

pri-mir-518a-1

chr19: 53730989-53731113 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	14	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.61 (321 / 476)	1.19 (380 / 476)
Cleavage homogeneity	0.45 (329 / 476)	0.91 (222 / 476)

Rank among
tested pri-miRs