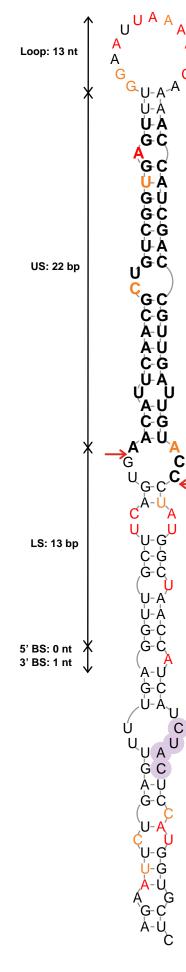
pri-mir-181a-1

chr1: 198859036-198859160 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	4	
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.94 (199 / 476)	1.82 (157 / 476)
Cleavage homogeneity	1.09 (40 / 476)	1.06 (95 / 476)



Rank among tested pri-miRs

Top

Bottom