8i	1.423	0	216.634	1	17.136	6.640	6.734	6.742	-5.740	
C8j	1.441	0	211.262	1	11.383	6.585	6.584	6.584	-7.087	
C8k	1.446	0	234.694	1	11.454	6.469	6.629	6.634	-5.259	
C81	1.445	0	222.985	1	11.399	6.733	6.607	6.603	-7.702	
C8m	1.464	0	201.415	1	5.745	6.456	6.431	6.431	-8.363	
C80	1.487	0	219.475	1	0.000	6.249	6.324	6.326	-7.610	
C8q	1.468	0	235.757	1	5.776	6.330	6.494	6.496	-7.264	
C8r	1.486	0	230.385	1	0.000	6.384	6.343	6.342	-5.824	
C8s	1.492	0	253.817	1	0.000	6.332	6.386	6.389	-5.118	
C8t	1.490	0	242.108	1	0.000	6.363	6.365	6.365	-6.107	
C8u	1.449	0	209.331	1	11.168	6.274	6.579	6.588	-7.638	
C8v	1.467	0	203.959	1	5.419	6.192	6.429	6.431	-6.114	
C8w	1.473	0	227.391	1	5.451	6.262	6.473	6.476	-6.808	
C8y	1.443	0	173.829	1	5.687	6.343	6.371	6.371	-5.698	
C8z	1.461	0	168.457	1	0.000	6.485	6.222	6.217	-7.186	
C9a	1.488	0	235.283	1	11.789	7.456	6.660	6.626	-4.567	
C9b	1.507	0	229.911	1	6.003	6.672	6.509	6.505	-8.047	
C9c	1.512	0	253.342	1	6.042	6.664	6.554	6.550	-6.565	
C9d	1.510	0	241.633	1	6.012	6.588	6.531	6.530	-6.829	
C9e	1.514	0	227.980	1	5.783	7.495	6.504	6.479	-8.491	
C9f	1.533	0	222.608	1	0.000	6.415	6.353	6.350	-7.209	
C9g	1.538	0	246.039	1	0.000	6.362	6.396	6.398	-7.857	
C9h	1.536	0	234.330	1	0.000	6.573	6.375	6.367	-3.818	
C9i	1.494	0	200.393	1	5.725	6.474	6.444	6.443	-6.385	
C9k	1.426	0	242.642	1	0.000	6.331	6.334	6.334	-7.072	
				Test set compo	ounds					
Name ^a	S_aasN	Atype_N_69	D/Dtr05	Atype_C_43	S_sCl	pIC ₅₀ ^b	pIC ₅₀ ^c	pIC ₅₀	gScore ^e	
B1d	1.292	1	114.277	1	0.000	6.578	6.560	-	-7.467	
B2a	1.377	0	220.231	1	5.368	6.192	6.410	-	-4.779	
B2b	1.386	0	155.555	1	5.832	6.335	6.314	-	-5.145	

6.730

6.119

6.165

6.212

7.208

7.194

6.373

6.310

6.246

6.157

6.382

6.442

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-

5.867

5.455

5.405

0.000

5.867

5.902

-5.807

-5.059

-5.669

-6.405

-7.767

-6.638

B4g	1.471	0	202.934	1	0.000	6.305	6.287	-	-6.724
B4k	1.450	0	173.286	1	0.000	6.241	6.225	-	-6.679
B5a	1.390	0	202.356	1	11.376	6.357	6.542	-	-6.864
B5h	1.430	0	203.092	1	0.000	6.188	6.267	-	-7.451
B6h	1.417	1	203.092	1	0.000	6.333	6.778	-	-7.422
B7c	1.370	0	251.837	1	5.928	6.469	6.476	-	-10.732
B7d	1.365	0	235.987	1	5.901	6.453	6.446	-	-6.259
B7j	1.351	0	228.049	1	11.264	6.585	6.565	-	-7.249
B7o	1.396	0	244.019	1	0.000	6.249	6.321	-	-9.204
B7q	1.381	0	256.290	1	5.528	6.331	6.479	-	-10.095
B7s	1.400	0	280.694	1	0.000	6.334	6.387	-	-5.986
B7u	1.362	0	228.136	1	10.871	6.276	6.560	-	-6.999
B7v	1.371	0	163.459	1	11.291	6.192	6.463	-	-7.110
B7w	1.381	0	252.540	1	5.396	6.266	6.469	-	-9.535
B7z	1.371	0	182.064	1	0.000	6.416	6.200	-	-5.960
B7zc	1.360	0	226.642	1	5.477	6.438	6.416	-	-6.613
B7zg	1.341	0	198.489	1	10.770	6.335	6.495	-	-6.300
B7zi	1.360	0	222.893	1	5.346	6.449	6.406	-	-6.311
B8c	1.420	0	280.107	1	5.970	6.609	6.552	-	-6.955
B9c	2.743	1	367.524	2	5.961	5.498	5.425	-	-6.286
B9d	2.732	1	341.884	2	5.934	5.177	5.374	-	-7.521
B9g	2.779	1	356.057	2	0.000	5.332	5.267	-	-6.359
B91	2.743	1	285.927	2	0.000	5.193	5.127	-	-6.677
C10d	2.852	1	324.741	2	6.003	5.233	5.406	-	-7.069
C10i	2.839	1	261.835	2	5.716	5.394	5.282	-	-7.128
C101	2.864	1	271.360	2	0.000	5.207	5.162	-	-7.050
C2a	1.364	1	96.707	1	5.606	7.367	6.712	-	-7.438
C2c	1.388	1	114.767	1	0.000	6.207	6.609	-	-8.810
C2d	1.386	1	103.058	1	0.000	6.572	6.588	-	-7.495
C3c	1.482	0	168.264	1	5.939	6.738	6.387	-	-5.626
C31	1.485	0	121.666	1	0.000	6.364	6.152	-	-8.189
C5a	1.513	0	170.912	1	11.662	7.237	6.557	-	-5.009
C5b	1.531	0	165.540	1	5.935	6.382	6.407	-	-4.452
C5d	1.535	0	177.263	1	5.944	6.375	6.430	-	-7.242
C5f	1.557	0	158.237	1	0.000	6.406	6.253	-	-5.180
C5g	1.563	0	181.669	1	0.000	6.313	6.296	-	-4.091
C51	1.540	0	142.374	1	0.000	6.193	6.216	-	-5.650
C6f	1.521	0	172.870	1	0.000	6.238	6.260	-	-5.831
C6l	1.504	0	157.007	1	0.000	6.135	6.223	-	-5.958
C7c	1.487	1	203.605	1	5.984	6.627	6.971	-	-5.008

C7d	1.485	1	191.896	1	5.954	6.697	6.949	-	-7.308
C7e	1.490	1	178.242	1	5.732	7.432	6.921	-	-5.504
C7j	1.487	1	145.284	1	0.000	6.604	6.713	-	-6.523
C71	1.491	1	157.007	1	0.000	6.031	6.735	-	-6.787
C8a	1.438	0	208.718	1	11.710	6.330	6.586	-	-6.040
C8ad	1.466	0	202.799	1	0.000	6.361	6.284	-	-4.987
C8ae	1.471	0	226.231	1	0.000	6.386	6.328	-	-8.514
C8c	1.461	0	226.777	1	6.000	6.498	6.481	-	-6.864
C8g	1.465	0	261.120	1	6.035	6.316	6.544	-	-5.993
C8h	1.464	0	249.411	1	6.004	6.567	6.521	-	-7.308
C8n	1.482	0	196.043	1	0.000	6.261	6.281	-	-6.977
C8p	1.486	0	207.766	1	0.000	6.331	6.303	-	-6.815
C8x	1.471	0	215.682	1	5.426	6.441	6.451	-	-7.620
C9j	1.512	0	195.021	1	0.000	6.344	6.294	-	-4.774
C91	1.515	0	206.744	1	0.000	6.272	6.316	-	-4.817

^aCompound names ^bObserved activity

^cCalculated activity using 2D-QSAR model ^dLOO predicted activity ^eGlide docking score (gScore)

Table S3. G-QSAR results consisting of compound names, values of important descriptors, and observed, calculated and LOO predicted activity values.

Training set compounds													
Name ^a	R3- SsClE- index	R2- chi5chain	R2- SsNH2count	R1-k1alpha	R1- chi3Cluster	pIC ₅₀ ^b	pIC ₅₀ ^c	pIC ₅₀ ^d					
B1a	0.780	0.000	0	0.000	0.000	6.409	6.728	6.790					
B1b	0.000	0.000	0	0.000	0.000	7.319	6.622	6.499					
B1c	0.000	0.000	0	0.000	0.000	6.201	6.622	6.696					
B2d	0.000	0.000	0	6.302	0.493	6.335	6.505	6.509					
B2f	0.000	0.000	0	5.035	0.204	6.241	6.267	6.267					
B2g	0.000	0.000	0	5.035	0.204	6.049	6.267	6.271					
B2h	0.000	0.000	0	5.035	0.204	6.171	6.267	6.269					
B2j	0.000	0.000	0	2.670	0.000	6.223	6.285	6.286					
B21	0.000	0.000	0	2.670	0.000	6.361	6.285	6.283					
B4a	0.780	0.000	0	6.302	0.493	6.250	6.611	6.624					
B4d	0.000	0.000	0	6.302	0.493	6.371	6.505	6.508					
B4e	0.780	0.000	0	5.035	0.204	6.182	6.373	6.379					

B4f	0.000	0.000	0	5.035	0.204	6.403	6.267	6.265
B4h	0.000	0.000	0	5.035	0.204	6.312	6.267	6.266
B4i	0.780	0.000	0	2.670	0.000	6.382	6.391	6.391
B4j	0.000	0.000	0	2.670	0.000	6.334	6.285	6.284
B41	0.000	0.000	0	2.670	0.000	6.193	6.285	6.286
B5b	0.000	0.000	0	6.302	0.493	6.420	6.505	6.507
B5c	0.000	0.000	0	6.302	0.493	6.538	6.505	6.504
B5d	0.000	0.000	0	6.302	0.493	6.252	6.505	6.511
B5e	0.780	0.000	0	5.035	0.204	6.171	6.373	6.380
B5f	0.000	0.000	0	5.035	0.204	6.231	6.267	6.268
B5g	0.000	0.000	0	5.035	0.204	6.439	6.267	6.264
B5i	0.780	0.000	0	2.670	0.000	6.158	6.391	6.399
B5j	0.000	0.000	0	2.670	0.000	6.331	6.285	6.284
B5k	0.000	0.000	0	2.670	0.000	6.223	6.285	6.286
B51	0.000	0.000	0	2.670	0.000	6.127	6.285	6.288
B6a	0.780	0.000	1	6.302	0.493	7.357	7.080	7.055
B6b	0.000	0.000	1	6.302	0.493	7.143	6.974	6.961
B6c	0.000	0.000	1	6.302	0.493	7.469	6.974	6.936
B6d	0.000	0.000	1	6.302	0.493	7.194	6.974	6.957
B6e	0.780	0.000	1	5.035	0.204	6.609	6.842	6.861
B6f	0.000	0.000	1	5.035	0.204	6.706	6.736	6.738
B6g	0.000	0.000	1	5.035	0.204	6.572	6.736	6.747
B6i	0.780	0.000	1	2.670	0.000	6.697	6.860	6.874
B6j	0.000	0.000	1	2.670	0.000	6.618	6.754	6.764
B6k	0.000	0.000	1	2.670	0.000	6.730	6.754	6.756
B61	0.000	0.000	1	2.670	0.000	6.025	6.754	6.806
B7a	0.780	0.000	0	6.302	0.493	6.333	6.611	6.621
B7b	0.000	0.000	0	6.302	0.493	6.193	6.505	6.512
B7e	0.780	0.000	0	6.302	0.493	6.194	6.611	6.627
B7f	0.000	0.000	0	6.302	0.493	6.272	6.505	6.510
B7g	0.000	0.000	0	6.302	0.493	6.315	6.505	6.509
B7h	0.000	0.000	0	6.302	0.493	6.572	6.505	6.503
B7i	0.780	0.000	0	6.302	0.493	6.650	6.611	6.609
B7k	0.000	0.000	0	6.302	0.493	6.469	6.505	6.506
B71	0.000	0.000	0	6.302	0.493	6.745	6.505	6.499
B7m	0.780	0.000	0	5.035	0.204	6.449	6.373	6.370
B7n	0.000	0.000	0	5.035	0.204	6.266	6.267	6.267
B7p	0.000	0.000	0	5.035	0.204	6.326	6.267	6.266
B7r	0.000	0.000	0	5.035	0.204	6.376	6.267	6.265
B7t	0.000	0.000	0	5.035	0.204	6.365	6.267	6.265

$\mathbf{D7}_{\mathbf{v}}$	0.000	0.000	Ο	5 025	0 204	6 2 2 5	6 267	()((
B/X D7u	0.000	0.000	0	5.055	0.204	0.333	0.207	6.266
В/у D7-с	0.780	0.000	0	2.670	0.000	0.431	0.391	6.389
B/Za D7-h	0.000	0.000	0	2.670	0.000	0.192	0.285	6.286
B/20 D7-4	0.000	0.000	0	2.670	0.000	0.334	0.285	6.284
B/Za	0.000	0.000	0	2.670	0.000	0.301	6.285	6.283
B/ze	0.000	0.000	0	2.670	0.000	6.384	6.285	6.283
B/ZI	0.000	0.000	0	2.670	0.000	6.322	6.285	6.284
B/zh	0.000	0.000	0	2.670	0.000	6.322	6.285	6.284
B/zj	0.000	0.000	0	2.670	0.000	6.578	6.285	6.279
B8a	0.780	0.000	0	6.302	0.493	7.377	6.611	6.582
B8b	0.000	0.000	0	6.302	0.493	7.523	6.505	6.480
B8d	0.000	0.000	0	6.302	0.493	6.623	6.505	6.502
B8e	0.780	0.000	0	5.035	0.204	6.331	6.373	6.374
B8f	0.000	0.000	0	5.035	0.204	6.314	6.267	6.266
B8g	0.000	0.000	0	5.035	0.204	6.305	6.267	6.266
B8h	0.000	0.000	0	5.035	0.204	6.279	6.267	6.267
B8i	0.780	0.000	0	2.670	0.000	6.271	6.391	6.395
B8j	0.000	0.000	0	2.670	0.000	6.334	6.285	6.284
B8k	0.000	0.000	0	2.670	0.000	6.331	6.285	6.284
B81	0.000	0.000	0	2.670	0.000	6.197	6.285	6.286
B9a	0.780	0.096	1	6.302	0.493	5.487	5.522	5.526
B9b	0.000	0.096	1	6.302	0.493	5.126	5.416	5.440
B9e	0.780	0.096	1	5.035	0.204	5.156	5.284	5.295
B9f	0.000	0.096	1	5.035	0.204	5.185	5.178	5.178
B9h	0.000	0.096	1	5.035	0.204	5.320	5.178	5.168
B9i	0.780	0.096	1	2.670	0.000	5.334	5.302	5.299
B9j	0.000	0.096	1	2.670	0.000	5.280	5.196	5.190
B9k	0.000	0.096	1	2.670	0.000	5.198	5.196	5.196
C10a	0.813	0.096	1	6.302	0.493	5.544	5.527	5.525
C10b	0.000	0.096	1	6.302	0.493	5.195	5.416	5.434
C10c	0.000	0.096	1	6.302	0.493	5.484	5.416	5.411
C10e	0.813	0.096	1	5.035	0.204	5.209	5.289	5.296
C10f	0.000	0.096	1	5.035	0.204	5.191	5.178	5.178
C10g	0.000	0.096	1	5.035	0.204	5.338	5.178	5.167
C10h	0.000	0.096	1	5.035	0.204	5.302	5.178	5.170
C10j	0.000	0.096	1	2.670	0.000	5.289	5.196	5.189
C10k	0.000	0.096	1	2.670	0.000	5.212	5.196	5 195
C2b	0.000	0.000	0	0.000	0.000	6.413	6.622	6.659
C3a	0.813	0.000	0	6.302	0.493	6.465	6.615	6 621
C3b	0.000	0.000	0	6.302	0.493	6.337	6.505	6 509
			~	0.2 02		5.551		0.507

C3d	0.000	0.000	0	6.302	0.493	6.340	6.505	6.509
C3e	0.813	0.000	0	5.035	0.204	6.124	6.377	6.386
C3f	0.000	0.000	0	5.035	0.204	6.243	6.267	6.267
C3g	0.000	0.000	0	5.035	0.204	6.050	6.267	6.271
C3h	0.000	0.000	0	5.035	0.204	6.171	6.267	6.269
C3i	0.813	0.000	0	2.670	0.000	6.162	6.395	6.404
C3j	0.000	0.000	0	2.670	0.000	6.234	6.285	6.286
C3k	0.000	0.000	0	2.670	0.000	6.209	6.285	6.286
C5c	0.000	0.000	0	6.302	0.493	6.259	6.505	6.511
C5e	0.813	0.000	0	5.035	0.204	6.182	6.377	6.384
C5h	0.000	0.000	0	5.035	0.204	5.312	6.267	6.284
C5i	0.813	0.000	0	2.670	0.000	7.215	6.395	6.366
C5j	0.000	0.000	0	2.670	0.000	6.339	6.285	6.284
C5k	0.000	0.000	0	2.670	0.000	6.249	6.285	6.285
C6a	0.813	0.000	0	6.302	0.493	6.354	6.615	6.626
C6b	0.000	0.000	0	6.302	0.493	6.410	6.505	6.507
C6c	0.000	0.000	0	6.302	0.493	6.533	6.505	6.504
C6d	0.000	0.000	0	6.302	0.493	6.267	6.505	6.510
C6e	0.813	0.000	0	5.035	0.204	6.167	6.377	6.385
C6g	0.000	0.000	0	5.035	0.204	6.452	6.267	6.264
C6h	0.000	0.000	0	5.035	0.204	6.193	6.267	6.268
C6i	0.813	0.000	0	2.670	0.000	6.158	6.395	6.404
C6j	0.000	0.000	0	2.670	0.000	6.410	6.285	6.283
C6k	0.000	0.000	0	2.670	0.000	6.246	6.285	6.285
C7a	0.813	0.000	1	6.302	0.493	7.409	7.084	7.054
C7b	0.000	0.000	1	6.302	0.493	7.194	6.974	6.957
C7f	0.000	0.000	1	5.035	0.204	6.717	6.736	6.737
C7g	0.000	0.000	1	5.035	0.204	6.578	6.736	6.747
C7h	0.000	0.000	1	5.035	0.204	6.342	6.736	6.763
C7i	0.813	0.000	1	2.670	0.000	7.215	6.864	6.833
C7k	0.000	0.000	1	2.670	0.000	6.745	6.754	6.754
C8aa	0.000	0.000	0	2.670	0.000	6.194	6.285	6.286
C8ab	0.000	0.000	0	2.670	0.000	6.437	6.285	6.282
C8ac	0.813	0.000	0	2.670	0.000	6.440	6.395	6.394
C8af	0.000	0.000	0	2.670	0.000	6.322	6.285	6.284
C8ag	0.813	0.000	0	2.670	0.000	6.345	6.395	6.397
C8ah	0.000	0.000	0	2.670	0.000	6.324	6.285	6.284
C8ai	0.000	0.000	0	2.670	0.000	6.446	6.285	6.282
C8aj	0.000	0.000	0	2.670	0.000	6.577	6.285	6.280
C8b	0.000	0.000	0	6.302	0.493	6.193	6.505	6.512

C8d	0.000	0.000	0	6.302	0.493	6.453	6.505	6.506
C8e	0.813	0.000	0	6.302	0.493	6.210	6.615	6.631
C8f	0.000	0.000	0	6.302	0.493	6.280	6.505	6.510
C8i	0.813	0.000	0	6.302	0.493	6.640	6.615	6.614
C8j	0.000	0.000	0	6.302	0.493	6.585	6.505	6.503
C8k	0.000	0.000	0	6.302	0.493	6.469	6.505	6.506
C81	0.000	0.000	0	6.302	0.493	6.733	6.505	6.499
C8m	0.813	0.000	0	5.035	0.204	6.456	6.377	6.375
C80	0.000	0.000	0	5.035	0.204	6.249	6.267	6.267
C8q	0.813	0.000	0	5.035	0.204	6.330	6.377	6.379
C8r	0.000	0.000	0	5.035	0.204	6.384	6.267	6.265
C8s	0.000	0.000	0	5.035	0.204	6.332	6.267	6.266
C8t	0.000	0.000	0	5.035	0.204	6.363	6.267	6.265
C8u	0.813	0.000	0	5.035	0.204	6.274	6.377	6.381
C8v	0.000	0.000	0	5.035	0.204	6.192	6.267	6.268
C8w	0.000	0.000	0	5.035	0.204	6.262	6.267	6.267
C8y	0.813	0.000	0	2.670	0.000	6.343	6.395	6.397
C8z	0.000	0.000	0	2.670	0.000	6.485	6.285	6.281
C9a	0.813	0.000	0	6.302	0.493	7.456	6.615	6.582
C9b	0.000	0.000	0	6.302	0.493	6.672	6.505	6.501
C9c	0.000	0.000	0	6.302	0.493	6.664	6.505	6.501
C9d	0.000	0.000	0	6.302	0.493	6.588	6.505	6.503
C9e	0.813	0.000	0	5.035	0.204	7.495	6.377	6.338
C9f	0.000	0.000	0	5.035	0.204	6.415	6.267	6.264
C9g	0.000	0.000	0	5.035	0.204	6.362	6.267	6.265
C9h	0.000	0.000	0	5.035	0.204	6.573	6.267	6.261
C9i	0.813	0.000	0	2.670	0.000	6.474	6.395	6.392
C9k	0.000	0.000	0	2.670	0.000	6.331	6.285	6.284

Test set compounds

Name ^a	R3- SsClE- index	R2- chi5chain	R2- SsNH2count	R1-k1alpha	R1- chi3Cluster	pIC ₅₀ ^b	pIC ₅₀ ^c	pIC ₅₀ ^d
B1d	0.000	0.000	0	0.000	0.000	6.578	6.622	-
B2a	0.780	0.000	0	6.302	0.493	6.192	6.611	-
B2b	0.000	0.000	0	6.302	0.493	6.335	6.505	-
B2c	0.000	0.000	0	6.302	0.493	6.730	6.505	-
B2e	0.780	0.000	0	5.035	0.204	6.119	6.373	-
B2i	0.780	0.000	0	2.670	0.000	6.165	6.391	-
B2k	0.000	0.000	0	2.670	0.000	6.212	6.285	-
B4b	0.000	0.000	0	6.302	0.493	7.208	6.505	-

B4c	0.000	0.000	0	6.302	0.493	7.194	6.505	-
B4g	0.000	0.000	0	5.035	0.204	6.305	6.267	-
B4k	0.000	0.000	0	2.670	0.000	6.241	6.285	-
B5a	0.780	0.000	0	6.302	0.493	6.357	6.611	-
B5h	0.000	0.000	0	5.035	0.204	6.188	6.267	-
B6h	0.000	0.000	1	5.035	0.204	6.333	6.736	-
B7c	0.000	0.000	0	6.302	0.493	6.469	6.505	-
B7d	0.000	0.000	0	6.302	0.493	6.453	6.505	-
B7j	0.000	0.000	0	6.302	0.493	6.585	6.505	-
B7o	0.000	0.000	0	5.035	0.204	6.249	6.267	-
B7q	0.780	0.000	0	5.035	0.204	6.331	6.373	-
B7s	0.000	0.000	0	5.035	0.204	6.334	6.267	-
B7u	0.780	0.000	0	5.035	0.204	6.276	6.373	-
B7v	0.000	0.000	0	5.035	0.204	6.192	6.267	-
B7w	0.000	0.000	0	5.035	0.204	6.266	6.267	-
B7z	0.000	0.000	0	2.670	0.000	6.416	6.285	-
B7zc	0.780	0.000	0	2.670	0.000	6.438	6.391	-
B7zg	0.780	0.000	0	2.670	0.000	6.335	6.391	-
B7zi	0.000	0.000	0	2.670	0.000	6.449	6.285	-
B8c	0.000	0.000	0	6.302	0.493	6.609	6.505	-
B9c	0.000	0.096	1	6.302	0.493	5.498	5.416	-
B9d	0.000	0.096	1	6.302	0.493	5.177	5.416	-
B9g	0.000	0.096	1	5.035	0.204	5.332	5.178	-
B91	0.000	0.096	1	2.670	0.000	5.193	5.196	-
C10d	0.000	0.096	1	6.302	0.493	5.233	5.416	-
C10i	0.813	0.096	1	2.670	0.000	5.394	5.307	-
C101	0.000	0.096	1	2.670	0.000	5.207	5.196	-
C2a	0.813	0.000	0	0.000	0.000	7.367	6.732	-
C2c	0.000	0.000	0	0.000	0.000	6.207	6.622	-
C2d	0.000	0.000	0	0.000	0.000	6.572	6.622	-
C3c	0.000	0.000	0	6.302	0.493	6.738	6.505	-
C31	0.000	0.000	0	2.670	0.000	6.364	6.285	-
C5a	0.813	0.000	0	6.302	0.493	7.237	6.615	-
C5b	0.000	0.000	0	6.302	0.493	6.382	6.505	-
C5d	0.000	0.000	0	6.302	0.493	6.375	6.505	-
C5f	0.000	0.000	0	5.035	0.204	6.406	6.267	-
C5g	0.000	0.000	0	5.035	0.204	6.313	6.267	-
C51	0.000	0.000	0	2.670	0.000	6.193	6.285	-
C6f	0.000	0.000	0	5.035	0.204	6.238	6.267	-
C61	0.000	0.000	0	2.670	0.000	6.135	6.285	-

C7c	0.000	0.000	1	6.302	0.493	6.627	6.974	-	
C7d	0.000	0.000	1	6.302	0.493	6.697	6.974	-	
C7e	0.813	0.000	1	5.035	0.204	7.432	6.846	-	
C7j	0.000	0.000	1	2.670	0.000	6.604	6.754	-	
C71	0.000	0.000	1	2.670	0.000	6.031	6.754	-	
C8a	0.813	0.000	0	6.302	0.493	6.330	6.615	-	
C8ad	0.000	0.000	0	2.670	0.000	6.361	6.285	-	
C8ae	0.000	0.000	0	2.670	0.000	6.386	6.285	-	
C8c	0.000	0.000	0	6.302	0.493	6.498	6.505	-	
C8g	0.000	0.000	0	6.302	0.493	6.316	6.505	-	
C8h	0.000	0.000	0	6.302	0.493	6.567	6.505	-	
C8n	0.000	0.000	0	5.035	0.204	6.261	6.267	-	
C8p	0.000	0.000	0	5.035	0.204	6.331	6.267	-	
C8x	0.000	0.000	0	5.035	0.204	6.441	6.267	-	
C9j	0.000	0.000	0	2.670	0.000	6.344	6.285	-	
C91	0.000	0.000	0	2.670	0.000	6.272	6.285	-	

^aCompound names ^bObserved activity

^cCalculated activity using G-QSAR model ^dLOO predicted activity

Name ^a	Jurs- FPSA_2	R2- chi3Cluster	R3-	pIC ₅₀ ^b	pIC ₅₀ ^c	pIC ₅₀ ^d
	11 SA-2	chijClusici	ElectronegativityCount			
B1b	0.925	0.000	0.662	0.018	0.173	0.244
B4b	1.175	0.000	0.662	1.265	1.138	1.106
B4c	1.261	0.000	0.638	1.321	1.333	1.336
B6a	1.606	0.408	0.735	1.306	1.419	1.480
B6b	1.629	0.408	0.662	0.938	1.094	1.108
B6c	1.639	0.408	0.638	1.139	0.996	0.987
B6d	1.887	0.408	0.668	1.996	2.121	2.186
B8a	1.487	0.612	0.735	0.085	0.062	0.040
B8b	1.792	0.612	0.662	1.232	0.824	0.748
C2a	0.985	0.000	0.597	0.083	0.038	0.010
C5a	1.302	0.000	0.597	1.292	1.259	1.252
C5i	1.320	0.000	0.597	1.335	1.327	1.327
C7a	1.705	0.408	0.597	1.312	1.020	0.986
C7b	1.761	0.408	0.555	0.963	0.998	1.008

Table S4. QAAR results consisting of compound names, values of important descriptors, and observed, calculated and LOO predicted activity values.

C7e	1.832	0.408	0.597	1.120	1.509	1.568
C7i	1.894	0.408	0.597	1.992	1.747	1.694
C9a	1.836	0.612	0.597	0.137	0.628	0.782
C9e	1.956	0.612	0.597	1.243	1.090	1.054

^aCompound names ^bObserved activity ^(CDK5-CDK2) ^cCalculated activity ^(CDK5-CDK2) using QAAR model ^dLOO predicted activity^(CDK5-CDK2)