

pri-mir-369

chr14: 101065574-101065698 (+)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	3	
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	19	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.26 (389 / 476)	1.34 (345 / 476)
Cleavage homogeneity	0.26 (383 / 476)	0.74 (313 / 476)

Rank among
tested pri-miRs

