

pri-mir-520f

chr19: 53682144-53682268 (+)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	21	Optimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	1.39 (27 / 476)	1.91 (115 / 476)	<div><div></div></div> Top Bottom
Cleavage homogeneity	1.18 (5 / 476)	1.1 (62 / 476)	

