## pri-mir-31

chr9: 21512085-21512209 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	8	Suboptimal
3' basal segment length (nt)	27	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

		w/o SRSF3	w/ SRSF3	
	Cleavage efficiency	1.28 (50 / 476)	1.65 (243 / 476)	
	Cleavage homogeneity	0.71 (241 / 476)	0.89 (232 / 476)	

Rank among tested pri-miRs

Top

Bottom

