

pri-mir-219a-1

chr6: 33207827-33207951 (+)

	Features	Optimality
Stem length (bp)	38	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	2	
Apical loop size (nt)	9	Suboptimal
3' basal segment length (nt)	11	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.65 (8 / 476)	2.06 (56 / 476)
Cleavage homogeneity	1.09 (41 / 476)	0.7 (331 / 476)

Rank among
tested pri-miRs