

**pri-mir-526b**

chr19: 53694376-53694500 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	11	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.83 (263 / 476)	1.23 (374 / 476)
Cleavage homogeneity	0.93 (127 / 476)	0.8 (292 / 476)

Rank among  
tested pri-miRs

 [Top](#)

 Bottom

