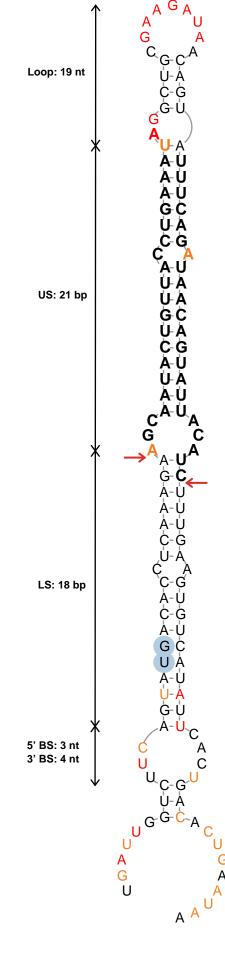
## pri-mir-3942

chr15: 35372247-35372371 (-)

	Features	Optimality
Stem length (bp)	39	Suboptimal
Lower stem length (bp)	18	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	19	Optimal
3' basal segment length (nt)	4	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.6 (326 / 476)	1.74 (194 / 476)
Cleavage homogeneity	0.15 (415 / 476)	0.5 (390 / 476)



Rank among tested pri-miRs

Top

**Bottom**