

pri-mir-5009
chr15: 89883916-89884040 (-)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	3	
Apical loop size (nt)	13	Optimal
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	0.44 (361 / 476)	0.87 (406 / 476)
Cleavage homogeneity	0.25 (386 / 476)	0.28 (433 / 476)

