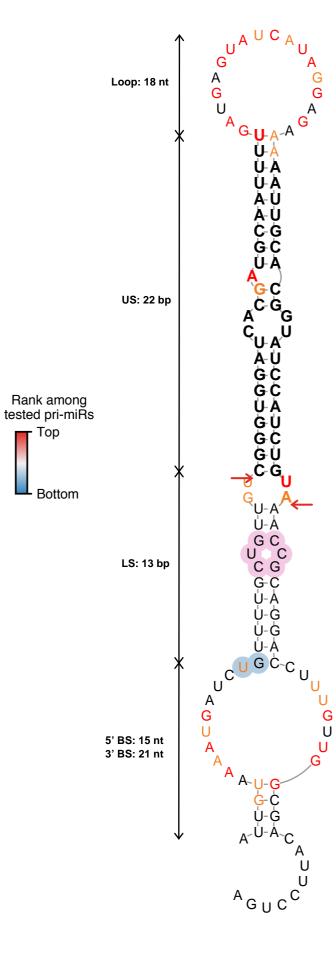
## pri-mir-363

chrX: 134169350-134169474 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	18	Optimal
3' basal segment length (nt)	21	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.98 (181 / 476)	1.24 (371 / 476)
Cleavage homogeneity	0.89 (151 / 476)	1.06 (99 / 476)



Top

**Bottom**