

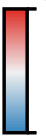
pri-mir-412

chr14: 101065434-101065558 (+)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	4	
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	28	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.28 (386 / 476)	1.56 (280 / 476)
Cleavage homogeneity	0.32 (373 / 476)	1.0 (153 / 476)

Rank among
tested pri-miRs



Top
Bottom

