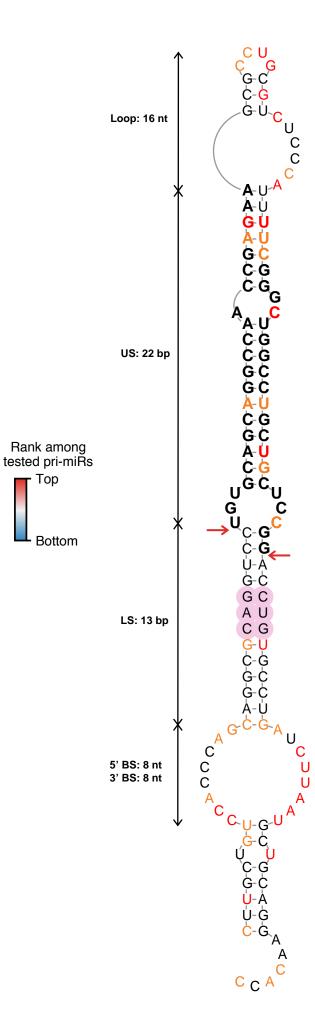
## pri-mir-3944

chr10: 133371546-133371670 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	16	Optimal
3' basal segment length (nt)	8	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

		w/o SRSF3	w/ SRSF3	
	Cleavage efficiency	0.94 (200 / 476)	1.3 (356 / 476)	
	Cleavage homogeneity	0.43 (339 / 476)	0.37 (419 / 476)	



Top