

chr19: 53741302-53741426 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	24	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.91 (220 / 476)	1.85 (147 / 476)
Cleavage homogeneity	0.44 (335 / 476)	1.0 (155 / 476)

Rank among
tested pri-miRs