

**pri-mir-330**

chr19: 45638975-45639099 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	6	
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	2	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.45 (17 / 476)	1.98 (83 / 476)
Cleavage homogeneity	1.21 (2 / 476)	1.19 (8 / 476)

Rank among tested pri-miRs

Top

Bottom

