

pri-mir-3690-1

chrX: 1293897-1294021 (+)

	Features	Optimality
Stem length (bp)	32	Suboptimal
Lower stem length (bp)	9	Suboptimal
Mismatched pairs in the stem	7	
Bulged bases in the stem	2	
Apical loop size (nt)	9	Suboptimal
3' basal segment length (nt)	4	Suboptimal
Mismatch at -6	Not possessed	
bGWG at -10/-11	Not possessed	

	w/o SRSF3	w/ SRSF3	Rank among tested pri-miRs
Cleavage efficiency	0.35 (374 / 476)	0.39 (431 / 476)	<div><div></div>Top</div> <div><div></div>Bottom</div>
Cleavage homogeneity	0.49 (312 / 476)	0.35 (421 / 476)	

