

Figure 1 displays a multi-panel plot showing the distribution of gene lists. The x-axis represents the Gene List Index, ranging from 0 to 17,976. The y-axis represents the number of gene lists, ranging from 0 to 17,976.

The top panel shows a red line graph representing the number of gene lists versus the Gene List Index. The curve starts at 0, rises sharply to a peak at index 3015, and then gradually declines, crossing the x-axis at index 9208. A horizontal dashed line is drawn at the peak value.

The middle panel shows a series of vertical bars representing the gene lists. The bars are concentrated in the region where the red line is above the x-axis, indicating a high density of gene lists in that region.

The bottom panel shows a green area graph representing the distribution of gene lists. The area is divided into two regions: "clus4" (left) and "NON.clus4" (right). The "clus4" region is characterized by a high density of gene lists, while the "NON.clus4" region shows a much lower density.

Key annotations include:

- Peak at 3015
- Zero crossing at 9208
- "clus4"
- "NON.clus4"

Gene List Index

Number of genes: 17976 (in list), 77 (in gene set)

