

Table 9. Chromosome counts after siRNA-mediated down-regulation

Condition	n*	Percentage of mitotic spreads								
		≤43	44	45	46	47	48	49	≥50	FI _{≥50} [†]
Untransfected	310	4.5	15.5	62.3	13.6	2.7	0.3	0.0	1.7	1.0
GAPD	380	9.2	18.2	55.0	11.6	3.9	0.8	0.0	1.3	0.8×
SMC1L1	304	7.2	16.1	46.7	13.2	6.6	0.7	0.7	8.9	5.5×
CSPG6	311	12.2	15.1	48.6	9.6	4.5	1.6	0.0	8.4	5.2×
STAG2	315	8.3	14.3	55.2	10.5	2.2	1.0	0.0	8.6	5.3×
MRE11A	307	5.2	16.0	54.7	12.1	4.9	2.0	0.7	4.6	2.8×

*Represents the total number of mitotic spreads quantified.

[†]FI_{≥50} represents the fold increase of the ≥50 population over that of the untransfected control.