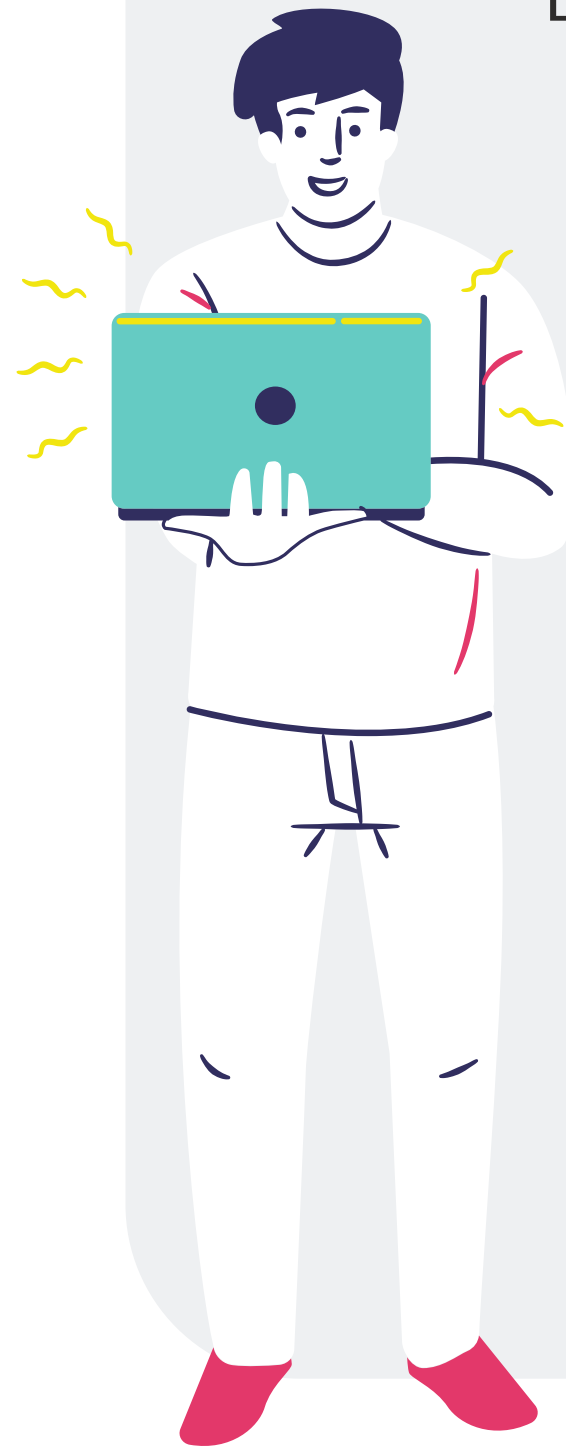


ANALYSIS OF PRAD HUB GENES

Leonardo Lavagna
Group 14



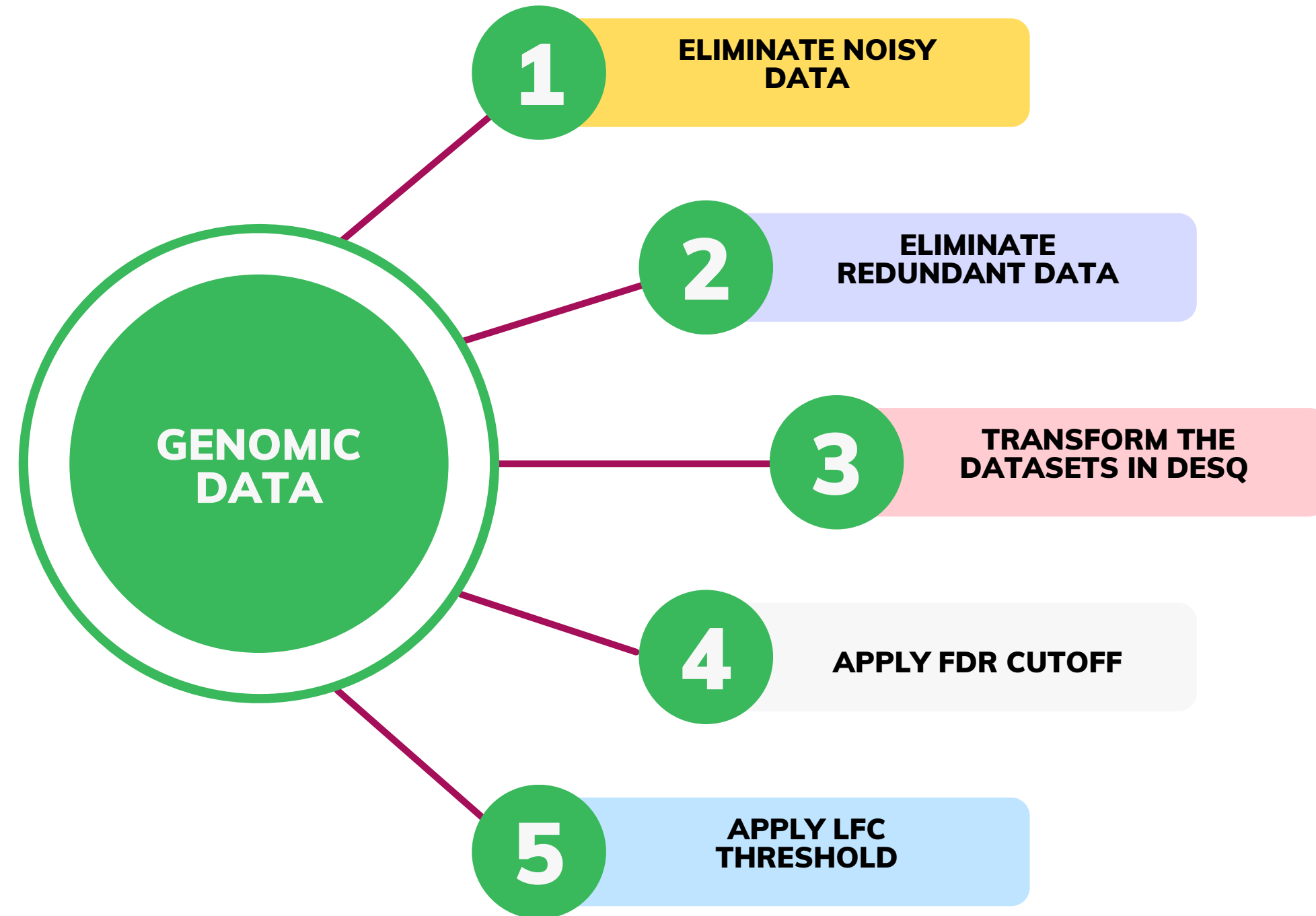
INTRODUCTION

Prostate Adenocarcinoma (**PRAD**) is a **common cancer affecting men** in late adulthood and it is **often treatable**. As a result of early diagnosis, the **mortality rate** of PRAD **fall**, although the **incidence** of PRAD continues to **rise**. It is very important to help develop new therapies and early screening tests.

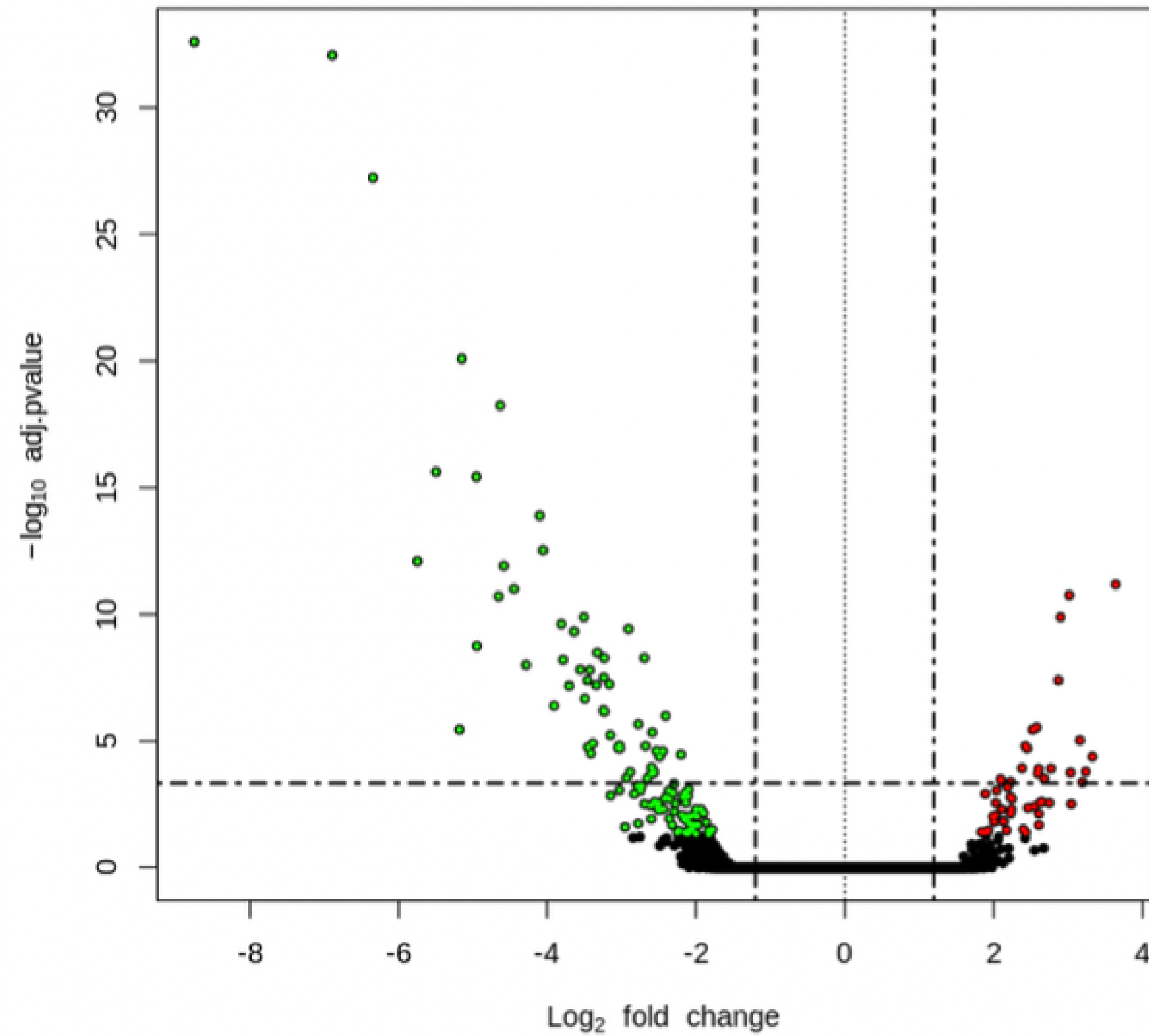
OUR GENOMIC DATA

From the **Cancer Genome Atlas** we retrieved a collection of **genomic data of about 18000 genes and 500 patients**. For our analysis we handled **two datasets** extracted from TCGA: the **tumor tissue** and the **normal tissue**.

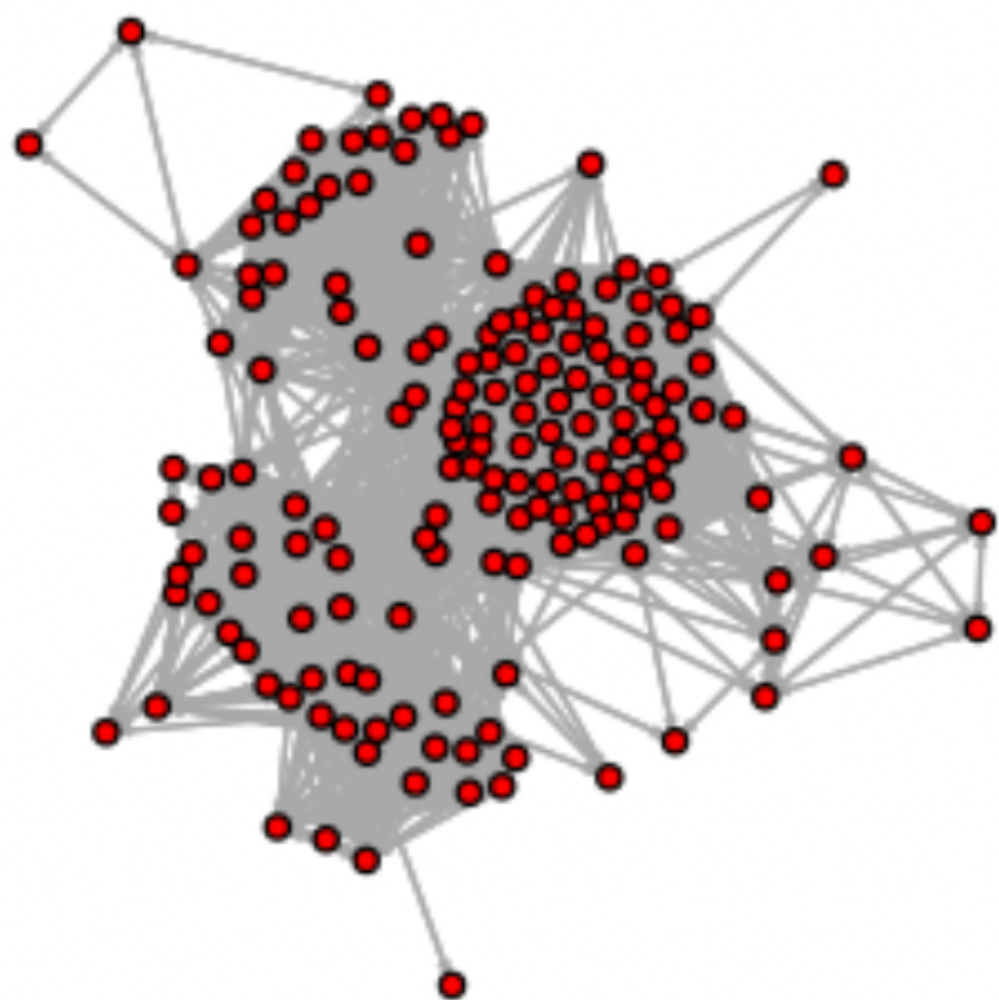
DATA CLEANING AND FEATURE ENGINEERING



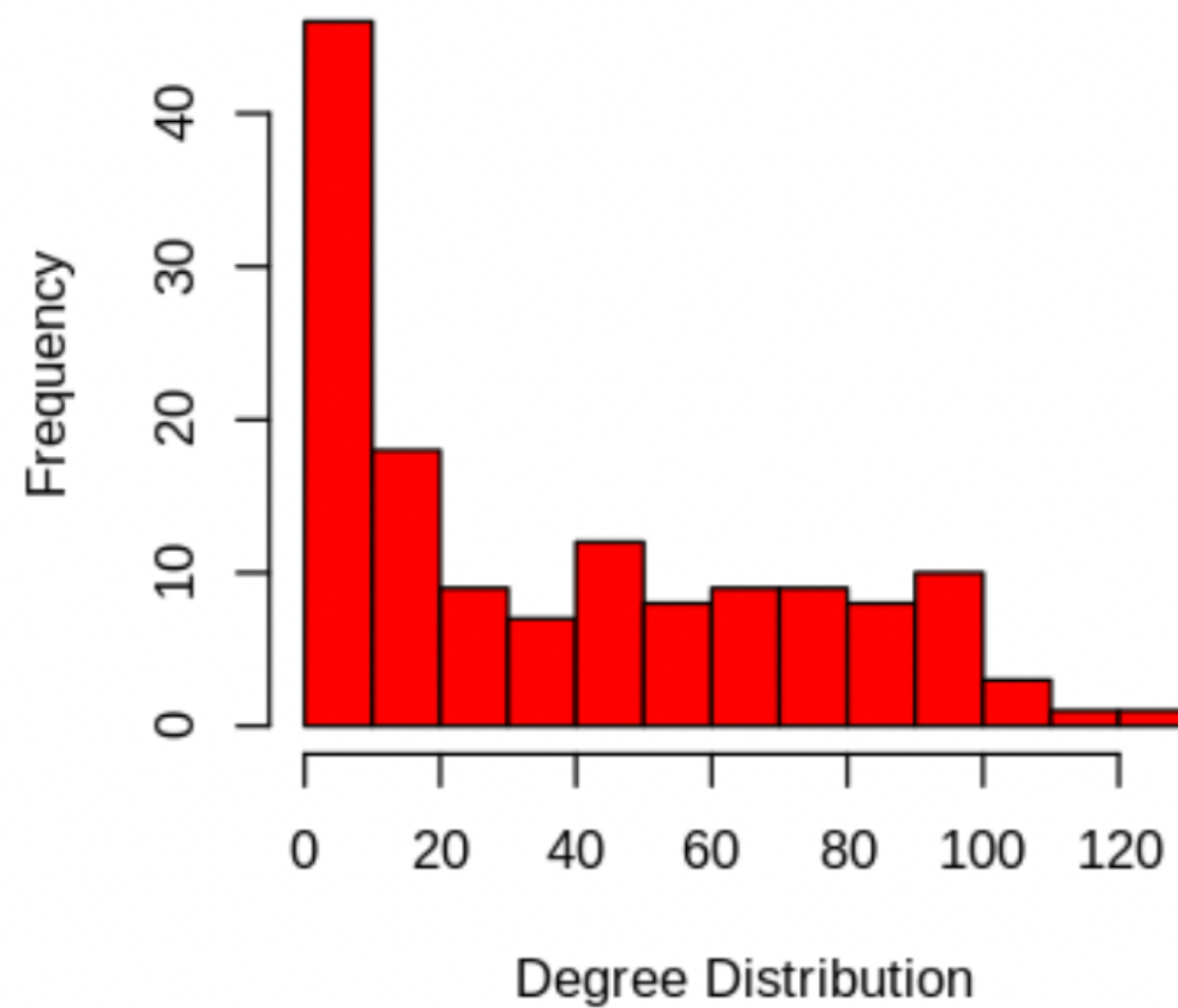
Volcano plot



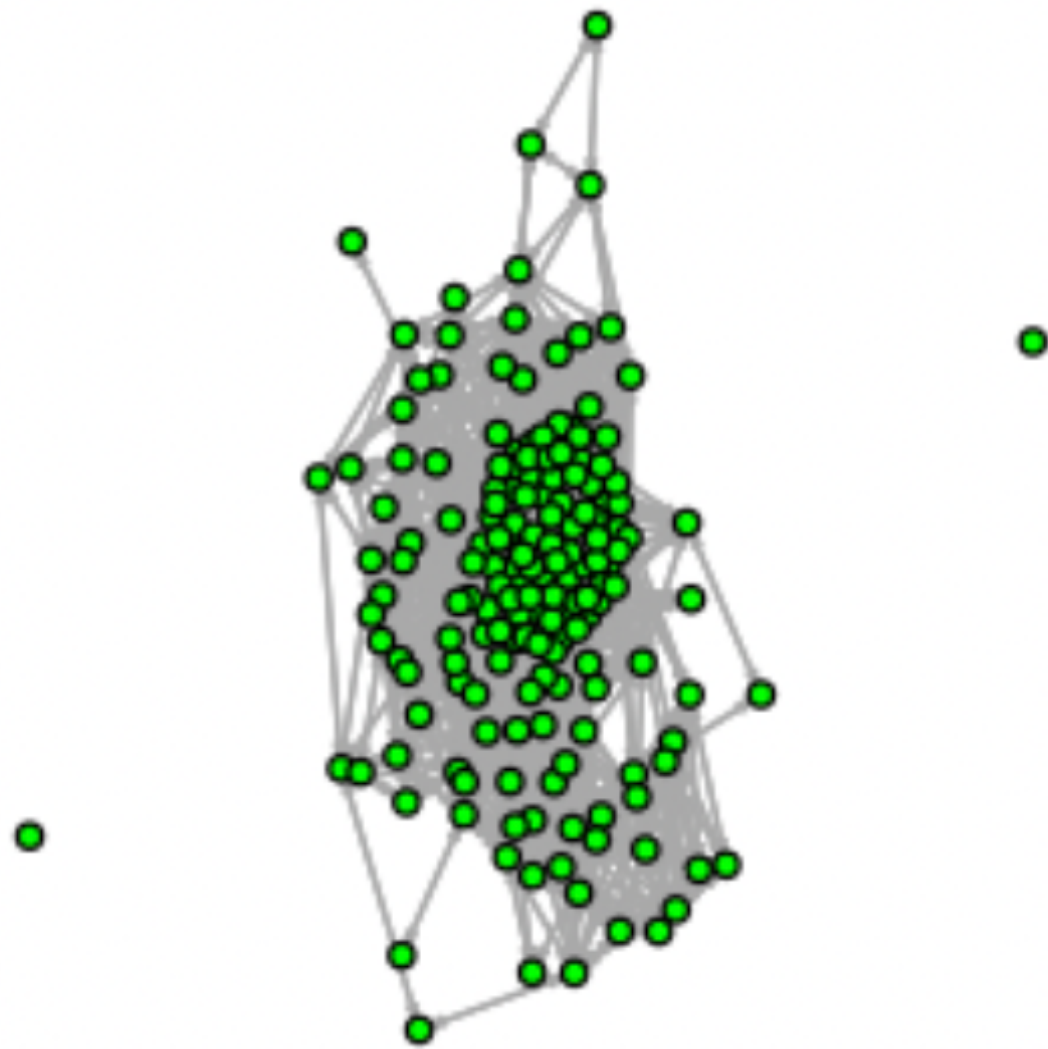
Co-expression network TUMOR



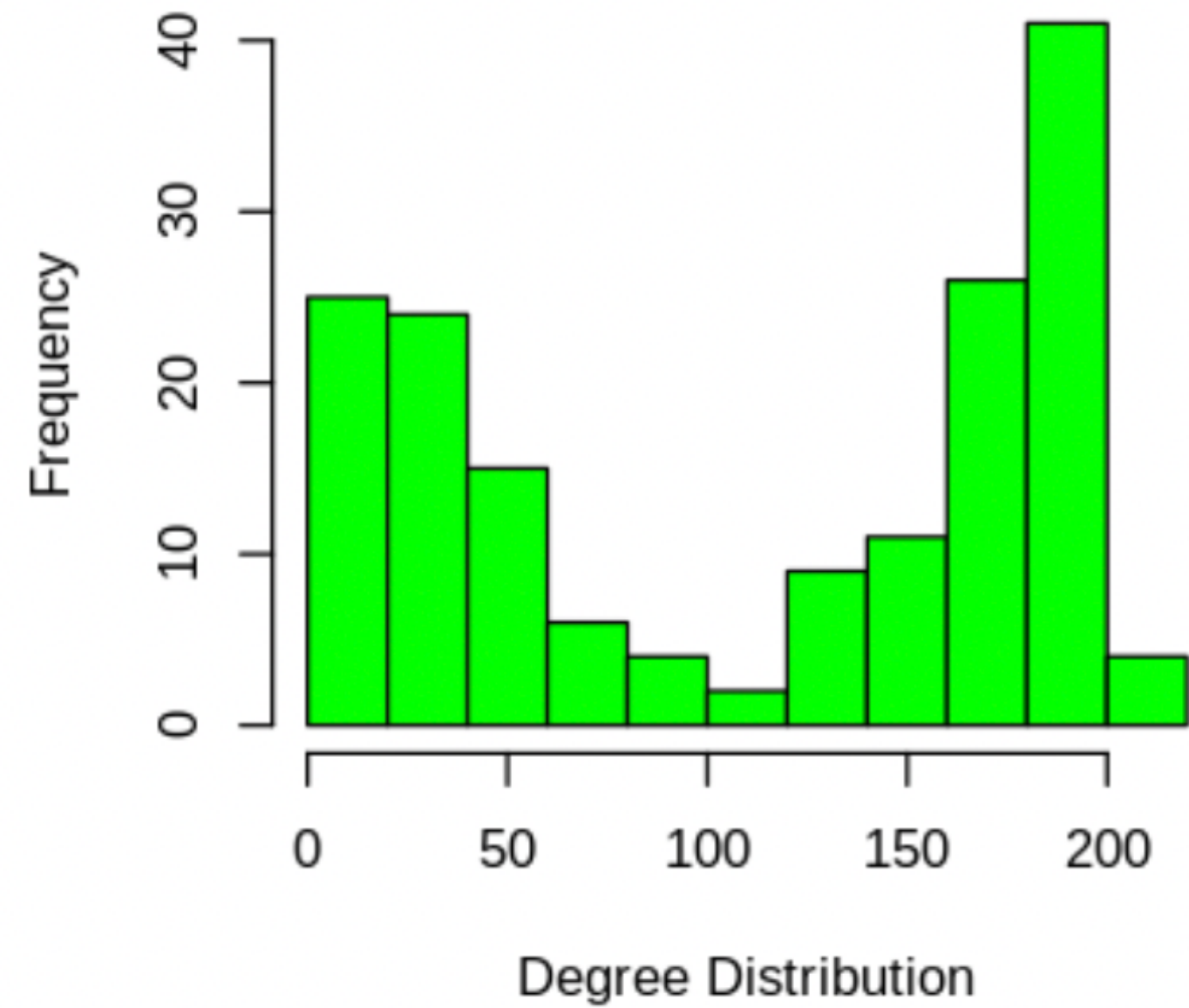
pearson Corr. TUMOR



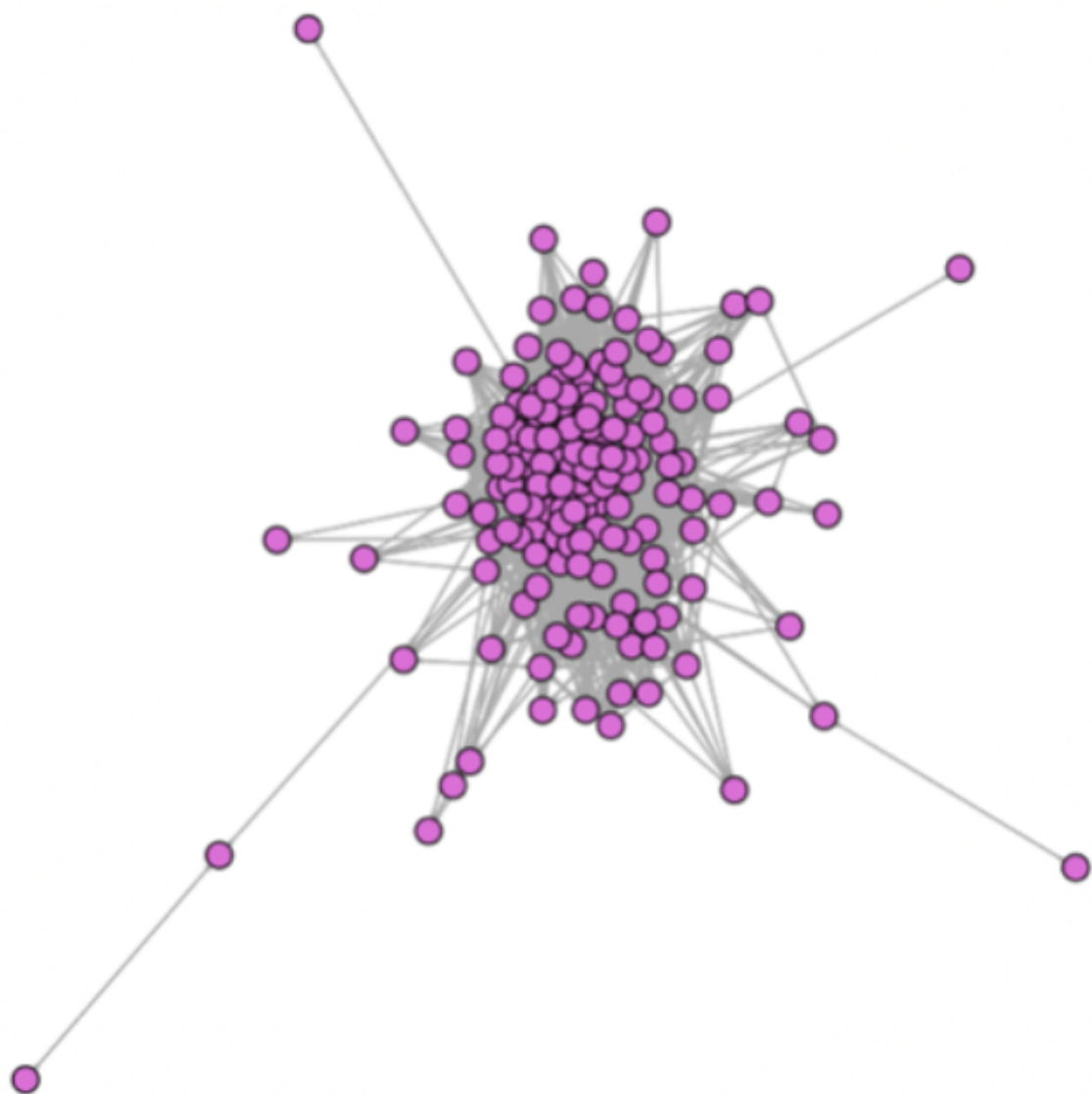
Co-expression network NORMAL



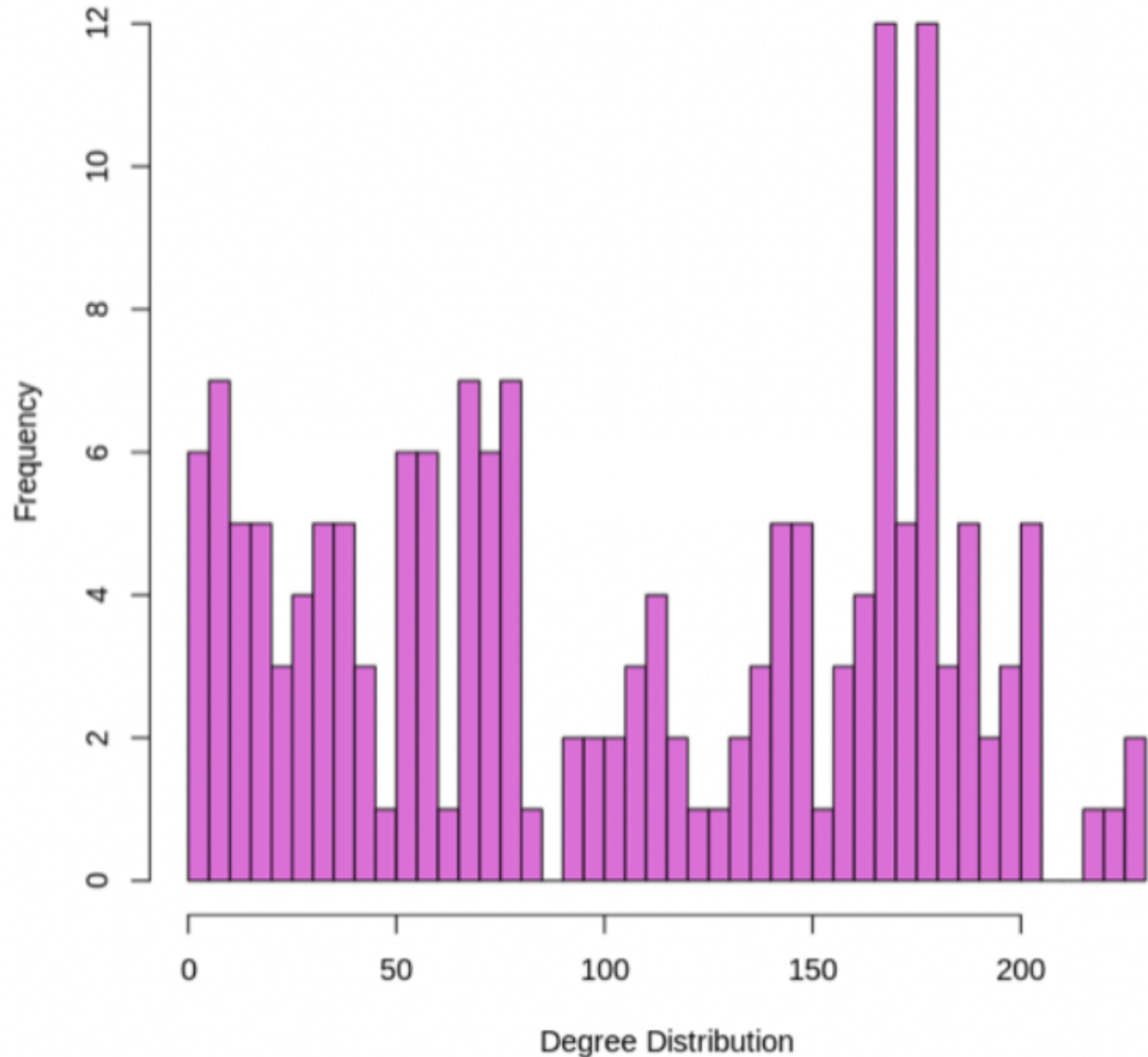
pearson Corr. NORMAL



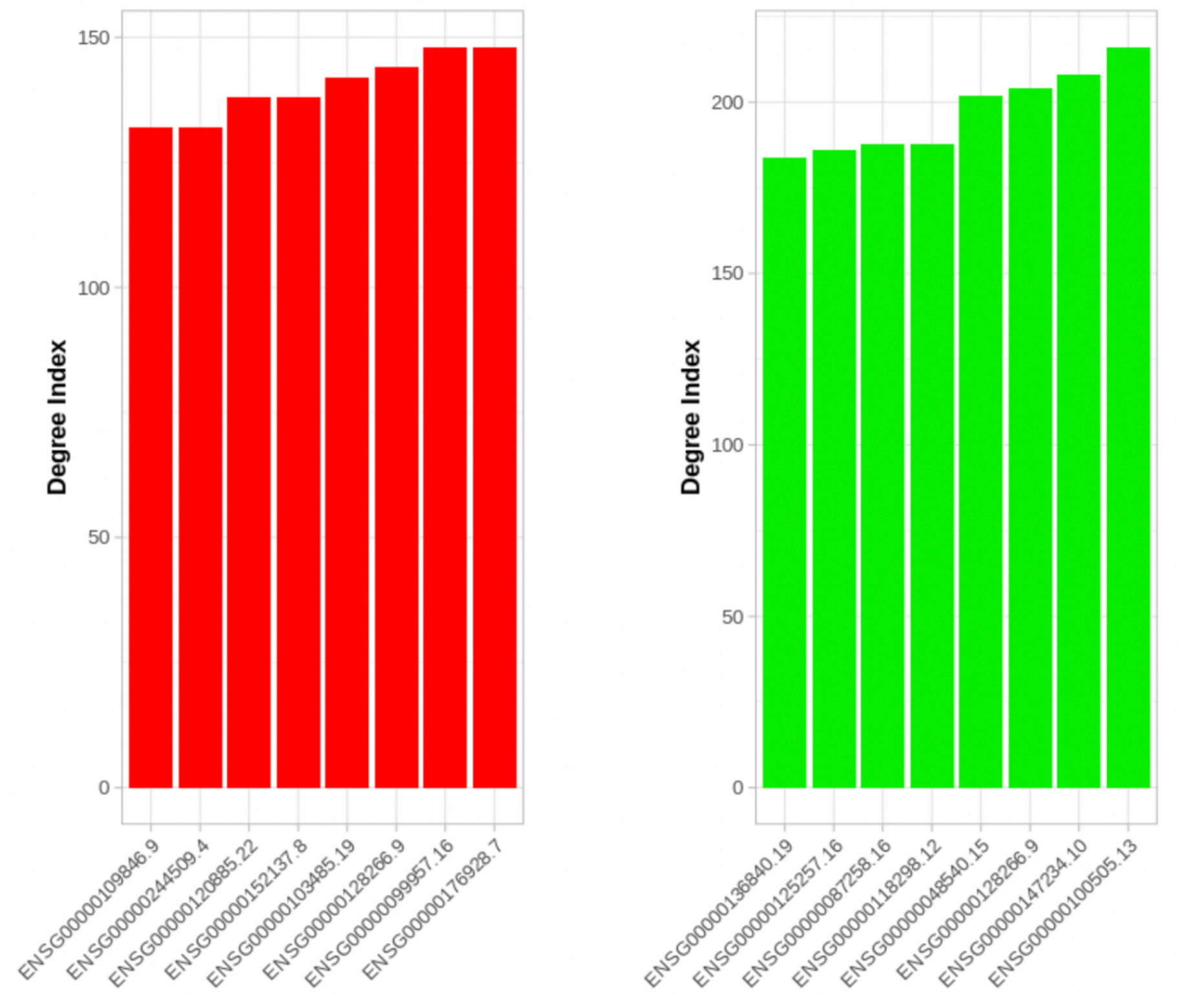
Differential Co-expression network in TUMOR vs NORMAL



Degree distribution in the Differential Co-expression

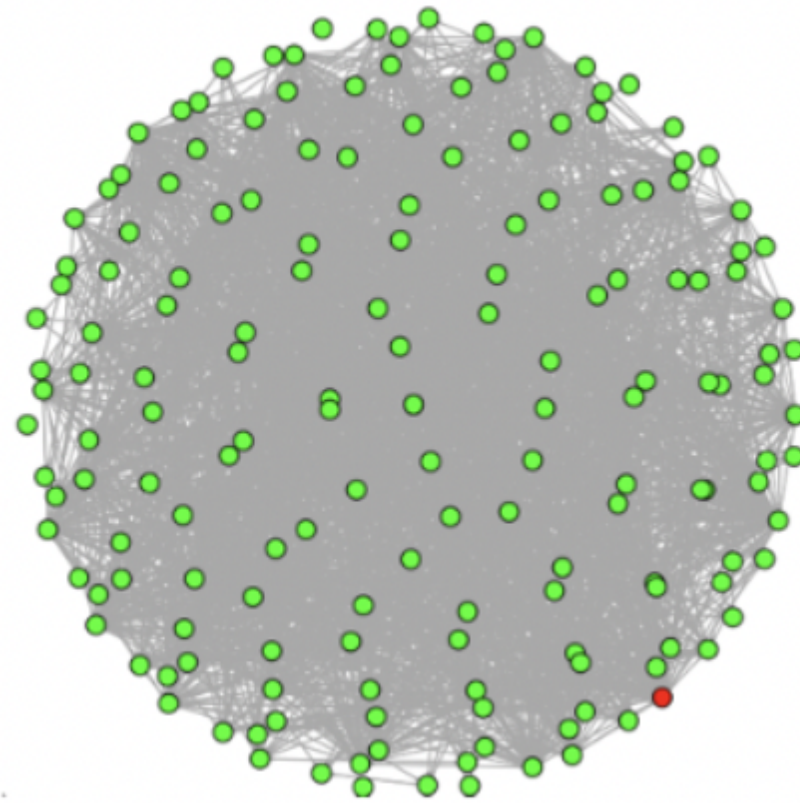


HUB GENES

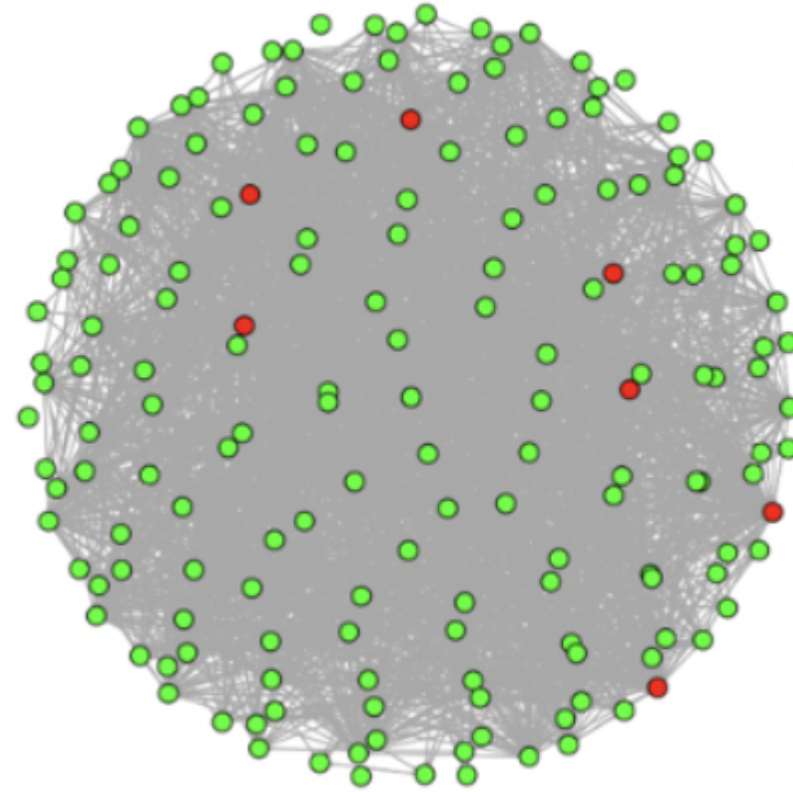


HUB GENES WITH DEGREE-BASED MEASURES

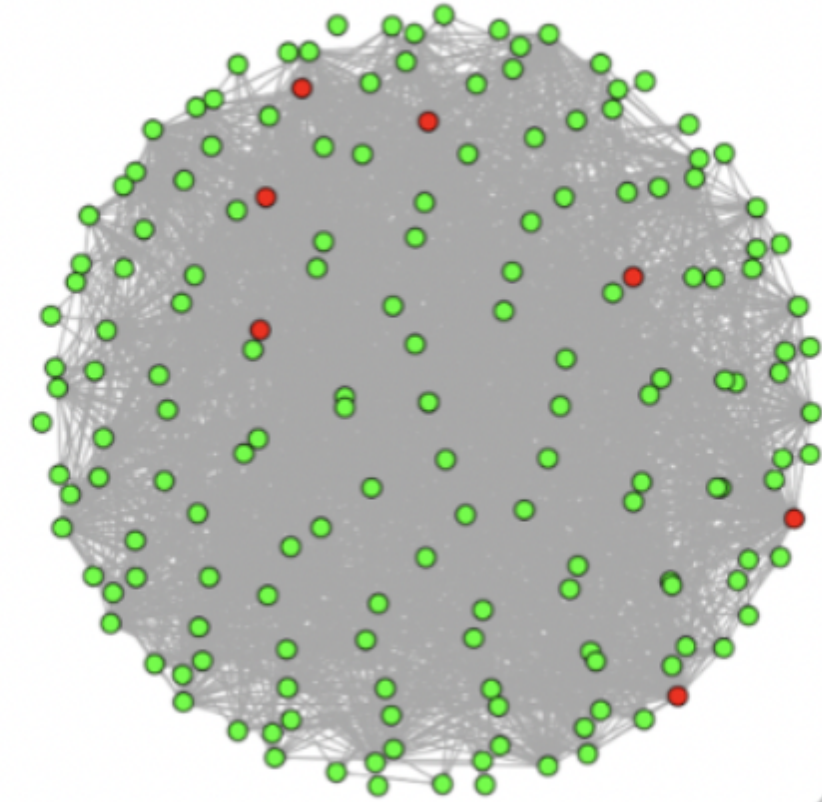
betweenness = 1 hubs in common



closeness = 7 hubs in common

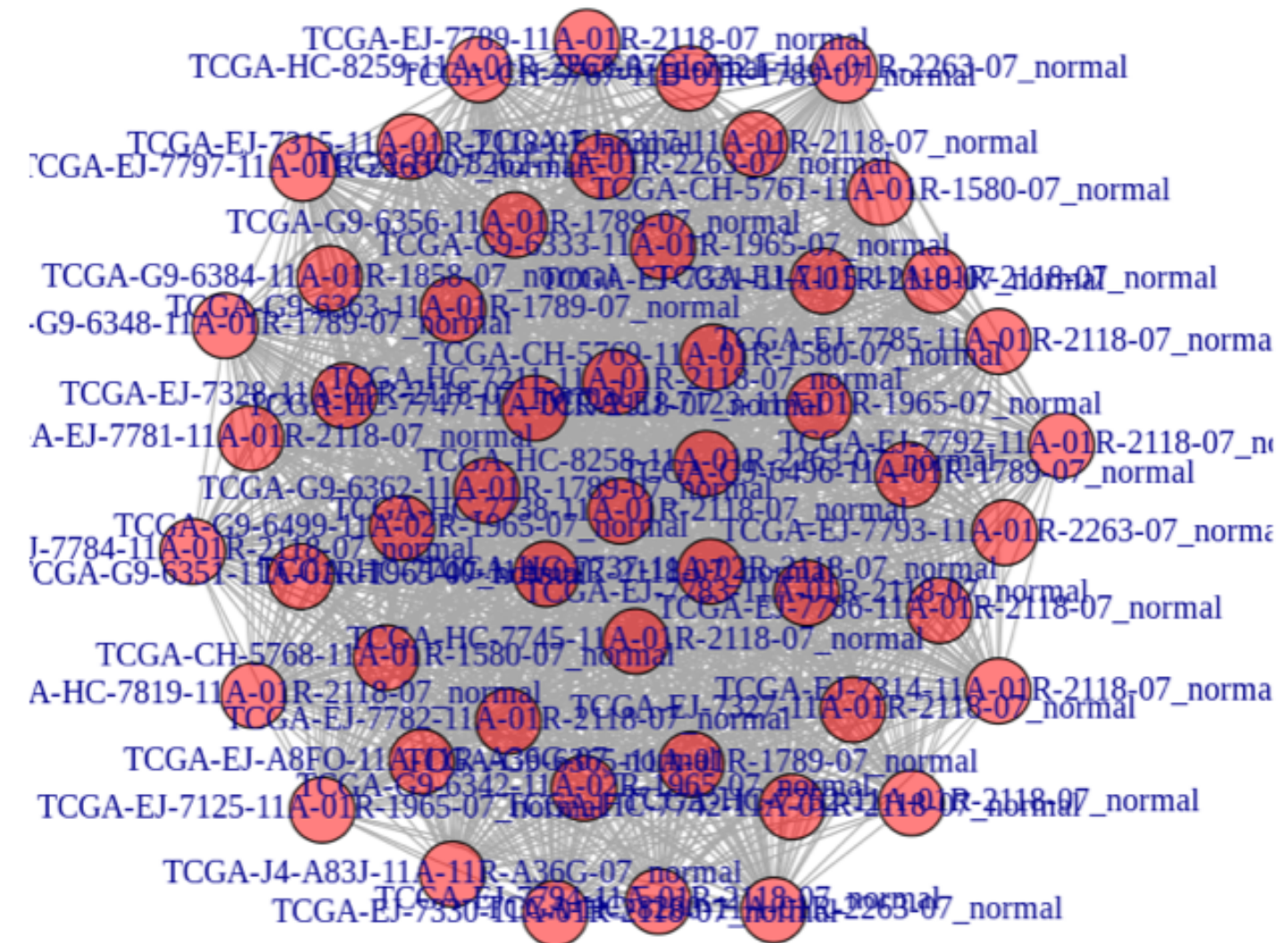
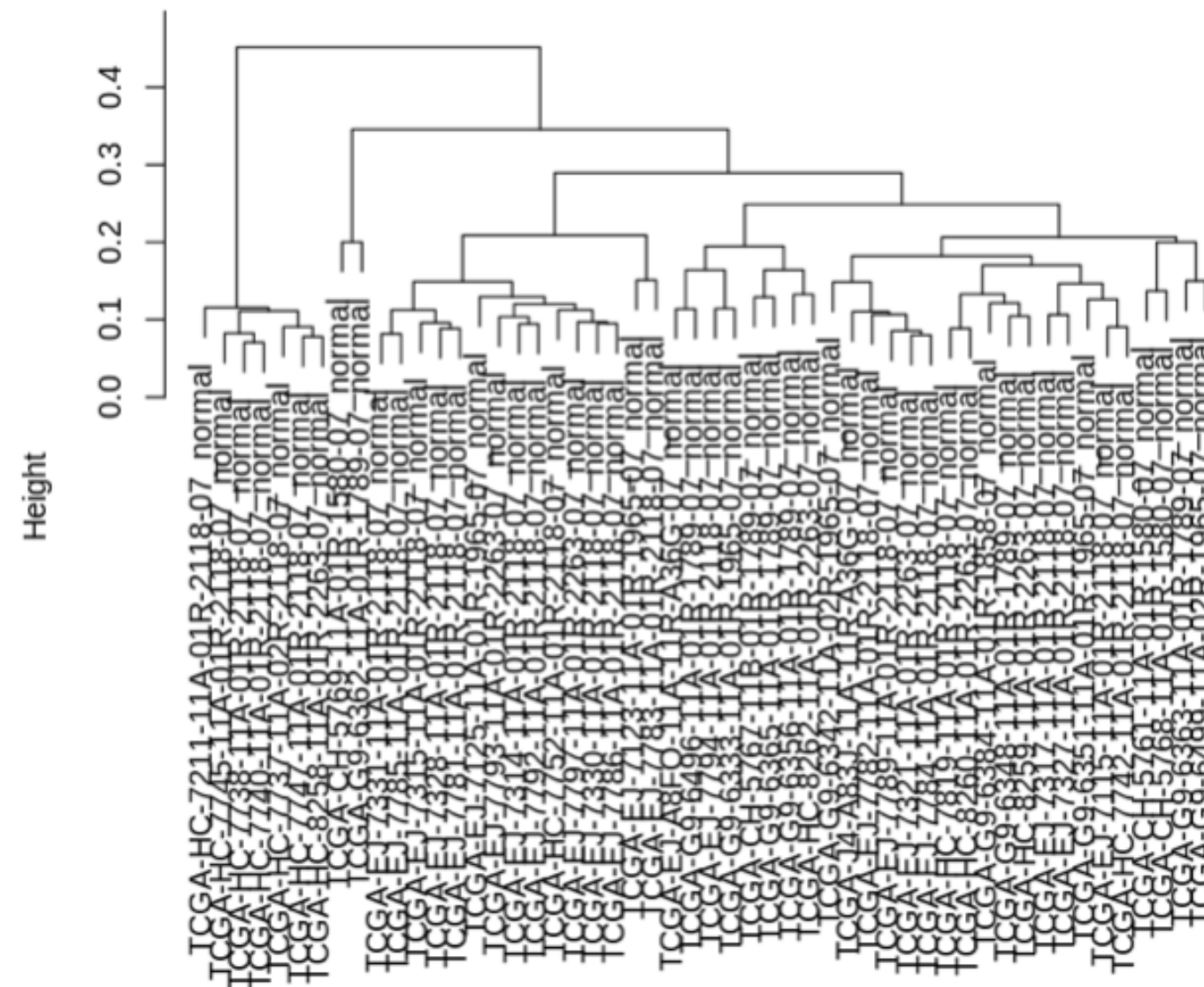


eigen = 7 hubs in common



PATIENT SIMILARITY NETWORK

Cluster Dendrogram



CONCLUSION

We identified **two hubs of genes** that have also been **found in the specialized literature**. Those are **ENSG00000120885.22** and **ENSG00000152137.8**. We used also **different similarity measures and correlations**. In every case we've obtained **similar results** and those hub genes were present.

THANK YOU!

