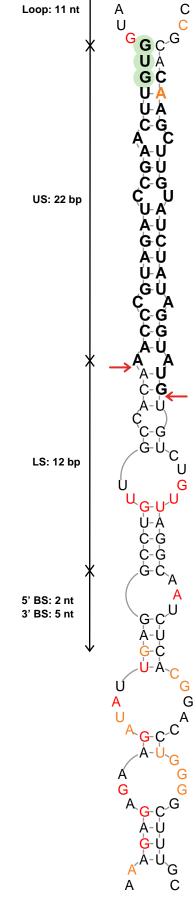
pri-mir-100

chr11: 122152204-122152328 (-)

	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	Optimal
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.87 (243 / 476)	1.65 (240 / 476)
Cleavage homogeneity	0.74 (224 / 476)	0.97 (176 / 476)



Rank among tested pri-miRs

Top

Bottom