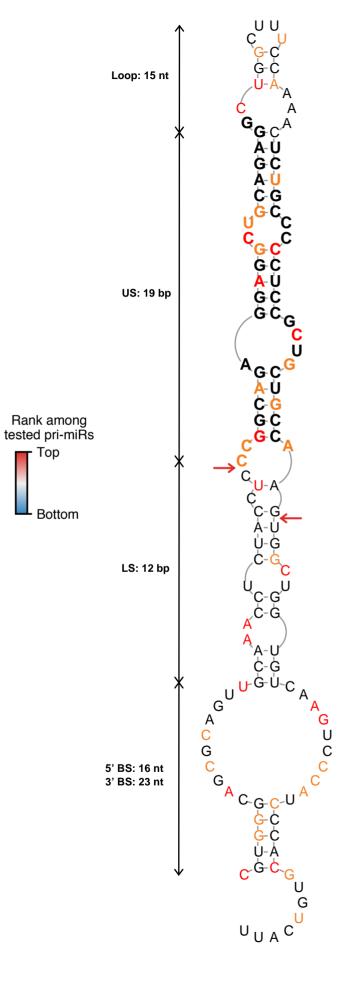
pri-mir-1913

chr6: 166509329-166509453 (-)

	Features	Optimality
Stem length (bp)	31	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	9	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	23	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.02 (424 / 476)	0.38 (432 / 476)
Cleavage homogeneity	0.09 (434 / 476)	0.3 (430 / 476)



Top