

pri-mir-122
chr18: 58451057-58451181 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	2	
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.28 (387 / 476)	1.64 (250 / 476)
Cleavage homogeneity	0.4 (350 / 476)	1.1 (63 / 476)

