

Figure 1 displays the distribution of gene lists across clusters. The top panel shows the cumulative distribution function (CDF) of the number of genes in the list, with a peak at 3432 and a zero crossing at 10017. The middle panel shows a barcode of vertical lines representing the number of genes in each cluster. The bottom panel shows the distribution of gene lists across clusters, with labels "clus4" and "NON.clus4".

