**R packages to install**

**CRAN packages:**

RColorBrewer

gplots

ggplot2

calibrate

pheatmap

Command to install: Please run the following command in R terminal

install.packages("package name", dependencies = TRUE)

**Bioconductor packages:**

genefilter

DESeq2

edgeR

Command to install:

if (!requireNamespace("BiocManager", quietly = TRUE))

 install.packages("BiocManager")

BiocManager::install("package name")

The above commands for both CRAN and Bioconductor packages will install the main as well as the dependent packages.