

pri-mir-520e

chr19: 53675696-53675820 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	5	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	15	Optimal
3' basal segment length (nt)	4	Suboptimal
Mismatch at -6	Possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.85 (256 / 476)	1.15 (387 / 476)
Cleavage homogeneity	0.45 (331 / 476)	0.53 (380 / 476)

