

pri-mir-4677

chr1: 243346158-243346282 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	7	Optimal
Apical loop size (nt)	8	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	0.0 (476 / 476)	0.0 (476 / 476)	

