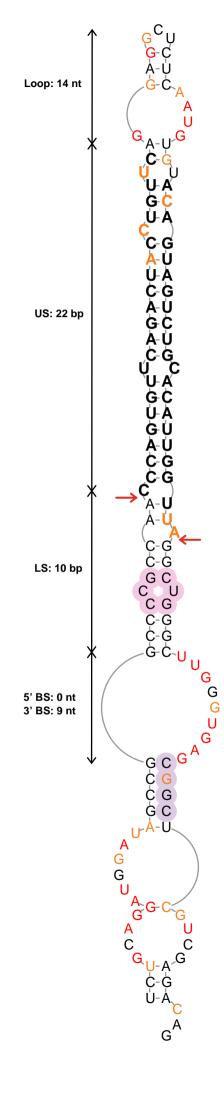
## pri-mir-199a-1

chr19: 10817396-10817520 (-)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	3	
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	9	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.76 (280 / 476)	2.05 (62 / 476)
Cleavage homogeneity	0.42 (340 / 476)	0.87 (249 / 476)



Rank among tested pri-miRs

Top

Bottom