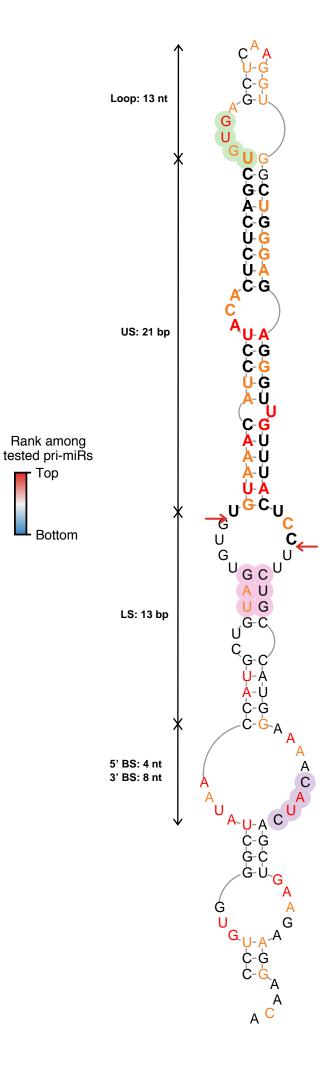
pri-mir-30c-1

chr1: 40757271-40757395 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	6	
Apical loop size (nt)	13	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	0.65 (314 / 476)	1.5 (293 / 476)
Cleavage homogeneity	0.81 (185 / 476)	1.05 (101 / 476)



Top

Bottom