

chr11: 79422134-79422258 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	16	Optimal
3' basal segment length (nt)	23	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.01 (427 / 476)	1.62 (258 / 476)
Cleavage homogeneity	0.22 (395 / 476)	0.73 (318 / 476)

Rank among
tested pri-miRs

