

pri-mir-30e

chr1: 40754343-40754467 (+)

	Features	Optimality
Stem length (bp)	34	Suboptimal
Lower stem length (bp)	12	Suboptimal
Mismatched pairs in the stem	1	Optimal
Bulged bases in the stem	4	
Apical loop size (nt)	16	Optimal
3' basal segment length (nt)	11	Optimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3
Cleavage efficiency	0.71 (295 / 476)	1.78 (174 / 476)
Cleavage homogeneity	0.93 (123 / 476)	0.98 (168 / 476)

Rank among
tested pri-miRs



Top

Bottom

