## pri-mir-301b

chr22: 21652960-21653084 (+)

	Features	Optimality
Stem length (bp)	31	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	5	
Apical loop size (nt)	17	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	1.42 (21 / 476)	2.1 (44 / 476)	
	Cleavage homogeneity	1.02 (85 / 476)	0.98 (169 / 476)	



