

pri-mir-3918

chr6: 158764642-158764766 (-)

	Features	Optimality
Stem length (bp)	39	Suboptimal
Lower stem length (bp)	16	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	6	
Apical loop size (nt)	9	Suboptimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	0.0 (476 / 476)	0.0 (476 / 476)	<div><div></div></div> Top Bottom
Cleavage homogeneity	0.07 (446 / 476)	0.04 (462 / 476)	

