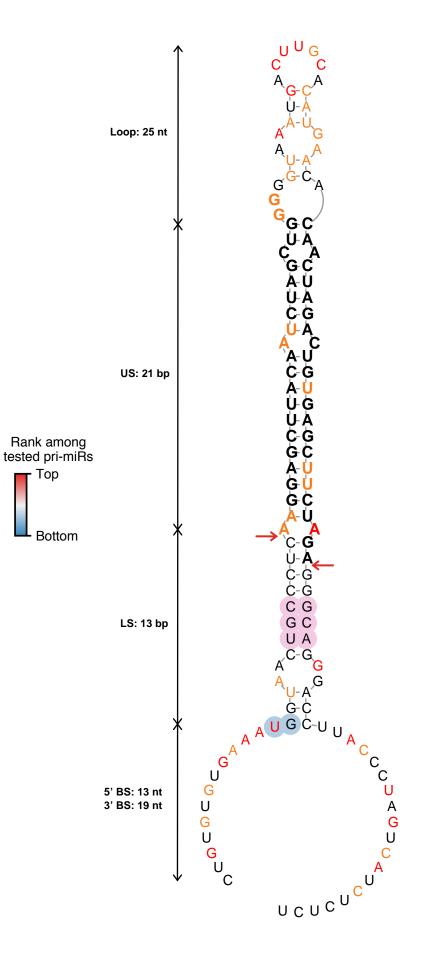
pri-mir-708

chr11: 79402001-79402125 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	25	Optimal
3' basal segment length (nt)	19	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.13 (114 / 476)	2.12 (38 / 476)
Cleavage homogeneity	0.98 (106 / 476)	1.02 (138 / 476)



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