Figure 3

A. Bar graph showing RPM of mRNA-seq and gPAR-CLIP across 5' UTR, CDS, and 3' UTR.

B. Line graph showing the proportion of Ts across CLS bins.

C. Bar graph showing the number of 3' UTR and 5' UTR blocks with conservation.

D. Graph showing RPM across ChrII position (kb) for ATG8 3' UTR.

E. Line graph showing the mean phastCons score across CLS bins for 5' UTR, CDS, and 3' UTR controls.