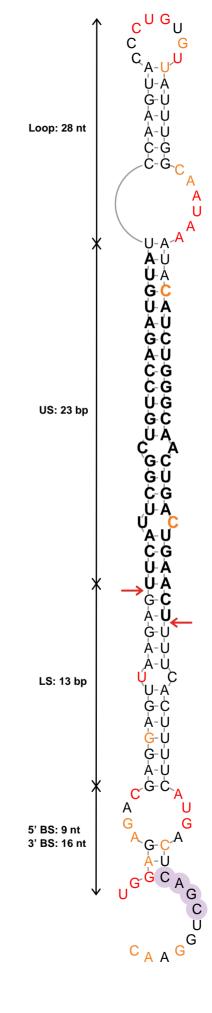
pri-mir-1298

chrX: 114715228-114715352 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	28	Optimal
3' basal segment length (nt)	16	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.01 (168 / 476)	2.1 (46 / 476)
Cleavage homogeneity	1.03 (76 / 476)	1.06 (92 / 476)



Rank among tested pri-miRs

Top

Bottom