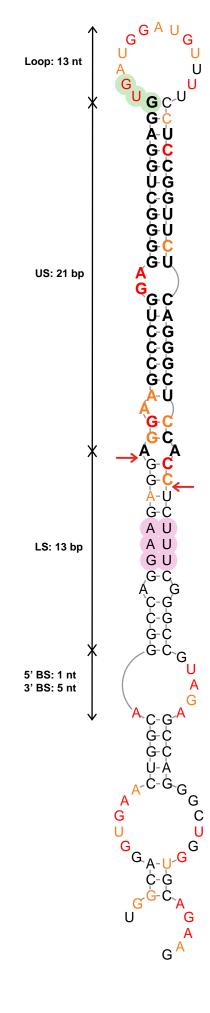
pri-mir-671

chr7: 151238419-151238543 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	2	
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	1.33 (38 / 476)	1.39 (334 / 476)
Cleavage homogeneity	0.9 (142 / 476)	0.82 (281 / 476)



Rank among tested pri-miRs

Top

Bottom