

pri-mir-191  
chr3: 49020599-49020723 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	17	Optimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.53 (346 / 476)	1.57 (277 / 476)
Cleavage homogeneity	0.46 (323 / 476)	0.86 (256 / 476)

