

pri-mir-3605

chr1: 33332374-33332498 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	11	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	17	Optimal
3' basal segment length (nt)	3	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.0 (476 / 476)
Cleavage homogeneity	0.04 (457 / 476)	0.0 (476 / 476)

