

pri-mir-1278

chr1: 193136484-193136608 (+)

	Features	Optimality
Stem length (bp)	31	Suboptimal
Lower stem length (bp)	8	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	29	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.69 (304 / 476)	1.4 (325 / 476)
Cleavage homogeneity	0.39 (355 / 476)	0.57 (368 / 476)

