

pri-mir-941-1

chr20: 63919427-63919551 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	7	
Apical loop size (nt)	8	Suboptimal
3' basal segment length (nt)	3	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.15 (400 / 476)	0.15 (443 / 476)
Cleavage homogeneity	0.07 (447 / 476)	0.13 (451 / 476)

Rank among tested pri-miRs

Top

Bottom

