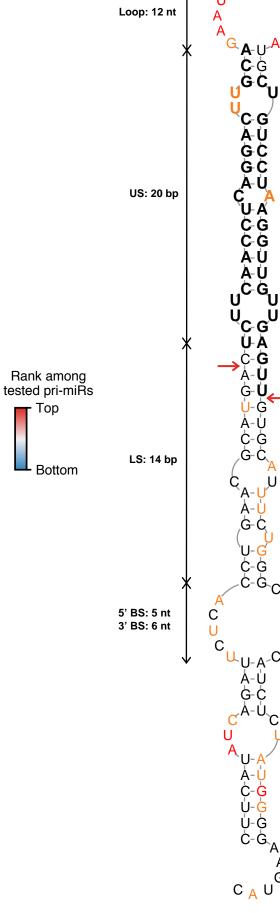
## pri-mir-676

chrX: 70022832-70022956 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	3	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.37 (371 / 476)	1.63 (252 / 476)
Cleavage homogeneity	0.22 (394 / 476)	0.49 (394 / 476)



Top

**Bottom**