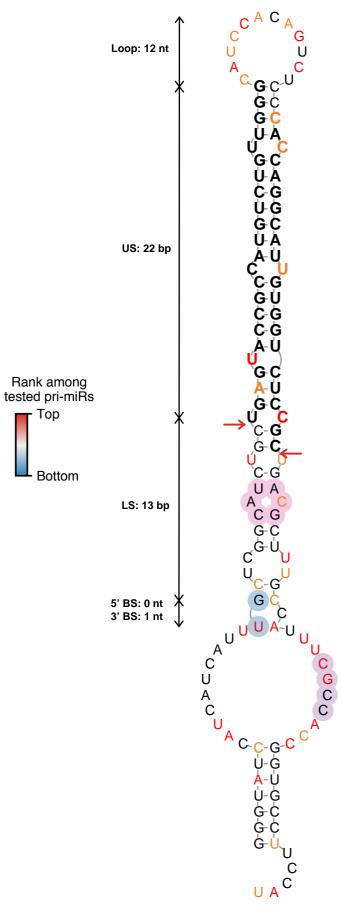
## pri-mir-1911

chrX: 114763164-114763288 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.04 (154 / 476)	2.0 (77 / 476)
Cleavage homogeneity	1.08 (45 / 476)	1.19 (11 / 476)



Top

**Bottom**