

pri-mir-383

chr8: 14853410-14853534 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	17	Optimal
3' basal segment length (nt)	18	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.85 (252 / 476)	1.4 (326 / 476)
Cleavage homogeneity	0.45 (330 / 476)	0.52 (384 / 476)

