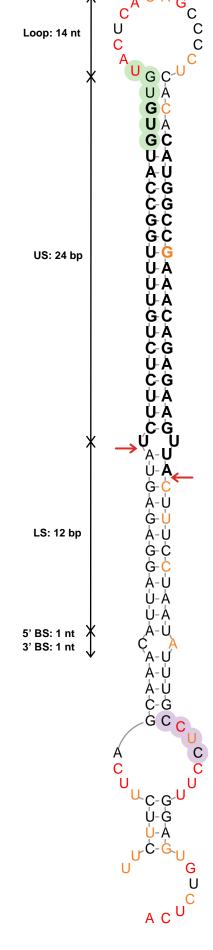
pri-mir-942

chr1: 117094627-117094751 (+)

	Features	Optimality
Stem length (bp)	36	Optimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	1.41 (322 / 476)
Cleavage homogeneity	0.09 (435 / 476)	0.47 (399 / 476)



Rank among tested pri-miRs

Top

Bottom