

pri-mir-2278

chr9: 94809951-94810075 (+)

	Features	Optimality
Stem length (bp)	30	Suboptimal
Lower stem length (bp)	8	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	3	
Apical loop size (nt)	20	Optimal
3' basal segment length (nt)	22	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.12 (406 / 476)	0.65 (420 / 476)
Cleavage homogeneity	0.04 (459 / 476)	0.51 (389 / 476)

