

**pri-mir-3150b**

chr8: 95072889-95073013 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	14	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	2.0 (4 / 476)	2.98 (2 / 476)
Cleavage homogeneity	0.84 (171 / 476)	0.55 (371 / 476)

Rank among  
tested pri-miRs

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**Bottom**

