

**pri-mir-448**

chrX: 114823445-114823569 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	31	Optimal
3' basal segment length (nt)	13	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.39 (28 / 476)	2.26 (15 / 476)
Cleavage homogeneity	0.55 (287 / 476)	0.52 (385 / 476)

Rank among  
tested pri-miRs