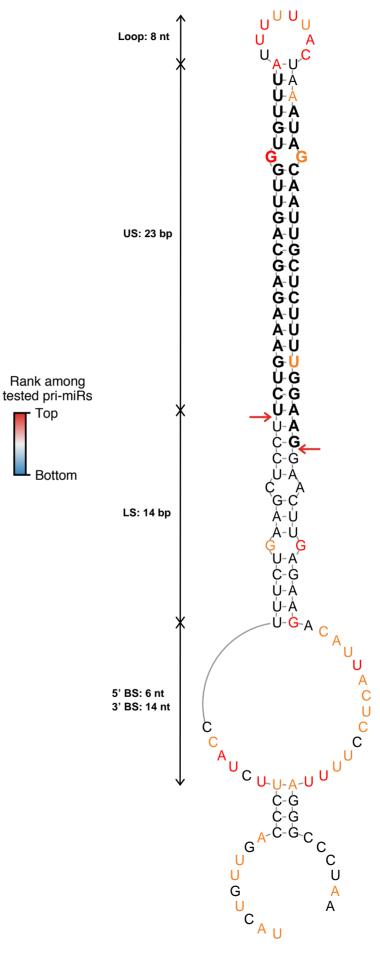
## pri-mir-4766

chr22: 40813856-40813980 (-)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	8	Suboptimal
3' basal segment length (nt)	14	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.15 (103 / 476)	1.98 (87 / 476)
Cleavage homogeneity	0.46 (321 / 476)	0.51 (388 / 476)



Top

**Bottom**