

pri-mir-429

chr1: 1168989-1169113 (+)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	10	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	10	Suboptimal
3' basal segment length (nt)	7	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	1.6 (269 / 476)
Cleavage homogeneity	0.12 (420 / 476)	0.88 (242 / 476)

