

pri-mir-516b-1

chr19: 53736832-53736956 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	5	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	4	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.97 (189 / 476)	1.63 (256 / 476)
Cleavage homogeneity	0.65 (261 / 476)	0.88 (247 / 476)

