## pri-mir-33a

chr22: 41900920-41901044 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	6	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	15	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.49 (12 / 476)	2.29 (9 / 476)
Cleavage homogeneity	1.03 (77 / 476)	1.22 (4 / 476)



