

pri-mir-518e

chr19: 53729823-53729947 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	3	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	22	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.88 (238 / 476)	1.63 (255 / 476)
Cleavage homogeneity	0.8 (194 / 476)	0.54 (378 / 476)

