

pri-mir-3940

chr19: 6416398-6416522 (-)

	Features	Optimality
Stem length (bp)	30	Suboptimal
Lower stem length (bp)	7	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	6	
Apical loop size (nt)	6	Suboptimal
3' basal segment length (nt)	19	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.0 (476 / 476)
Cleavage homogeneity	0.05 (456 / 476)	0.0 (476 / 476)

