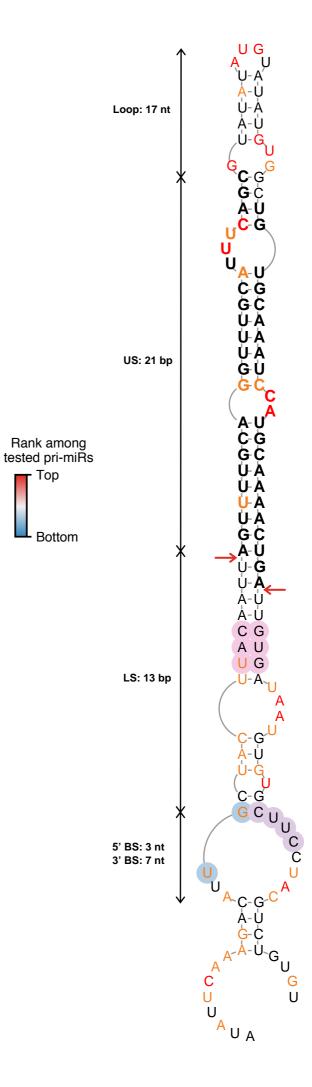
## pri-mir-19b-2

chrX: 134169652-134169776 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	0	Optimal
Bulged bases in the stem	10	
Apical loop size (nt)	17	Optimal
3' basal segment length (nt)	7	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.85 (254 / 476)	1.43 (315 / 476)
Cleavage homogeneity	1.08 (44 / 476)	1.04 (120 / 476)



Top

Bottom