


pri-mir-217

chr2: 55982949-55983073 (-)

	Features	Optimality
Stem length (bp)	33	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	10	
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	21	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.43 (362 / 476)	1.67 (228 / 476)
Cleavage homogeneity	0.58 (276 / 476)	0.99 (166 / 476)

Rank among
tested pri-miRs



Top

Bottom

