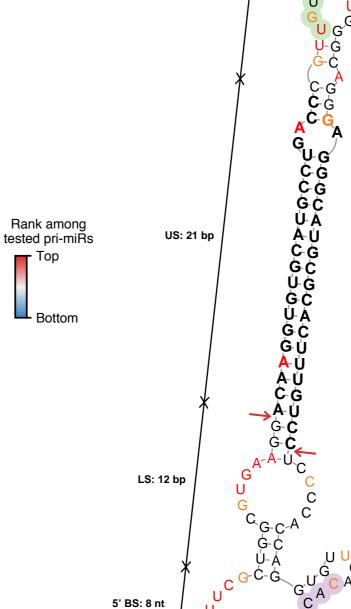
## pri-mir-4761

chr22: 19963735-19963859 (+)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	35	Optimal
3' basal segment length (nt)	15	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.96 (191 / 476)	1.84 (151 / 476)
Cleavage homogeneity	0.67 (254 / 476)	0.84 (272 / 476)



3' BS: 15 nt

Loop: 35 nt