

pri-mir-520d

chr19: 53720081-53720205 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	2	
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	6	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	1.21 (77 / 476)	1.79 (171 / 476)	<div><div></div></div> Top Bottom
Cleavage homogeneity	0.74 (227 / 476)	0.82 (282 / 476)	

