

**pri-mir-331**

chr12: 95308413-95308537 (+)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	3	Optimal
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	5	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	1.58 (272 / 476)
Cleavage homogeneity	0.2 (397 / 476)	0.75 (307 / 476)

Rank among  
tested pri-miRs

Bottom

