pri-mir-2355

chr2: 207109967-207110091 (-)

	Features	Optimality
Stem length (bp)	38	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	4	
Apical loop size (nt)	9	Suboptimal
3' basal segment length (nt)	1	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.48 (353 / 476)	1.79 (167 / 476)
Cleavage homogeneity	0.45 (327 / 476)	0.8 (291 / 476)



