

pri-mir-516a-2
chr19: 53761120-53761244 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	0	Optimal
Bulged bases in the stem	5	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.06 (145 / 476)	1.25 (369 / 476)
Cleavage homogeneity	1.13 (28 / 476)	0.92 (221 / 476)

