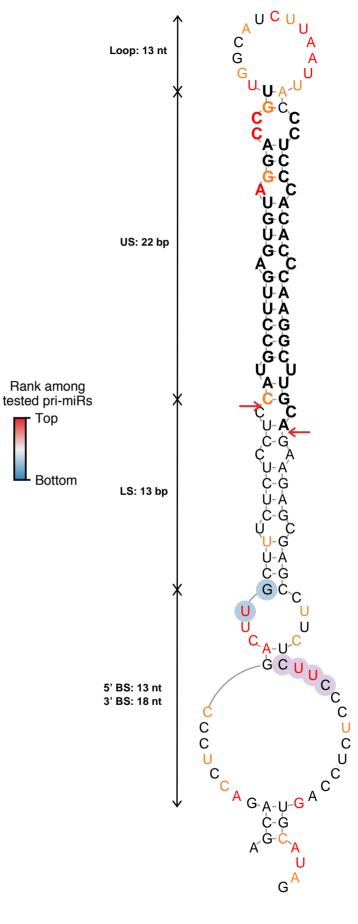
pri-mir-532

chrX: 50003137-50003261 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	8	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	18	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.21 (79 / 476)	1.95 (96 / 476)
Cleavage homogeneity	0.95 (111 / 476)	0.95 (198 / 476)



Top