

pri-mir-30d
chr8: 134804846-134804970 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	3	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.09 (132 / 476)	1.99 (82 / 476)
Cleavage homogeneity	1.17 (11 / 476)	1.18 (21 / 476)

