

**pri-mir-518d**

chr19: 53734861-53734985 (+)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	9	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	10	Suboptimal
3' basal segment length (nt)	24	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.66 (311 / 476)	0.81 (410 / 476)
Cleavage homogeneity	0.51 (304 / 476)	0.76 (302 / 476)

Rank among  
tested pri-miRs