

pri-mir-106b

chr7: 100093969-100094093 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.05 (149 / 476)	1.85 (149 / 476)
Cleavage homogeneity	0.67 (252 / 476)	1.12 (52 / 476)

