

pri-mir-576

chr4: 109488684-109488808 (+)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	0	Optimal
Apical loop size (nt)	14	Optimal
3' basal segment length (nt)	8	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.03 (161 / 476)	1.87 (135 / 476)
Cleavage homogeneity	1.02 (86 / 476)	1.16 (27 / 476)

Rank among
tested pri-miRs