

pri-mir-3121

chr1: 180438287-180438411 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	11	Suboptimal
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	0.0 (476 / 476)	1.81 (159 / 476)
Cleavage homogeneity	0.1 (430 / 476)	0.67 (345 / 476)

