

pri-mir-218-2

chr5: 168768136-168768260 (-)

	Features	Optimality
Stem length (bp)	29	Suboptimal
Lower stem length (bp)	8	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	20	Optimal
3' basal segment length (nt)	10	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	0.89 (232 / 476)	1.85 (145 / 476)	<div><div></div>Top</div> <div><div></div>Bottom</div>
Cleavage homogeneity	0.53 (292 / 476)	1.07 (85 / 476)	

