

pri-mir-526a-1

chr19: 53706236-53706360 (+)

	Features	Optimality
Stem length (bp)	27	Suboptimal
Lower stem length (bp)	6	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	22	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.88 (237 / 476)	1.4 (330 / 476)
Cleavage homogeneity	0.81 (184 / 476)	1.04 (118 / 476)

Rank among tested pri-miRs

Top

Bottom