

Figure 1 displays a multi-panel plot showing gene expression data across a gene list index (0 to 18113).

The top panel shows the $\log_2(\text{RPKM})$ expression level (y-axis) versus the Gene List Index (x-axis). A blue line represents the expression profile. A vertical dashed green line marks the "Zero crossing at 7260", and a vertical dashed blue line marks the "Peak at 12438".

The middle panel is a barcode plot showing the number of genes in each cluster (y-axis) versus the Gene List Index (x-axis). The plot displays a dense series of vertical black lines, indicating the presence of genes in the clusters.

The bottom panel is a stacked area chart showing the proportion of genes in two clusters: "clus2" (green area) and "NON.clus2" (black area) versus the Gene List Index (x-axis). The "clus2" cluster is dominant in the early part of the gene list, while the "NON.clus2" cluster is dominant in the later part.

The x-axis for all panels is labeled "Gene List Index" and "Number of genes: 18113 (in list), 179 (in gene set)".

[illegible]