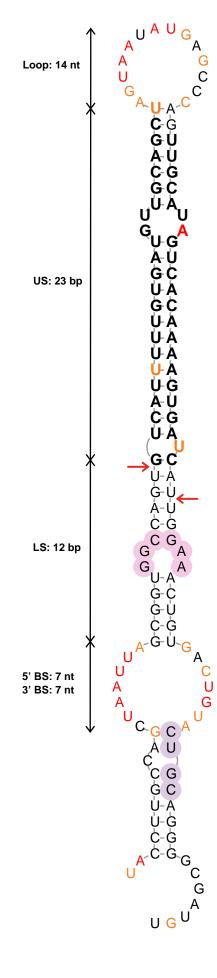
pri-mir-153-2

chr7: 157574315-157574439 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	7	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

		w/o SRSF3	w/ SRSF3	
	Cleavage efficiency	1.11 (124 / 476)	1.75 (187 / 476)	
	Cleavage homogeneity	0.93 (122 / 476)	1.05 (106 / 476)	



Rank among tested pri-miRs

Top

Bottom