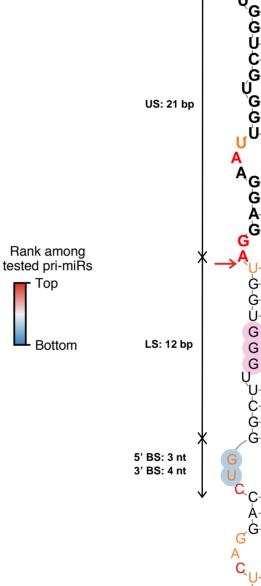
pri-mir-766

chrX: 119646728-119646852 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	8	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	4	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.25 (392 / 476)	1.06 (392 / 476)
Cleavage homogeneity	0.1 (432 / 476)	0.48 (398 / 476)



Loop: 13 nt