

pri-mir-302d

chr4: 112647972-112648096 (-)

	Features	Optimality
Stem length (bp)	31	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	3	Optimal
Apical loop size (nt)	19	Optimal
3' basal segment length (nt)	24	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.91 (224 / 476)	1.92 (113 / 476)
Cleavage homogeneity	0.83 (178 / 476)	0.67 (346 / 476)

