

Table 6. Summary of cell cycle analyses following siRNA-mediated down-regulation

Percentage of Cells in						
Condition	<G₀/G₁	G₀/G₁	S-Phase	G₂/M	>G₂M	FI >G₂/M*
Untransfected	1.02	38.62	35.55	23.04	1.55	1.0
GAPD	0.12	43.27	43.42	11.69	1.72	1.11×
SMC1L1	1.65	28.26	42.97	17.29	9.55	6.16×
CSPG6	1.37	28.65	41.86	19.18	8.62	5.56×
STAG2	1.58	28.87	44.48	18.81	5.78	3.72×
MRE11A	2.35	51.59	28.08	14.12	3.08	1.99×

*FI, fold increase over the untransfected control cells.