

pri-mir-379

chr14: 101022041-101022165 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	9	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.43 (363 / 476)	1.42 (319 / 476)
Cleavage homogeneity	0.75 (220 / 476)	1.11 (56 / 476)

Rank among
tested pri-miRs

 Top

Bottom

