

**pri-mir-374b**

chrX: 74218519-74218643 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	4	Suboptimal
3' basal segment length (nt)	11	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.21 (441 / 476)
Cleavage homogeneity	0.0 (476 / 476)	0.0 (476 / 476)

Rank among  
tested pri-miRs