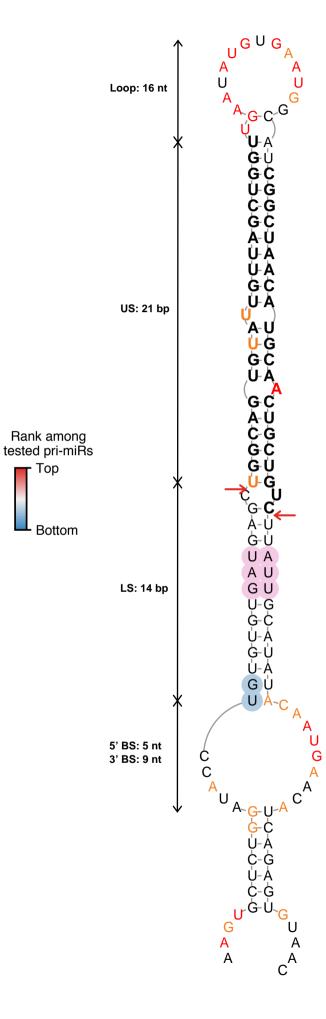
pri-mir-449a

chr5: 55170513-55170637 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	2	
Apical loop size (nt)	16	Optimal
3' basal segment length (nt)	9	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.14 (108 / 476)	1.71 (210 / 476)
Cleavage homogeneity	1.12 (29 / 476)	0.96 (187 / 476)



Top