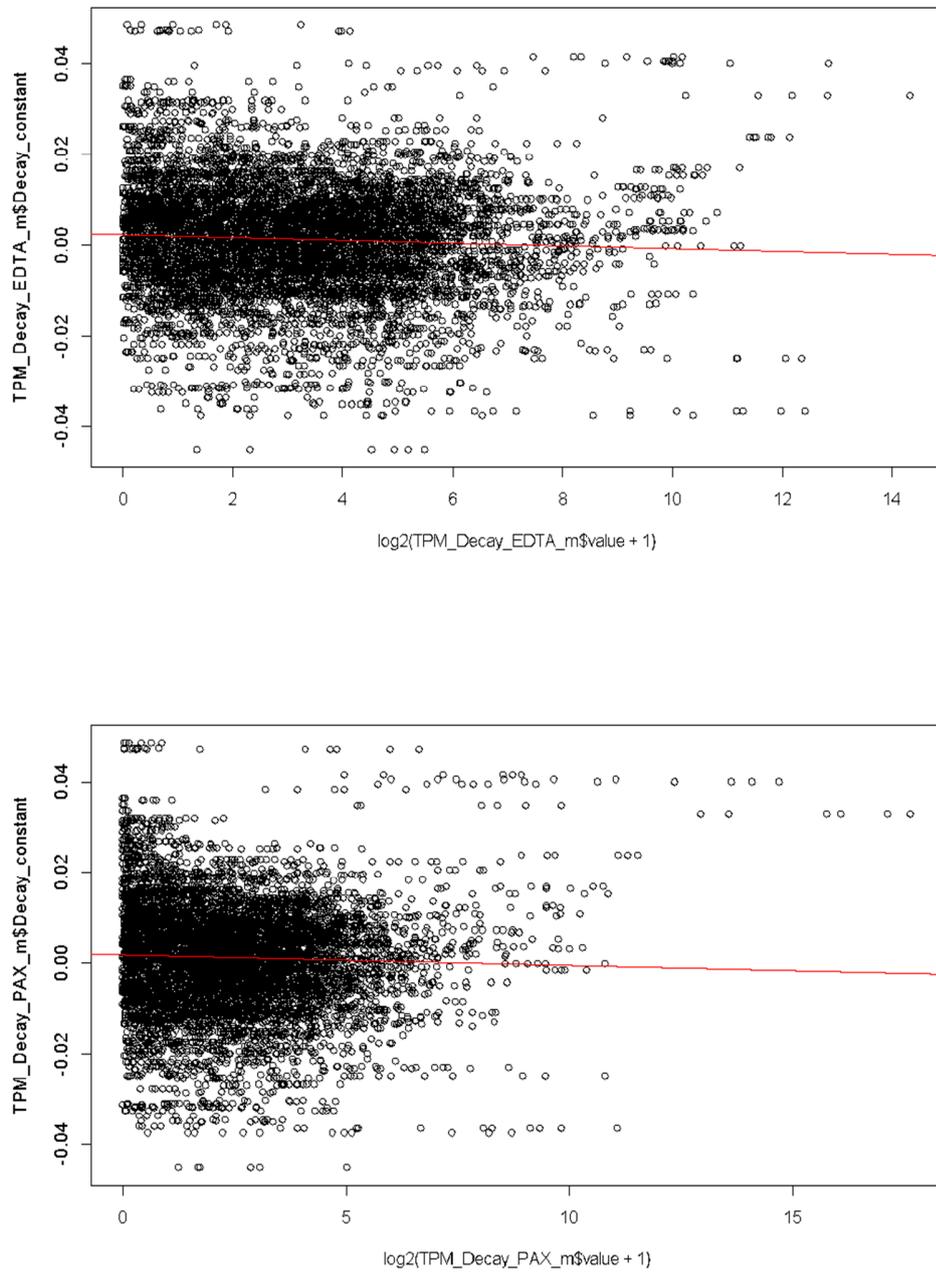
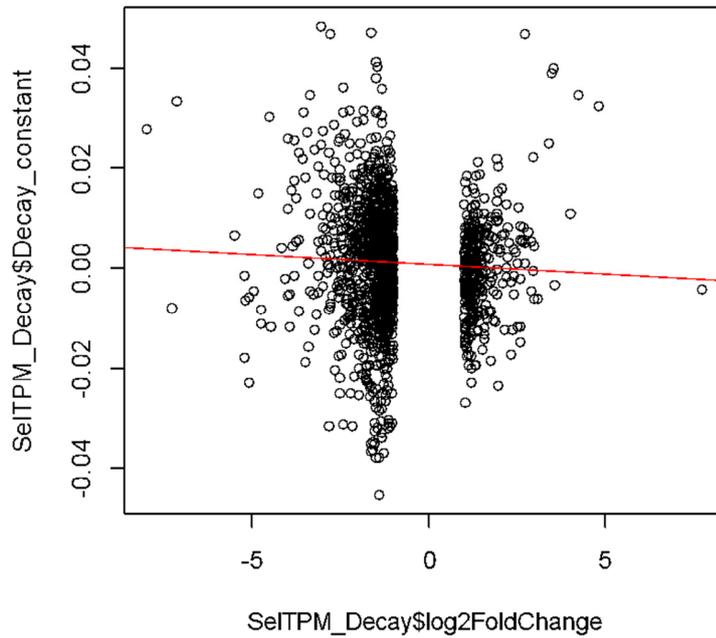


## SUPPLEMENTARY FIGURES



**Figure S2.** No appreciable correlation between expression (TPM) and decay constant exists in either EDTA or PAXgene samples.



**Figure S3. Slow degrading DEGs are not overrepresented in genes upregulated in the EDTA samples.** On the left side are genes with higher relative abundance in EDTA. The gap shows exclusion of DEGs with  $|L2FC| \leq 1$ . On the right side are the genes with higher relative abundance in PAXgene. DEGs above 0 on the Y axis are degrading faster than the ones below 0. From this representation, we can reject the hypothesis that slow decaying DEGs dominate the upregulated genes in EDTA.