

**pri-mir-523**

chr19: 53698370-53698494 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	15	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	0.75 (285 / 476)	1.8 (164 / 476)
Cleavage homogeneity	0.77 (205 / 476)	1.14 (42 / 476)

Rank among  
tested pri-miRs