

pri-mir-522

chr19: 53751196-53751320 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	2	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	23	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.42 (366 / 476)	1.91 (114 / 476)
Cleavage homogeneity	0.51 (302 / 476)	0.86 (257 / 476)

