

pri-mir-561

chr2: 188297486-188297610 (+)

	Features	Optimality
Stem length (bp)	35	Optimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	11	Suboptimal
3' basal segment length (nt)	2	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.16 (97 / 476)	2.21 (23 / 476)
Cleavage homogeneity	0.82 (181 / 476)	0.93 (215 / 476)

Rank among
tested pri-miRs

 Top

 Bottom

