

**pri-mir-514b**

chrX: 147250126-147250250 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	8	Suboptimal
3' basal segment length (nt)	24	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.08 (135 / 476)	2.09 (48 / 476)
Cleavage homogeneity	1.16 (13 / 476)	1.22 (3 / 476)

Rank among  
tested pri-miRs

