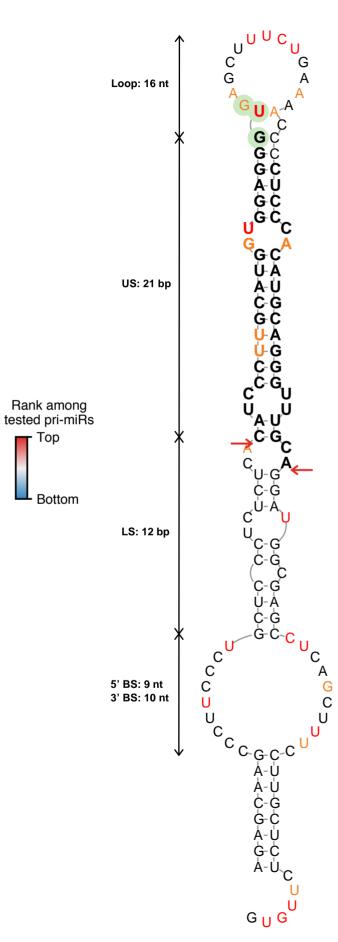
## pri-mir-188

chrX: 50003487-50003611 (+)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	16	Optimal
3' basal segment length (nt)	10	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.17 (94 / 476)	2.06 (58 / 476)
Cleavage homogeneity	1.15 (16 / 476)	1.19 (13 / 476)



Top

**Bottom**