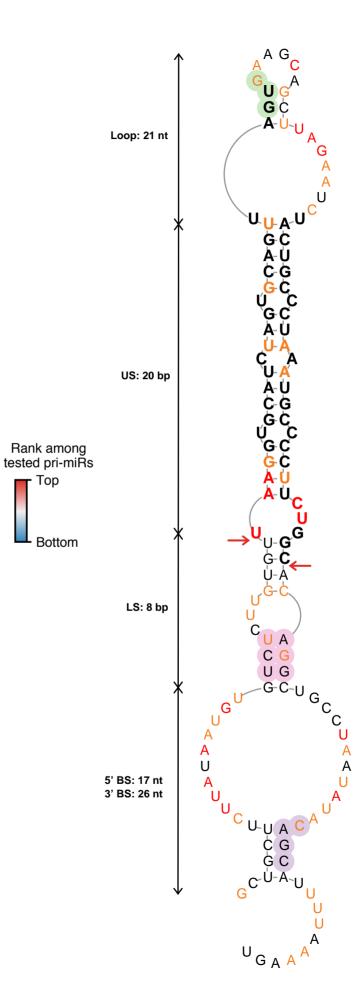
pri-mir-18b

chrX: 134170010-134170134 (-)

	Features	Optimality
Stem length (bp)	28	Suboptimal
Lower stem length (bp)	8	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	5	Optimal
Apical loop size (nt)	21	Optimal
3' basal segment length (nt)	26	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.7 (299 / 476)	1.44 (310 / 476)
Cleavage homogeneity	0.9 (147 / 476)	1.05 (100 / 476)



Top