

**pri-mir-944**

chr3: 189829907-189830031 (+)

	Features	Optimality
Stem length (bp)	38	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	0	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.92 (211 / 476)	2.09 (49 / 476)
Cleavage homogeneity	0.98 (105 / 476)	0.88 (241 / 476)

Rank among  
tested pri-miRs