


pri-mir-212

chr17: 2050256-2050380 (-)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	11	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	3	
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	23	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.04 (421 / 476)	0.05 (453 / 476)
Cleavage homogeneity	0.05 (452 / 476)	0.04 (464 / 476)

Rank among
tested pri-miRs



Top

Bottom

