

**pri-mir-449b**

chr5: 55170633-55170757 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	5	
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	11	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.31 (382 / 476)	1.18 (384 / 476)
Cleavage homogeneity	0.16 (410 / 476)	0.85 (265 / 476)

Rank among  
tested pri-miRs

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