

**pri-mir-515-1**

chr19: 53678986-53679110 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	7	Suboptimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.89 (235 / 476)	1.56 (279 / 476)
Cleavage homogeneity	0.91 (136 / 476)	1.08 (82 / 476)

Rank among  
tested pri-miRs

