

**pri-mir-106a**

chrX: 134170173-134170297 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	
Apical loop size (nt)	11	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	1.19 (83 / 476)	1.42 (318 / 476)
Cleavage homogeneity	0.5 (305 / 476)	0.57 (369 / 476)

Rank among  
tested pri-miRs

