

pri-mir-616

chr12: 57519150-57519274 (-)

	Features	Optimality
Stem length (bp)	36	Suboptimal
Lower stem length (bp)	13	Optimal
Mismatched pairs in the stem	3	
Bulged bases in the stem	1	Optimal
Apical loop size (nt)	13	Optimal
3' basal segment length (nt)	22	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	1.27 (54 / 476)	1.76 (183 / 476)
Cleavage homogeneity	0.91 (139 / 476)	1.04 (116 / 476)

Rank among
tested pri-miRs