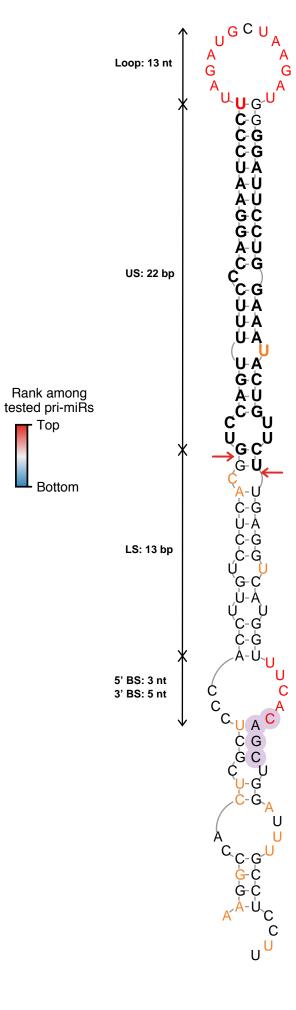
pri-mir-145

chr5: 149430631-149430755 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	3	
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.94 (198 / 476)	1.79 (172 / 476)
Cleavage homogeneity	0.34 (369 / 476)	1.02 (132 / 476)



Top

Bottom