

pri-mir-769

chr19: 46018932-46019056 (+)

	Features	Optimality
Stem length (bp)	31	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	2	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	0	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.3 (43 / 476)	1.37 (337 / 476)
Cleavage homogeneity	0.45 (333 / 476)	0.39 (416 / 476)

Rank among
tested pri-miRs

