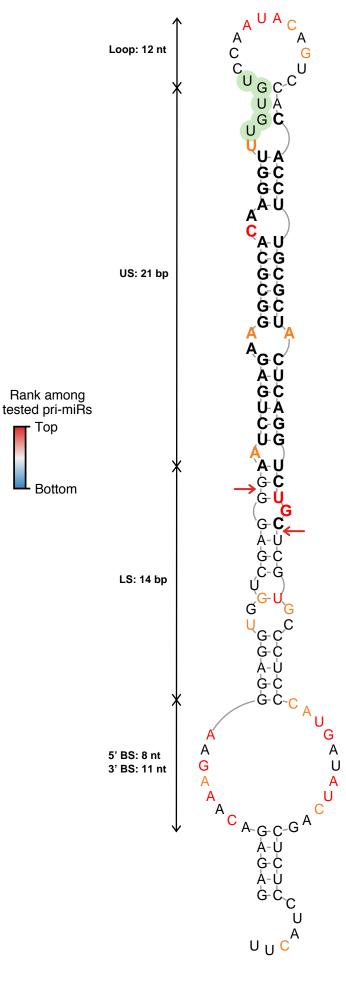
## pri-mir-3200

chr22: 30731541-30731665 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	8	
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	11	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

		w/o SRSF3	w/ SRSF3	
	Cleavage efficiency	0.97 (190 / 476)	1.49 (297 / 476)	
	Cleavage homogeneity	1.03 (78 / 476)	0.97 (177 / 476)	



Top