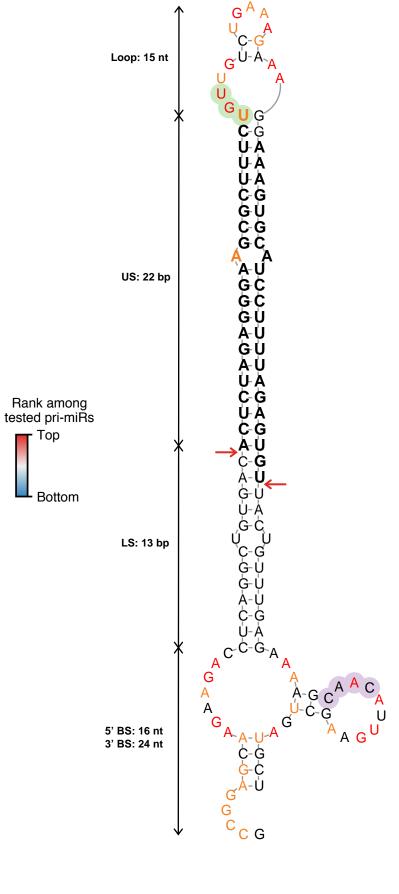
pri-mir-519a-1

chr19: 53752381-53752505 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	24	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.98 (179 / 476)	1.96 (90 / 476)
Cleavage homogeneity	0.93 (126 / 476)	1.13 (47 / 476)



Top