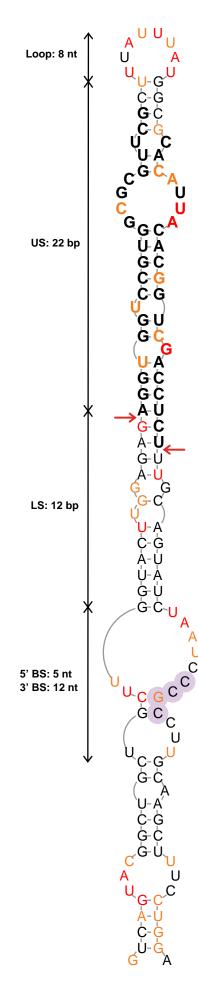
pri-mir-323a

chr14: 101025715-101025839 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	3	
Apical loop size (nt)	8	Suboptimal
3' basal segment length (nt)	12	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.63 (316 / 476)	1.63 (257 / 476)
Cleavage homogeneity	0.27 (382 / 476)	0.81 (284 / 476)



Rank among tested pri-miRs

Top

Bottom