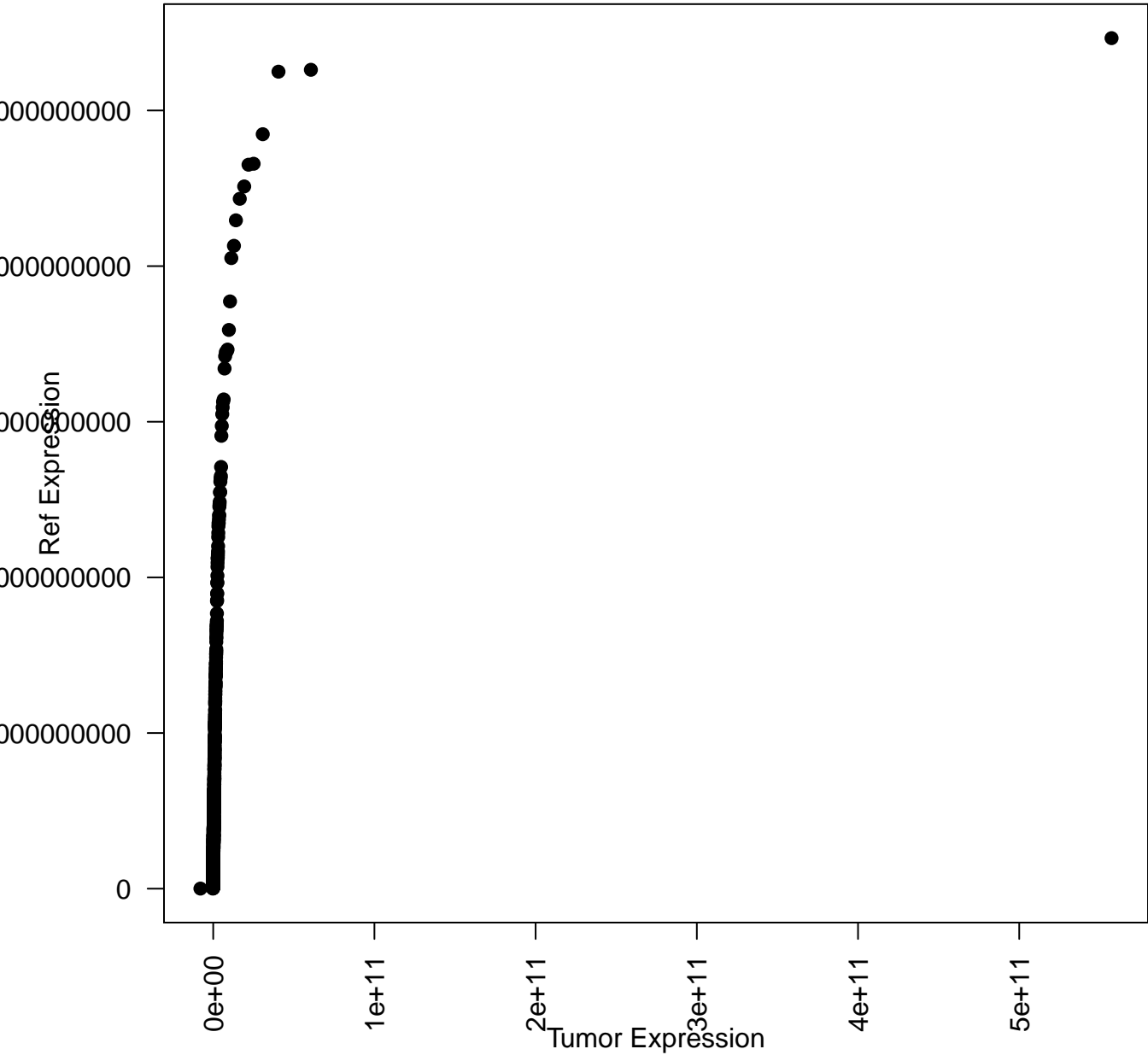
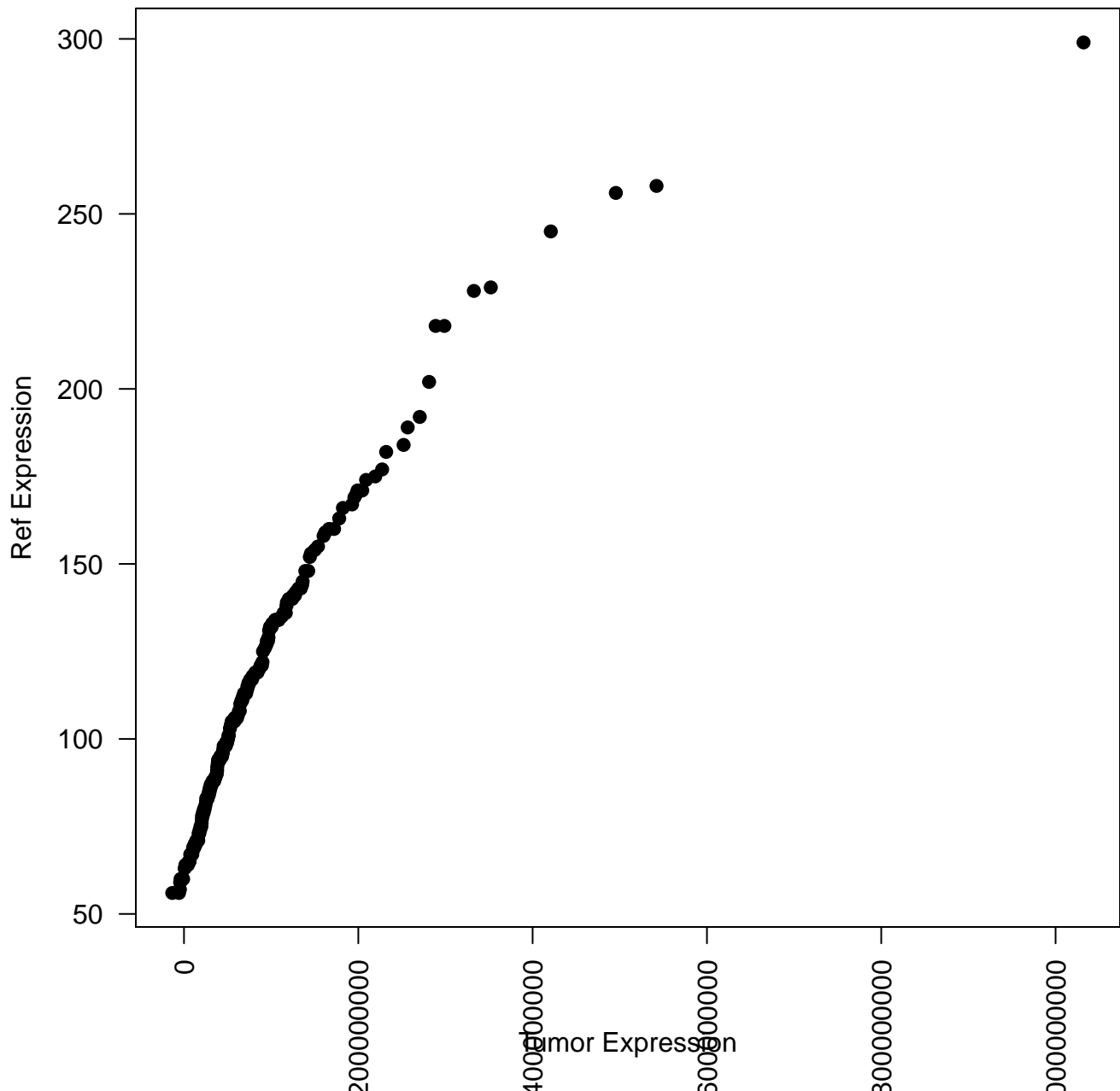
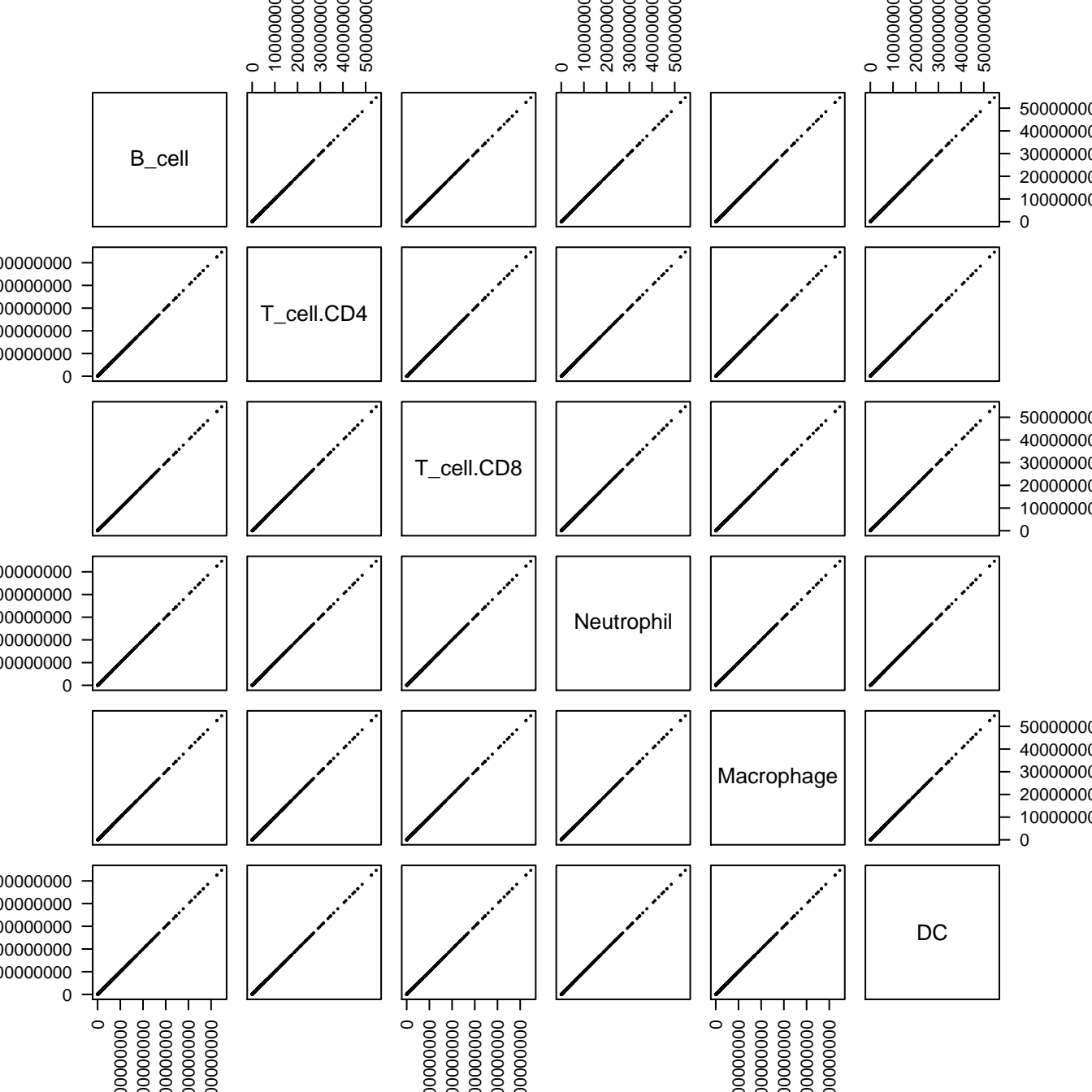


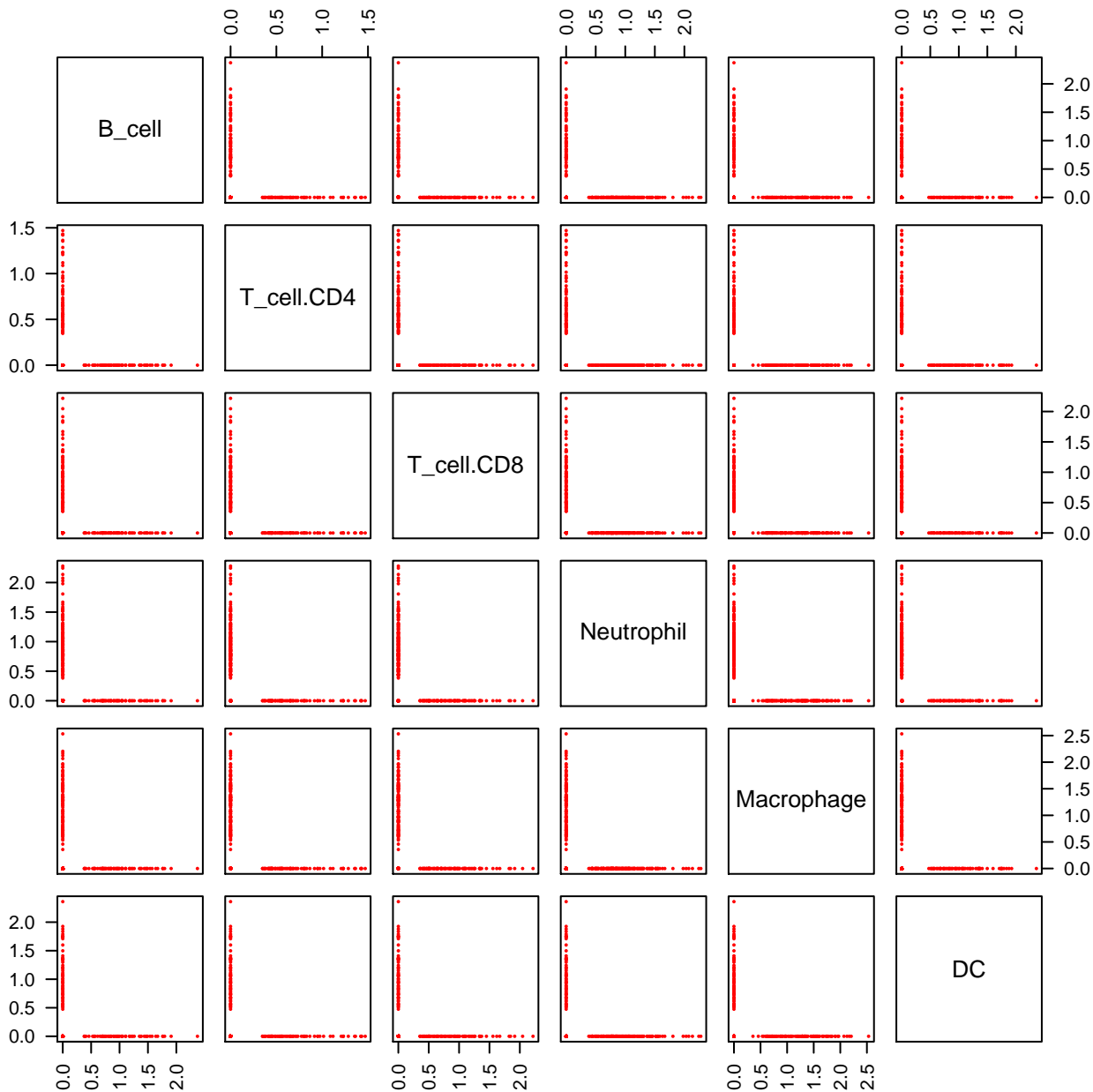
A scatter plot showing the relationship between Tumor Expression (x-axis) and an unlabeled y-axis. The x-axis is labeled 'Tumor Expression' and has major ticks at 0e+00, 1e+11, 2e+11, 3e+11, 4e+11, and 5e+11. The y-axis is unlabeled. The data points are black dots. There is a dense cluster of points at low tumor expression values (near 0e+00) with a wide range of y-values. As tumor expression increases, the y-values generally increase, forming a curve that rises sharply and then levels off. There are two distinct points at approximately 1e+11 on the x-axis with high y-values. A single outlier point is located at approximately 5.5e+11 on the x-axis with a very high y-value.

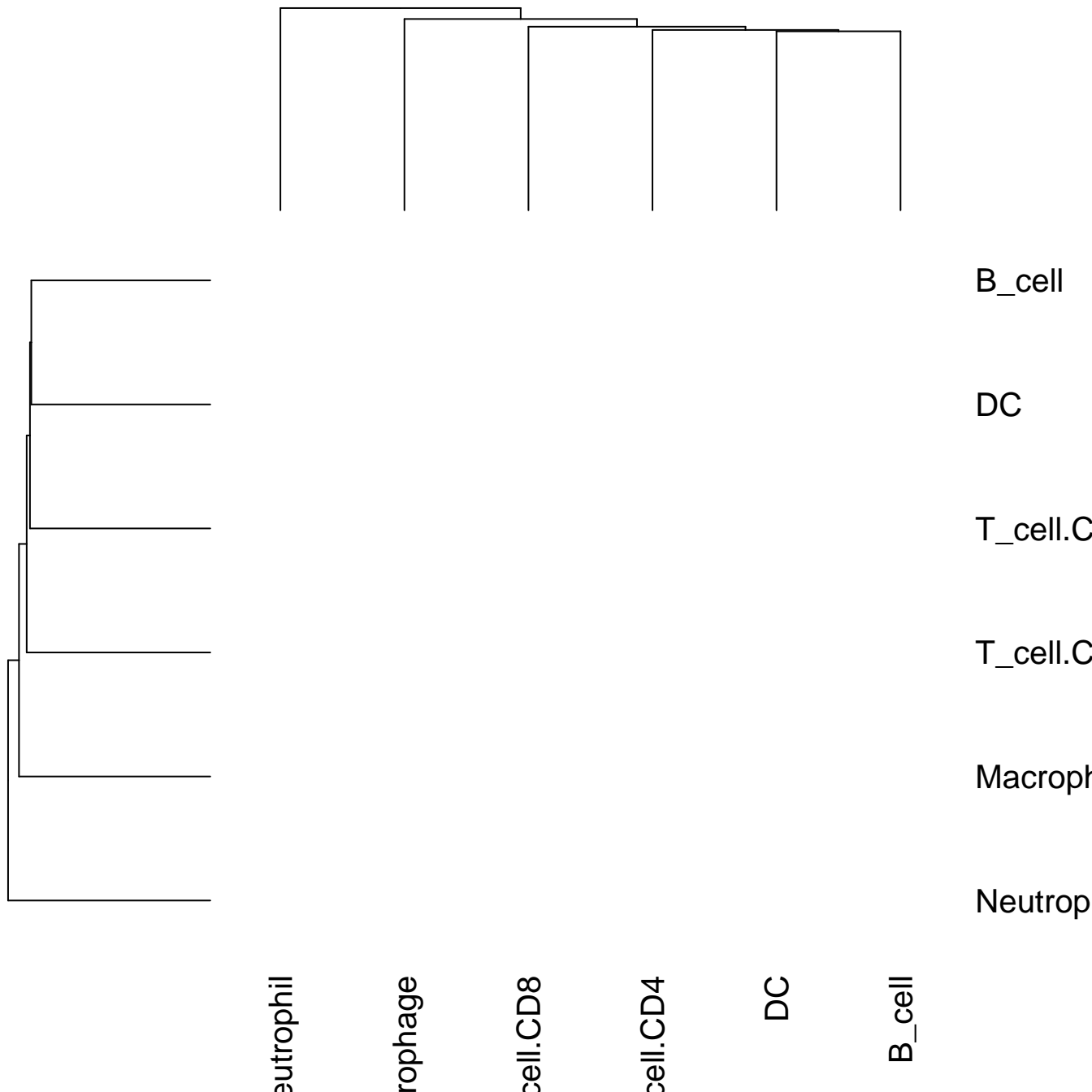


Gene-Gene Q-Q plot









Histogram of $\text{cor}(\text{Fmat})[\text{upper.tri}(\text{cor}(\text{Fmat}))]$

