

**pri-mir-10b**

chr2: 176150300-176150424 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.89 (230 / 476)	1.42 (320 / 476)
Cleavage homogeneity	0.75 (211 / 476)	0.78 (296 / 476)

Rank among  
tested pri-miRs

 Top

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