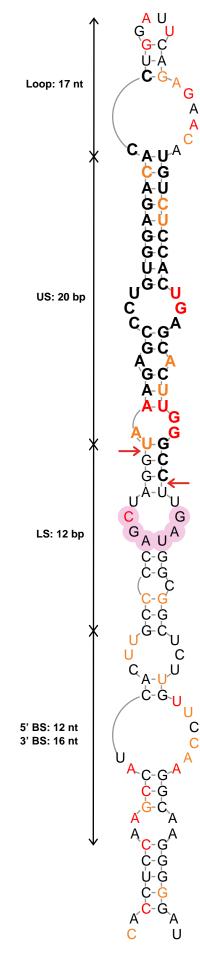
pri-mir-1276

chr15: 85770474-85770598 (-)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	8	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	17	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	0.9 (226 / 476)	1.43 (316 / 476)
Cleavage homogeneity	1.03 (69 / 476)	0.97 (180 / 476)



Rank among tested pri-miRs

Top

Bottom