



Figure S11. Correlation of PAS with transcript level (FPKM). **A)** The box plot depicts FPKM distribution of transcripts having top 12 polyadenylation signals. Transcripts with one annotated 3'UTR is considered for analysis. **B)** Transcripts with particular PAS, which showed distinct FPKM distribution profile are chosen and looked for cell-type specific FPKM distribution.