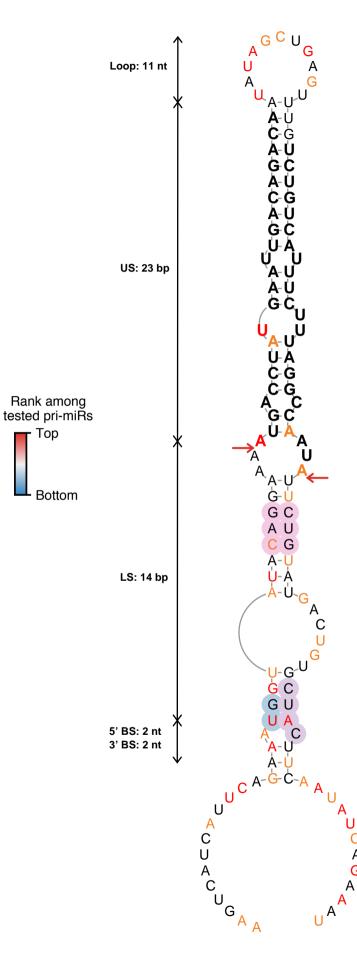
## pri-mir-215

chr1: 220117843-220117967 (-)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	7	Optimal
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	2	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.39 (368 / 476)	1.29 (362 / 476)
Cleavage homogeneity	0.36 (365 / 476)	0.88 (245 / 476)



Top

Bottom