

pri-mir-526a-2

chr19: 53726896-53727020 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	5	
Apical loop size (nt)	6	Suboptimal
3' basal segment length (nt)	7	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.37 (372 / 476)	1.41 (323 / 476)
Cleavage homogeneity	0.42 (342 / 476)	1.09 (71 / 476)

Rank among tested pri-miRs

Top

Bottom

