

pri-mir-885

chr3: 10394462-10394586 (-)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	Optimal
Apical loop size (nt)	6	Suboptimal
3' basal segment length (nt)	26	Optimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.39 (370 / 476)	2.22 (21 / 476)
Cleavage homogeneity	0.49 (311 / 476)	0.88 (239 / 476)

