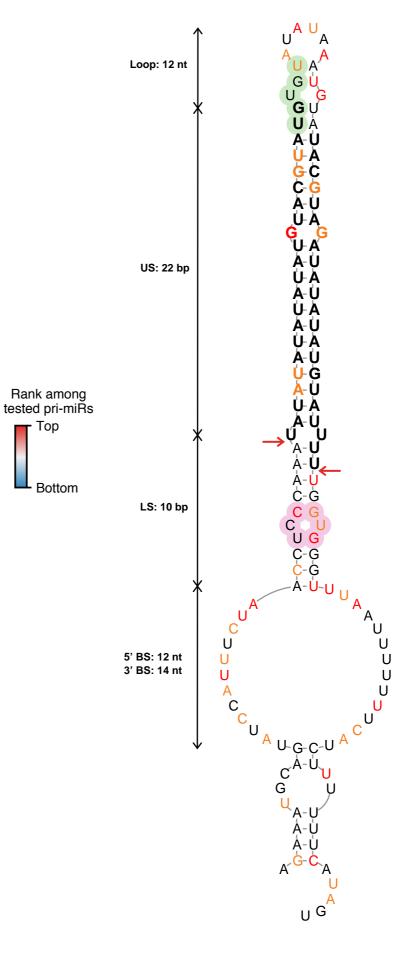
pri-mir-1277

chrX: 118386372-118386496 (+)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	14	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

		w/o SRSF3	w/ SRSF3
	Cleavage efficiency	0.92 (215 / 476)	2.11 (42 / 476)
	Cleavage homogeneity	0.76 (208 / 476)	0.97 (174 / 476)



Top

Bottom