

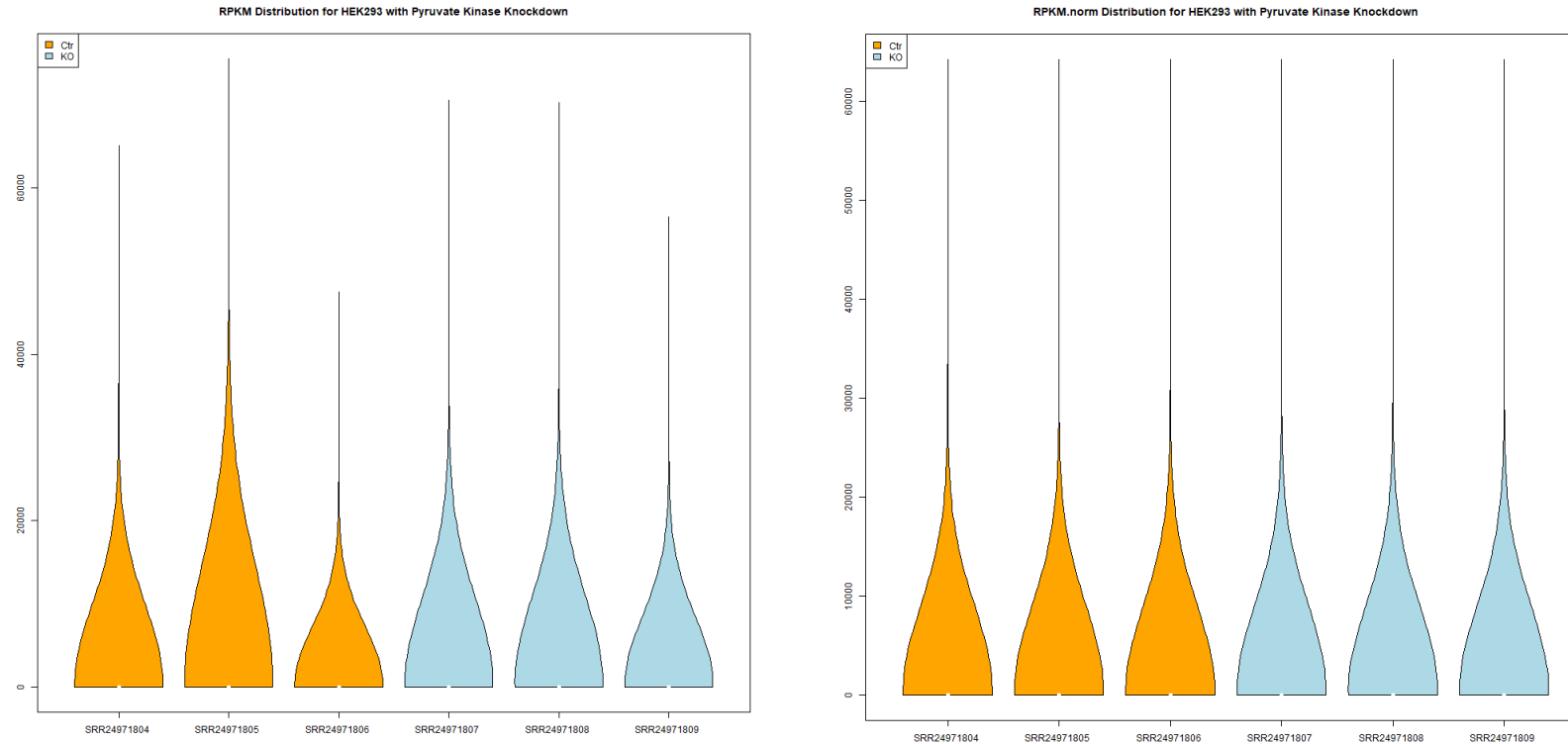
# RNA-seq data analysis for HEK293 cells with knockdown of pyruvate kinase

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Jianxun Wang

08/2023

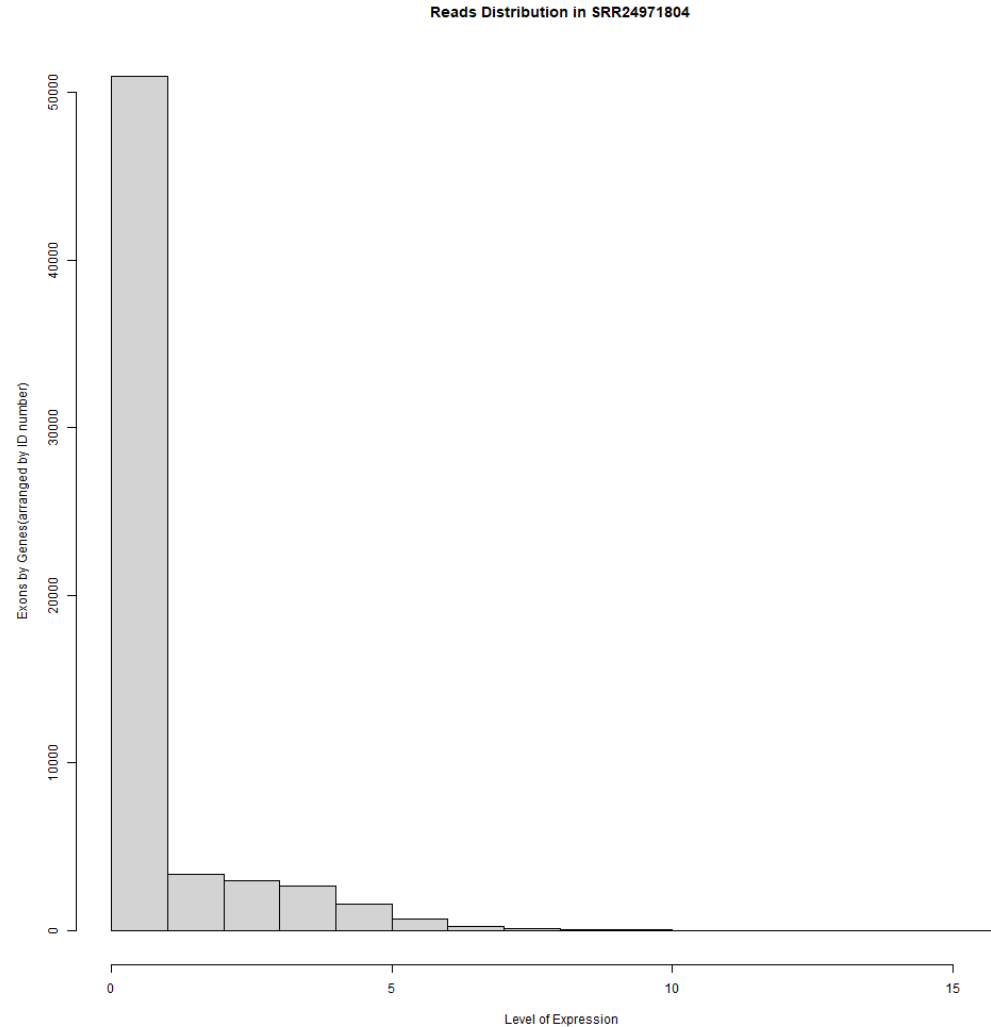
# Violin plot of RPKM showing RNA Seq reads is not in normal distribution



The T test is not fitted for this RNA Seq data

# Histogram confirms that the reads distribution is not normally distributed

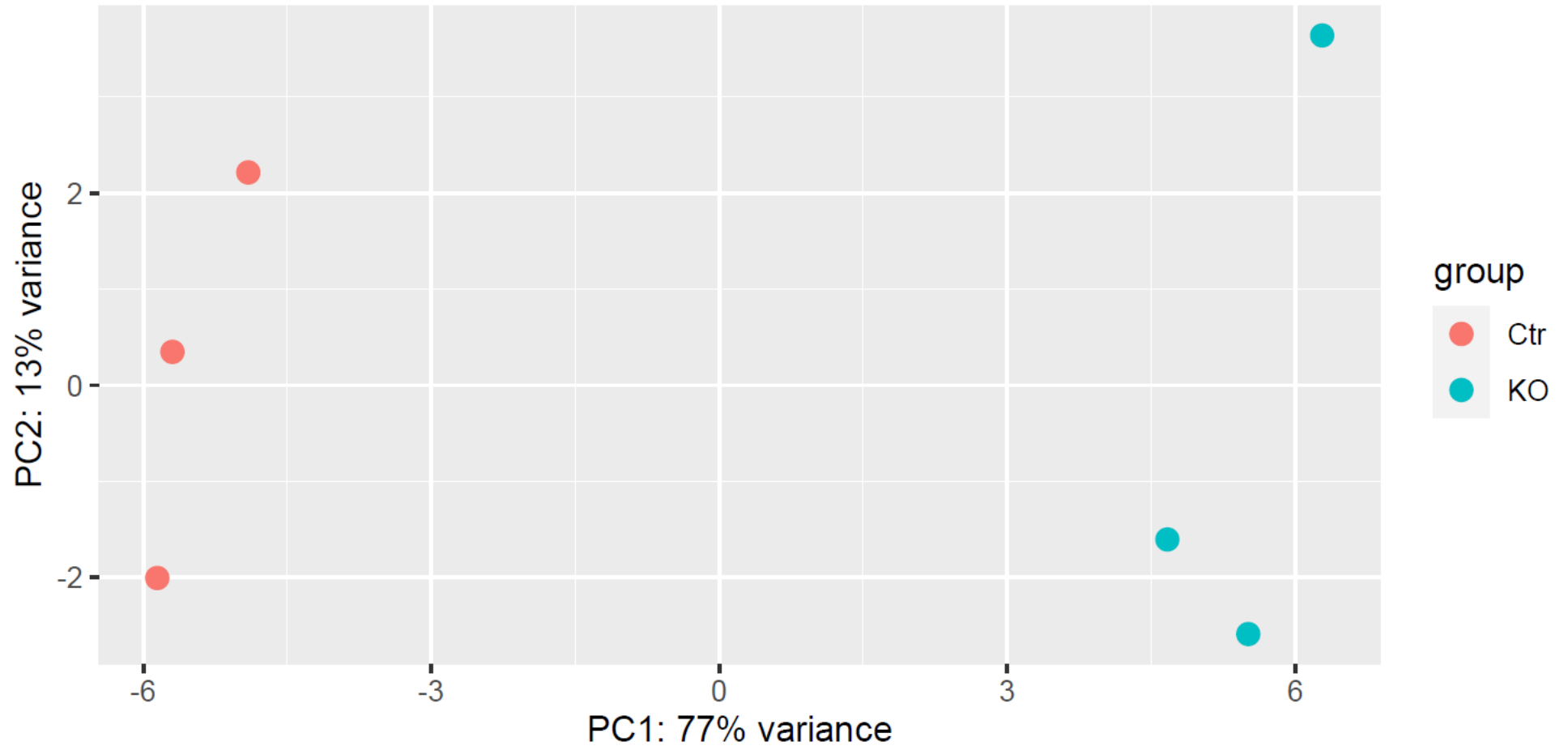
The T test is not fitted for this RNA Seq data



# PCA showing genes are significantly differentially expressed between control and KO groups

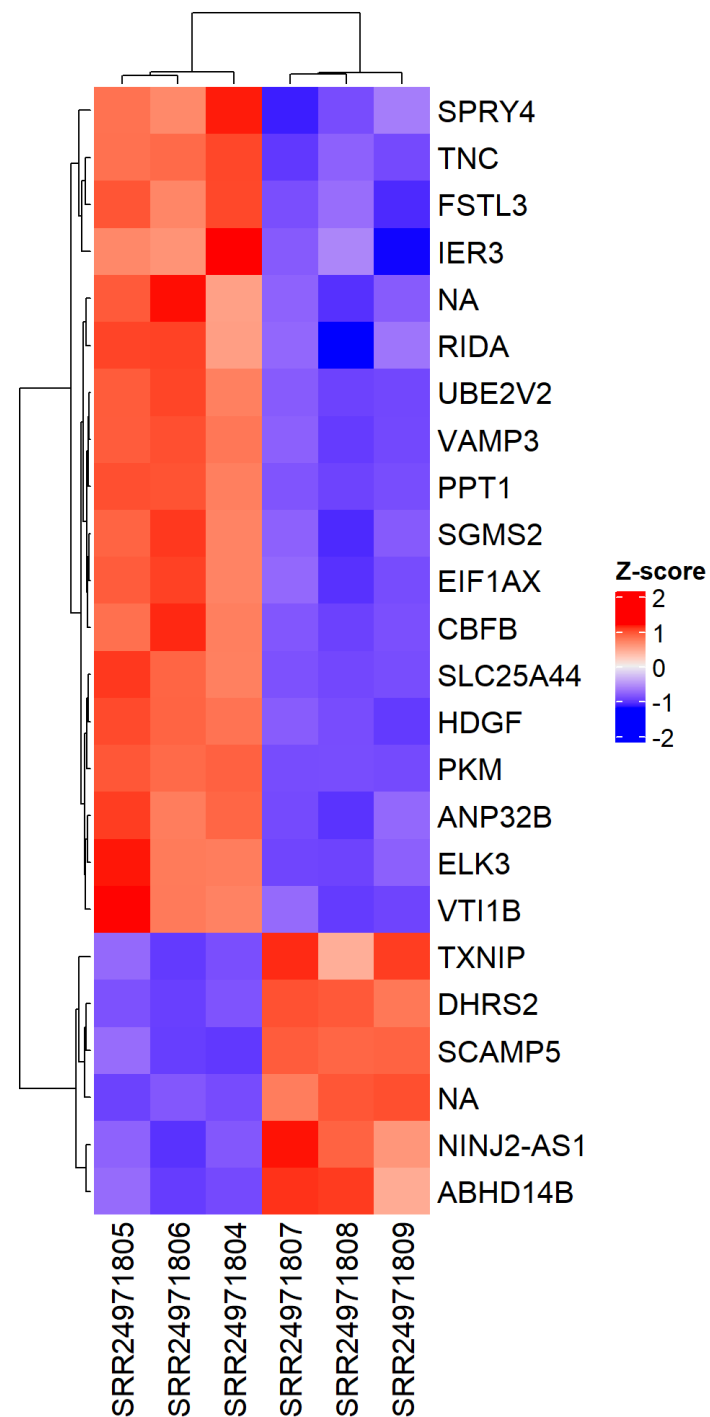
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DESeq package is used to analyze this RNA Seq data, in which the data with negative binomial distribution statistically assayed by Wald test



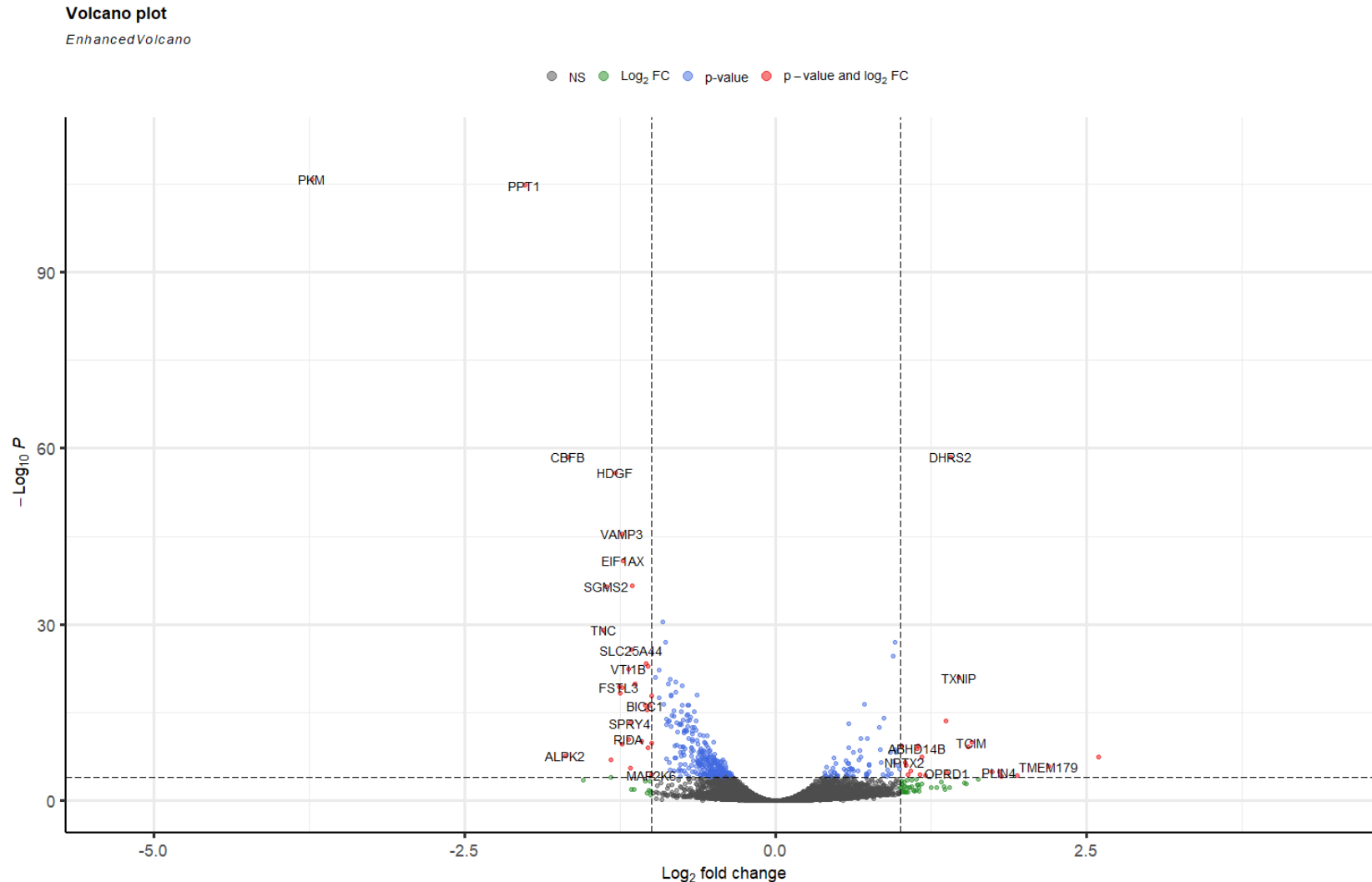
# Heatmap for genes up and down regulated

DESeq package is used to analyze this RNA Seq data, in which the data with negative binomial distribution statistically assayed by Wald test

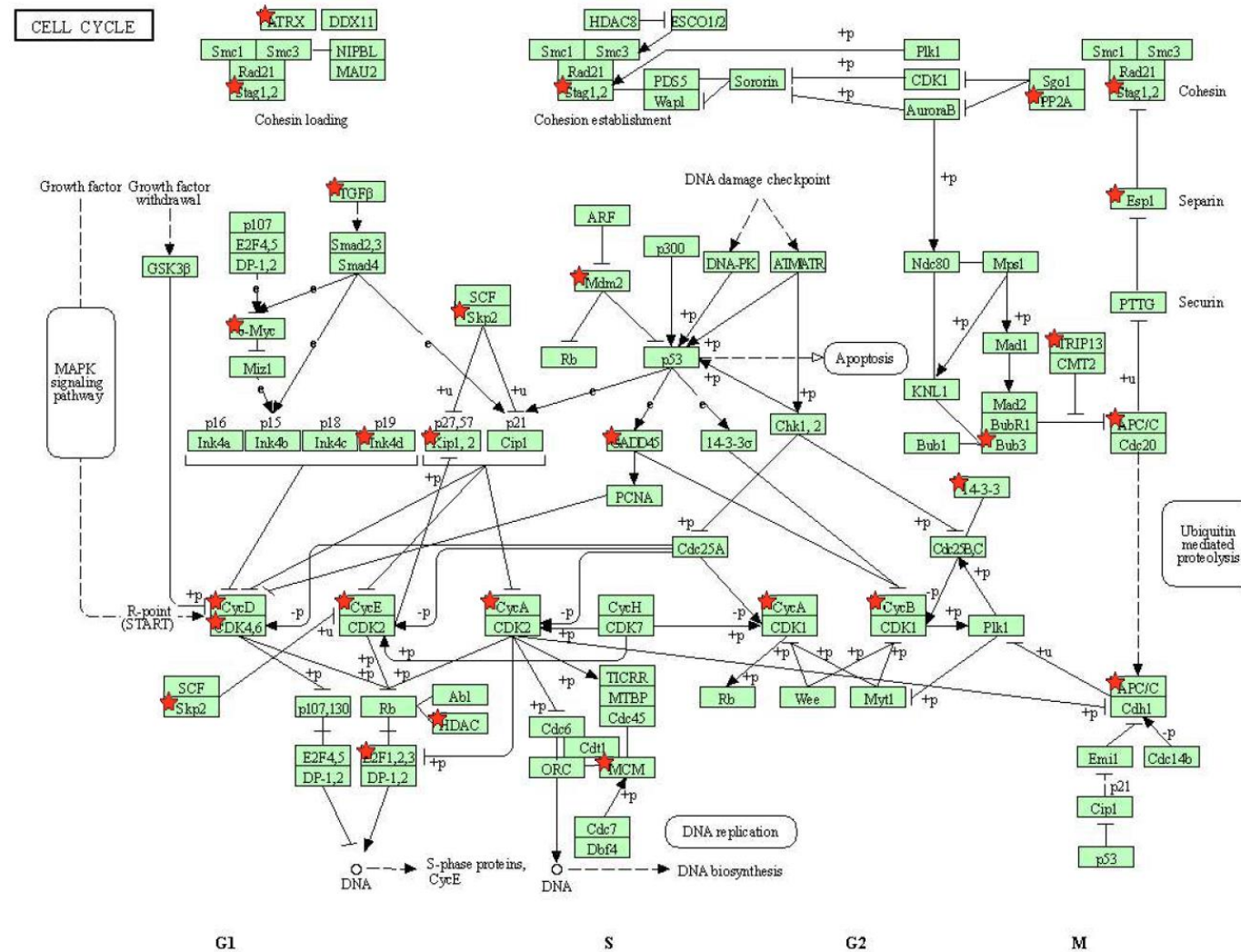


# Volcano plot for genes up and down regulated

DESeq package is used to analyze this RNA Seq data, in which the data with negative binomial distribution statistically assayed by Wald test



# The up and down regulated genes are involved in cell cycle pathways



The up and down regulated genes are involved in cell senescence

