

pri-mir-181d

chr19: 13874882-13875006 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	4	Optimal
Apical loop size (nt)	16	Optimal
3' basal segment length (nt)	7	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.09 (412 / 476)	0.09 (450 / 476)
Cleavage homogeneity	0.17 (404 / 476)	0.14 (447 / 476)

