

pri-mir-377

chr14: 101062026-101062150 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	14	Suboptimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	2	
Apical loop size (nt)	12	Suboptimal
3' basal segment length (nt)	14	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.17 (93 / 476)	1.79 (170 / 476)
Cleavage homogeneity	0.85 (169 / 476)	0.84 (274 / 476)

