

pri-mir-3186

chr17: 81451081-81451205 (-)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	3	
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	12	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.0 (476 / 476)
Cleavage homogeneity	0.08 (439 / 476)	0.05 (459 / 476)

