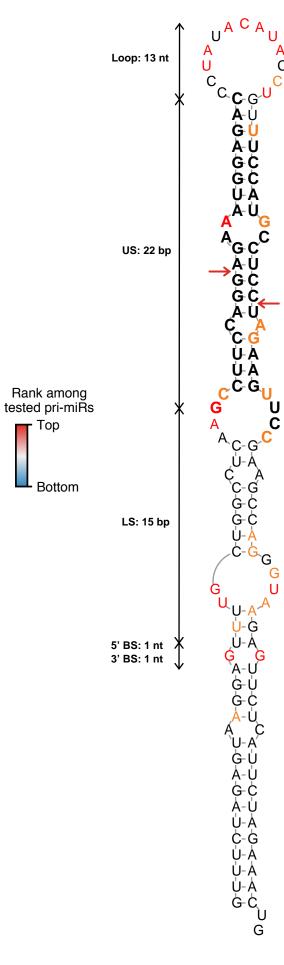
pri-mir-5581

chr1: 37500901-37501025 (-)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	10	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.0 (476 / 476)
Cleavage homogeneity	0.17 (407 / 476)	0.05 (460 / 476)



Тор

Bottom