

pri-mir-588

chr6: 126484615-126484739 (+)

	Features	Optimality
Stem length (bp)	35	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	4	
Bulged bases in the stem	2	
Apical loop size (nt)	9	Suboptimal
3' basal segment length (nt)	26	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.0 (476 / 476)	0.0 (476 / 476)
Cleavage homogeneity	0.0 (476 / 476)	0.0 (476 / 476)

Rank among tested pri-miRs

