

chr11: 43559645-43559769 (+)

	Features	Optimality
Stem length (bp)	37	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	5	
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	2	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.03 (423 / 476)	0.88 (405 / 476)
Cleavage homogeneity	0.08 (441 / 476)	0.33 (425 / 476)

