

pri-mir-296

chr20: 58817590-58817714 (-)

	Features	Optimality
Stem length (bp)	30	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	3	
Apical loop size (nt)	16	Optimal
3' basal segment length (nt)	13	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.32 (40 / 476)	2.0 (78 / 476)
Cleavage homogeneity	0.56 (283 / 476)	0.6 (360 / 476)

