

pri-mir-582

chr5: 59703595-59703719 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	16	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.97 (187 / 476)	1.54 (284 / 476)
Cleavage homogeneity	1.02 (84 / 476)	1.0 (156 / 476)

