

pri-mir-378a

chr5: 149732799-149732923 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	6	Optimal
Apical loop size (nt)	14	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	1.56 (10 / 476)	2.28 (12 / 476)
Cleavage homogeneity	1.12 (32 / 476)	1.03 (124 / 476)

