


chr10: 133247502-133247626 (-)

	Features	Optimality
Stem length (bp)	39	Suboptimal
Lower stem length (bp)	15	Suboptimal
Mismatched pairs in the stem	2	
Bulged bases in the stem	3	
Apical loop size (nt)	6	Suboptimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.02 (426 / 476)	0.57 (423 / 476)
Cleavage homogeneity	0.12 (419 / 476)	0.49 (392 / 476)



Top

Bottom

