

pri-mir-1-2

chr18: 21828981-21829105 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	24	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.14 (105 / 476)	1.62 (260 / 476)
Cleavage homogeneity	1.02 (83 / 476)	1.05 (110 / 476)

