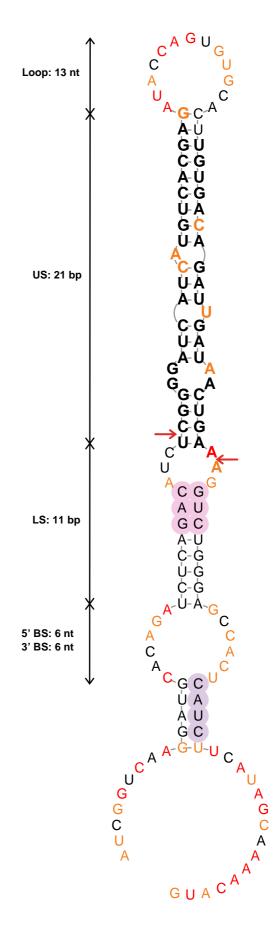
pri-mir-542

chrX: 134541329-134541453 (-)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	2	
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.15 (399 / 476)	0.98 (399 / 476)
Cleavage homogeneity	0.1 (429 / 476)	0.44 (409 / 476)



Rank among tested pri-miRs

Top

Bottom