

pri-mir-5194

chr8: 130008326-130008450 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	7	
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	8	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.02 (425 / 476)	0.14 (445 / 476)
Cleavage homogeneity	0.04 (458 / 476)	0.2 (440 / 476)

