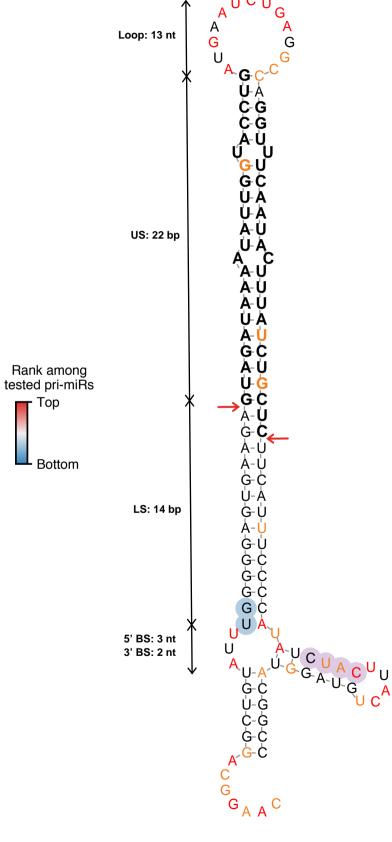
pri-mir-577

chr4: 114656743-114656867 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	0	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	1.11 (122 / 476)	1.86 (138 / 476)
Cleavage homogeneity	0.99 (100 / 476)	1.08 (79 / 476)



Top

Bottom