

**pri-mir-520a**

chr19: 53690865-53690989 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	24	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.82 (264 / 476)	1.31 (354 / 476)
Cleavage homogeneity	0.98 (102 / 476)	0.97 (181 / 476)

Rank among  
tested pri-miRs

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