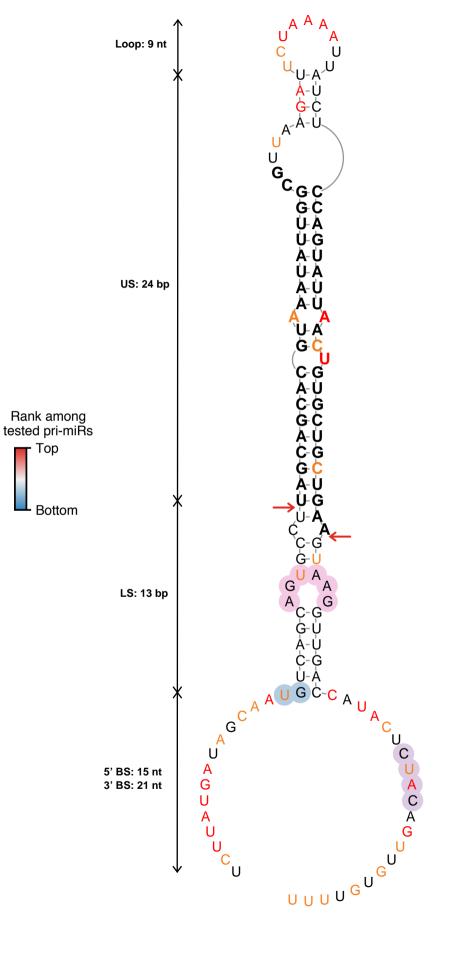
## pri-mir-16-1

chr13: 50048952-50049076 (-)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	6	
Apical loop size (nt)	9	Suboptimal
3' basal segment length (nt)	21	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

		w/o SRSF3	w/ SRSF3
	Cleavage efficiency	0.92 (212 / 476)	1.72 (204 / 476)
	Cleavage homogeneity	0.83 (174 / 476)	0.85 (266 / 476)



Top