

pri-mir-376a-1

chr14: 101040757-101040881 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	3	Optimal
Apical loop size (nt)	5	Suboptimal
3' basal segment length (nt)	7	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.85 (255 / 476)	1.59 (270 / 476)
Cleavage homogeneity	0.56 (281 / 476)	0.83 (276 / 476)

Rank among
tested pri-miRs

