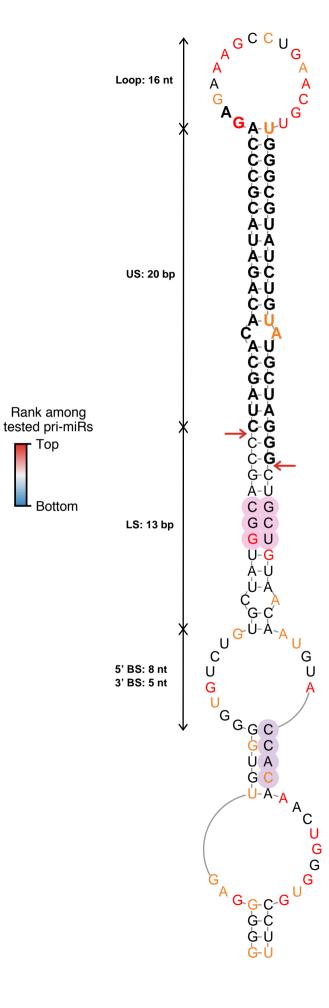
pri-mir-585

chr5: 169263579-169263703 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	16	Optimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.12 (118 / 476)	2.18 (27 / 476)
Cleavage homogeneity	0.93 (129 / 476)	1.16 (25 / 476)



Top

Bottom