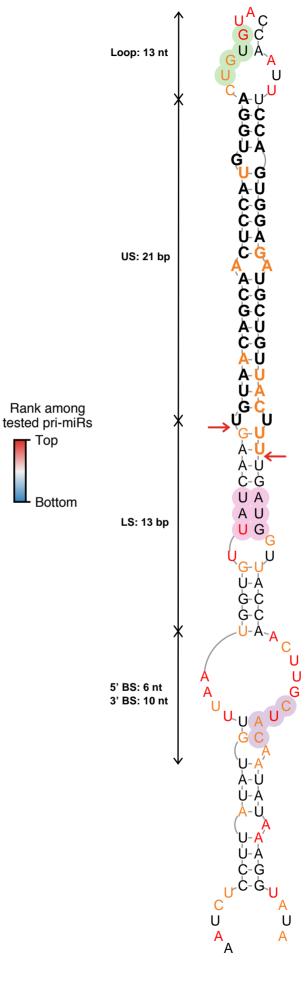
pri-mir-194-1

chr1: 220118135-220118259 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	10	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.78 (276 / 476)	1.5 (294 / 476)
Cleavage homogeneity	0.78 (203 / 476)	0.83 (275 / 476)



Top

Bottom