

pri-mir-197

chr1: 109598872-109598996 (+)

	Features	Optimality
Stem length (bp)	31	Suboptimal
Lower stem length (bp)	8	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	17	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.03 (158 / 476)	1.8 (162 / 476)
Cleavage homogeneity	0.74 (222 / 476)	0.83 (278 / 476)

