

pri-mir-32
chr9: 109046199-109046323 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	3	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.18 (87 / 476)	1.94 (100 / 476)
Cleavage homogeneity	1.15 (18 / 476)	0.95 (196 / 476)

