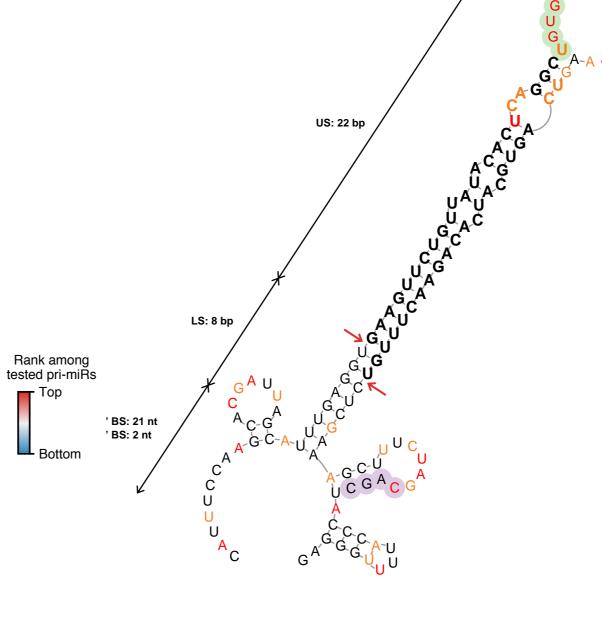
pri-mir-148b

chr12: 54337210-54337334 (+)

	Features	Optimality
Stem length (bp)	30	Suboptimal
Lower stem length (bp)	8	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	2	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

		w/o SRSF3	w/ SRSF3	
	Cleavage efficiency	0.6 (327 / 476)	1.38 (336 / 476)	
	Cleavage homogeneity	0.52 (297 / 476)	1.01 (148 / 476)	



Loop: 13 nt