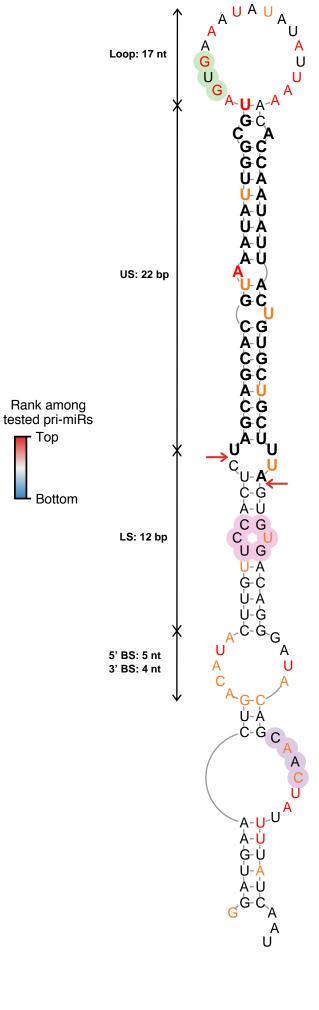
pri-mir-16-2

chr3: 160404727-160404851 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	
Apical loop size (nt)	17	Optimal
3' basal segment length (nt)	4	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.97 (184 / 476)	1.72 (206 / 476)
Cleavage homogeneity	0.81 (189 / 476)	0.94 (199 / 476)



Top