

Table ST1. Average number of additional change points after introducing a chromosomal aberration.

size	deletion	duplication	translocation
3	2.33	1.65	1.01
2	2.28	2.35	1.09
1	2.24	2.43	0.85
0.75	2.83	2.70	0.79
0.5	3.25	3.20	1.14
0.4	3.38	3.21	1.03
0.3	3.38	3.25	1.08
0.2	3.35	3.20	1.15
0.1	2.85	2.96	1.16
0.05	1.74	2.31	0.90
0.01	0.33	0.35	0.32