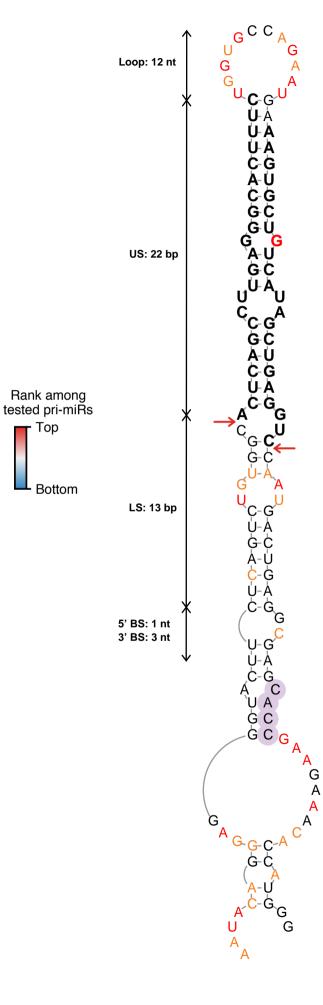
pri-mir-512-1

chr19: 53666662-53666786 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	3	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.53 (345 / 476)	1.6 (265 / 476)
Cleavage homogeneity	0.34 (371 / 476)	0.86 (255 / 476)



Top

Bottom