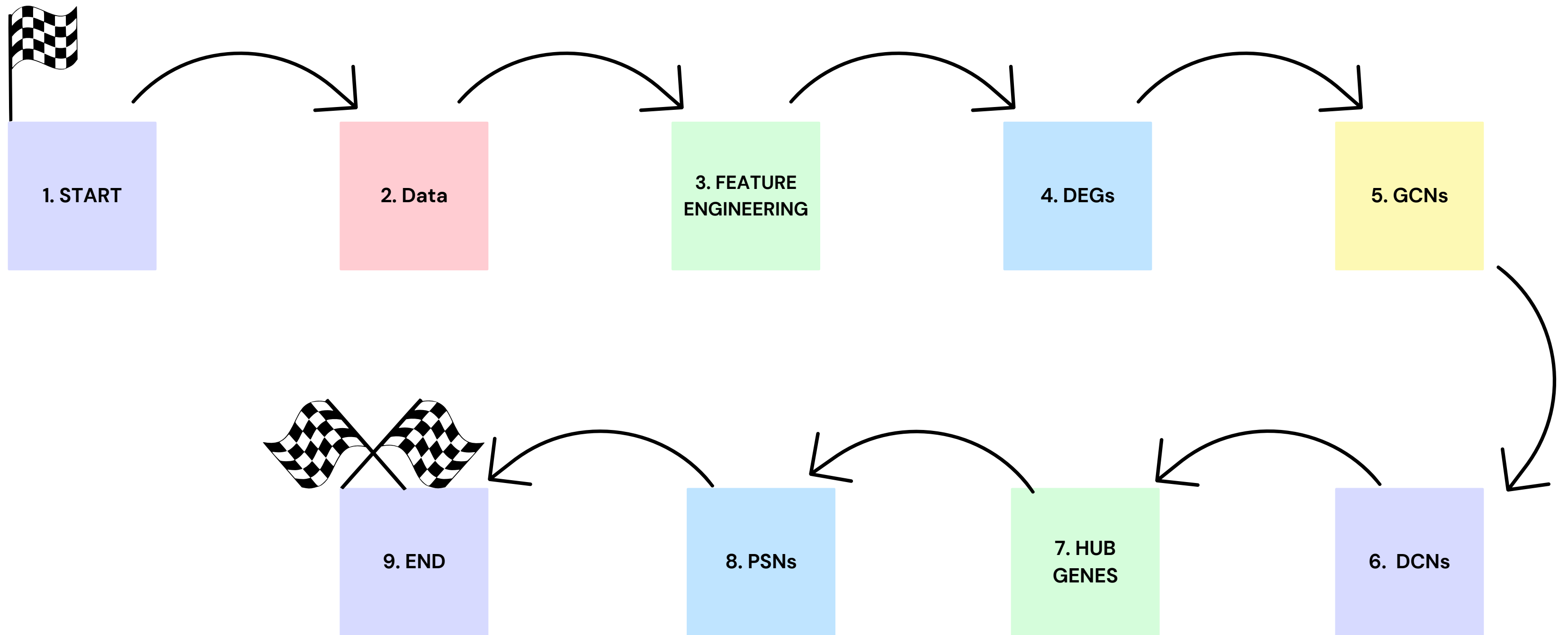


ANALYSIS OF PRAD HUB GENES

Leonardo Lavagna
Group 14





1. START

2. Data

3. FEATURE
ENGINEERING

4. DEGs

5. GCNs

6. DCNs

7. HUB
GENES

8. PSNs

9. END

Overview

Prostate Adenocarcinoma (PRAD) is a **common cancer affecting men** in their 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.

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Genomic Data

From the **Cancer Genome Atlas** we retrieved a collection of genomic data of about **575 genes and 500 patients**. For our analysis we handled **two datasets extracted** from TCGA: the **tumor tissue** and the **normal tissue**.

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DATA CLEANING AND
PROCESSING

ELIMINATE
NOISY DATA

GENOMIC DATA

FILTER
PATIENTS

FILTER
GENES

1. START

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4. DEGs

5. GCNs

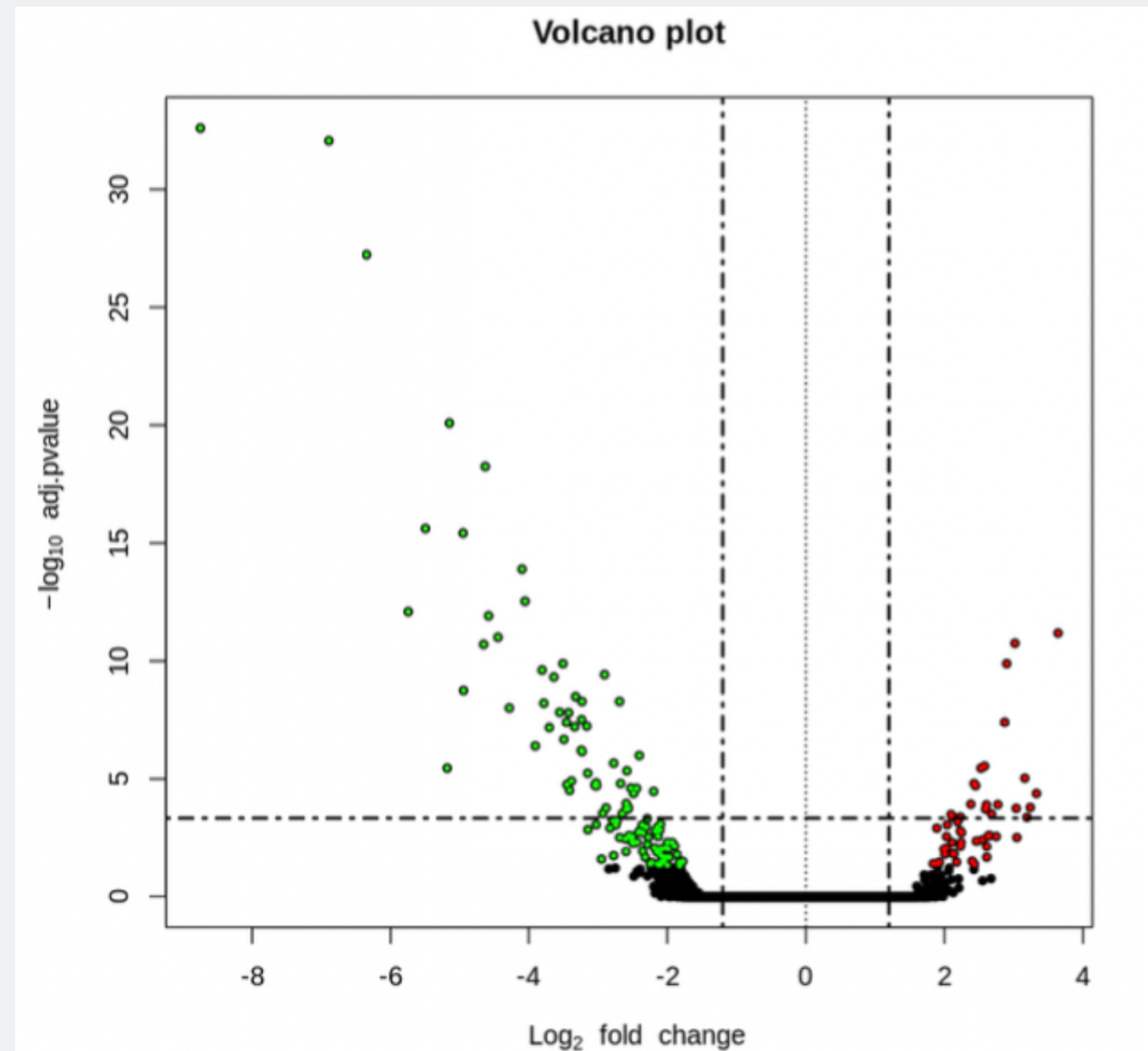
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DEGs WITH DIFFERENT
KINDS OF THRESHOLDS



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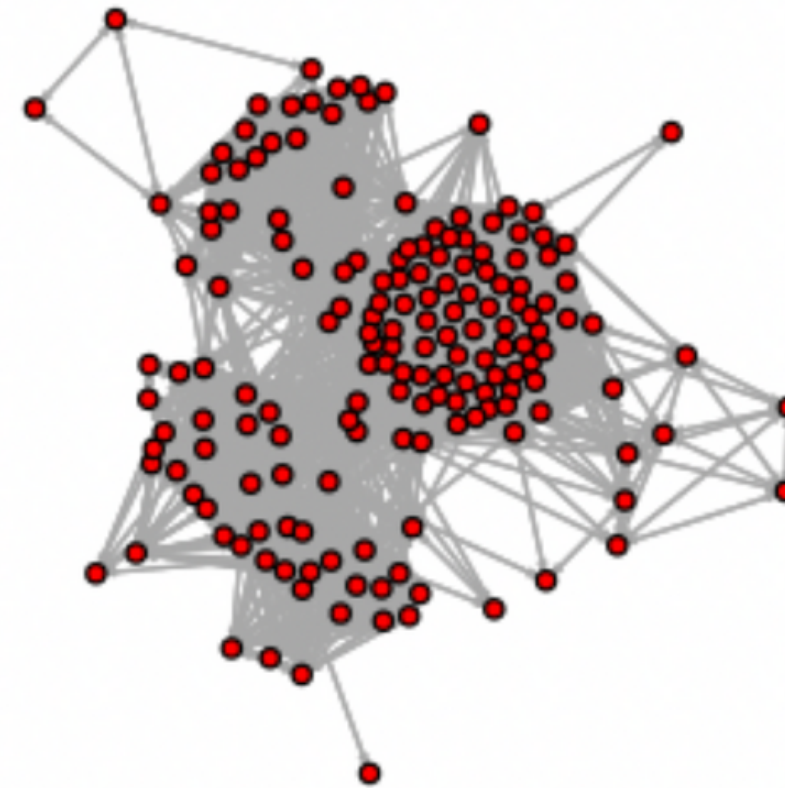
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GCNs: TUMOR vs NORMAL

Co-expression network TUMOR



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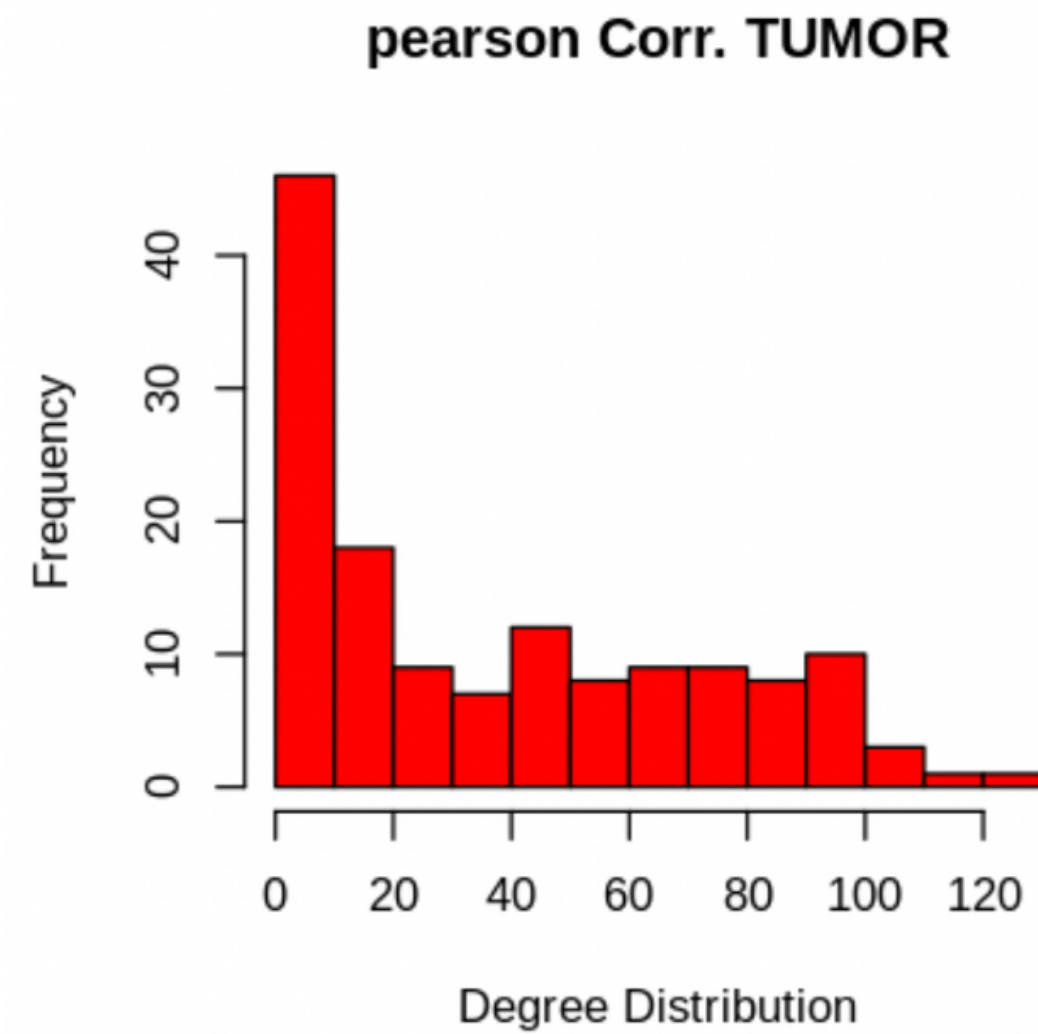
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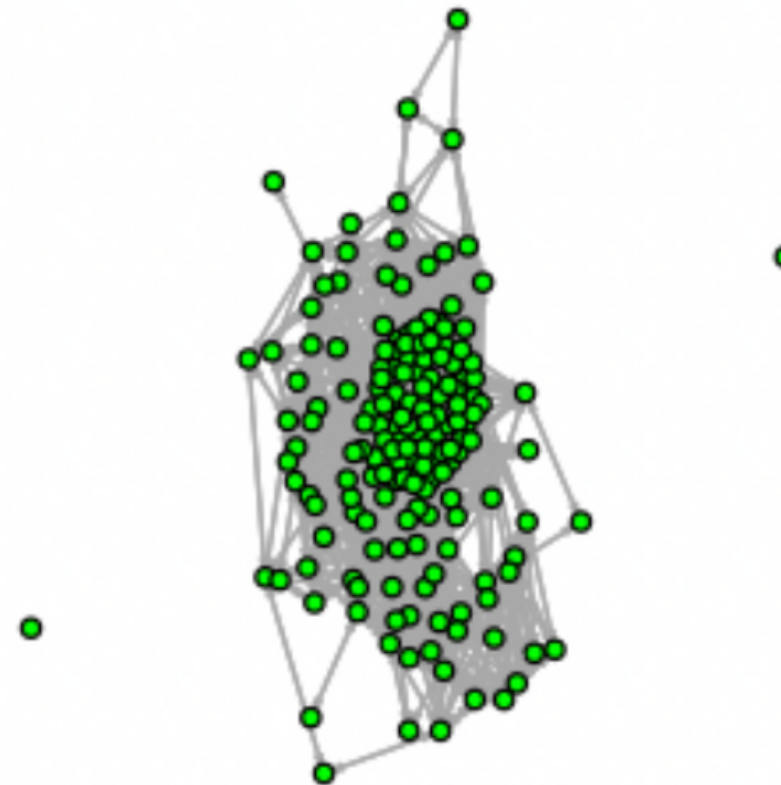
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GCNs: TUMOR vs NORMAL

Co-expression network NORMAL



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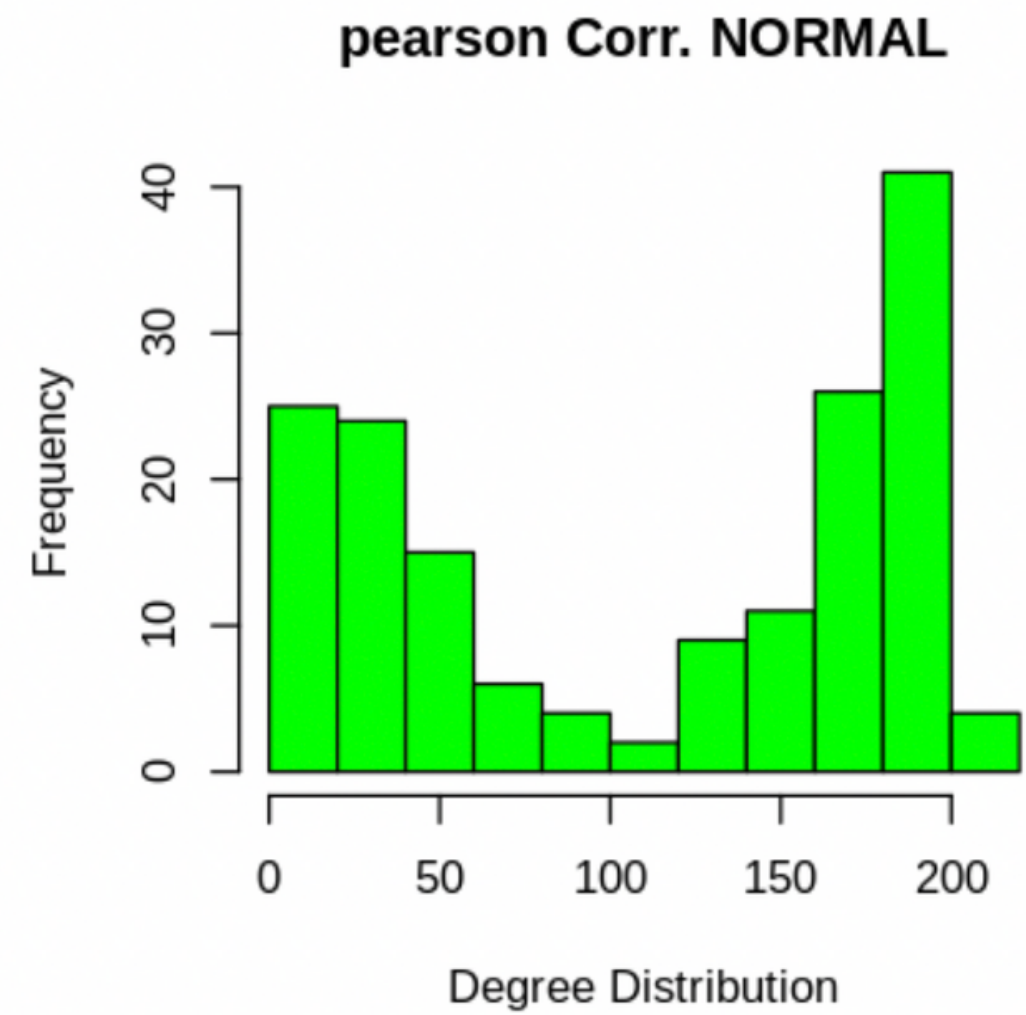
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GCNs: TUMOR vs NORMAL



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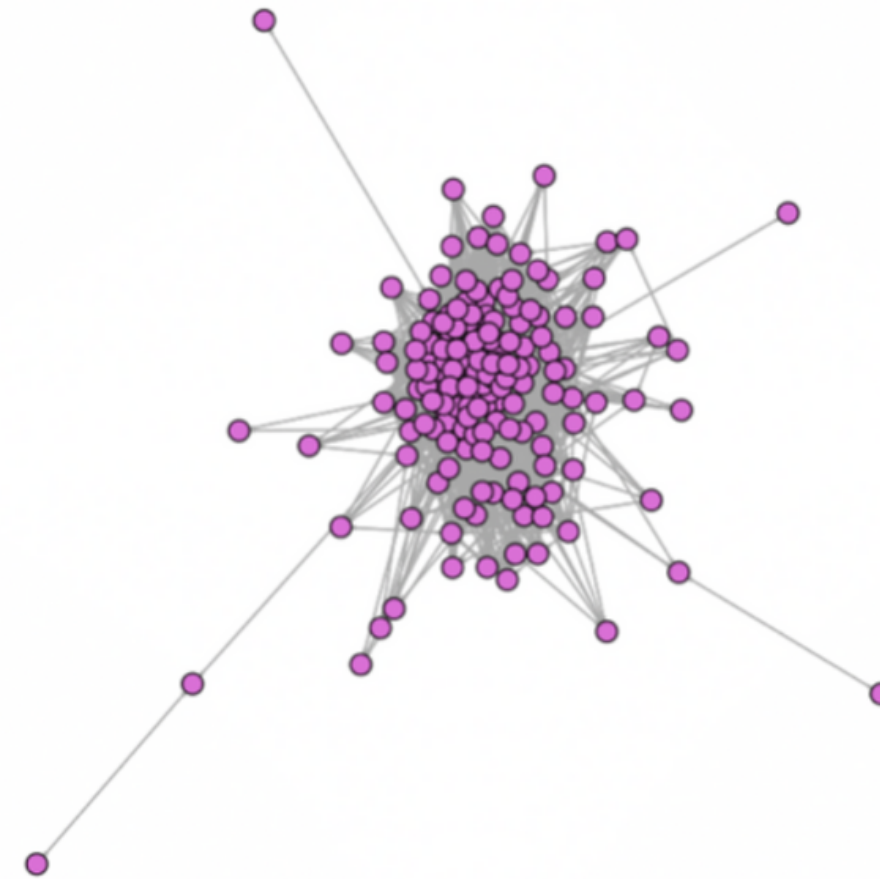
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DCNs: TUMOR vs NORMAL

Differential Co-expression network in TUMOR vs NORMAL



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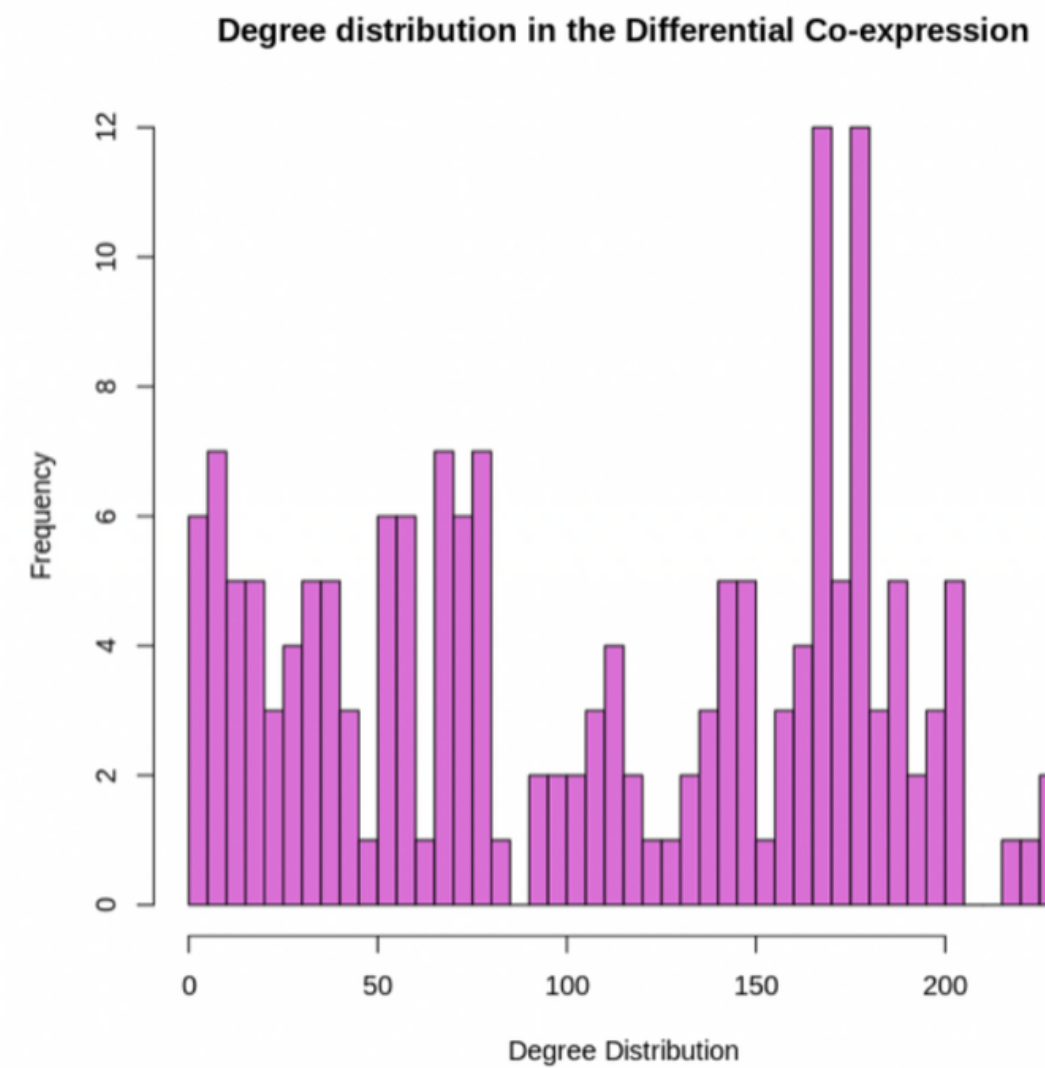
6. DCNs

7. HUB
GENES

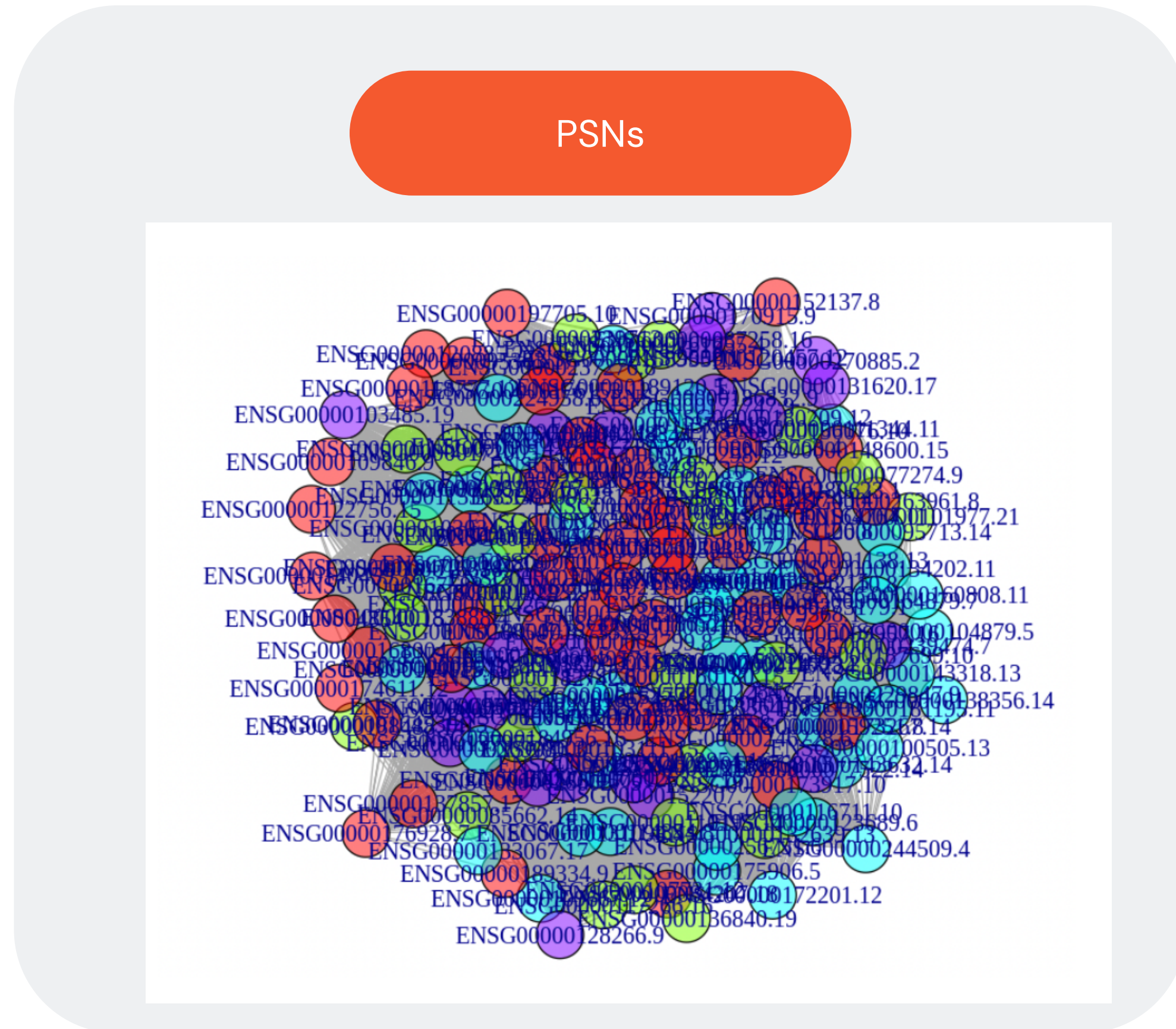
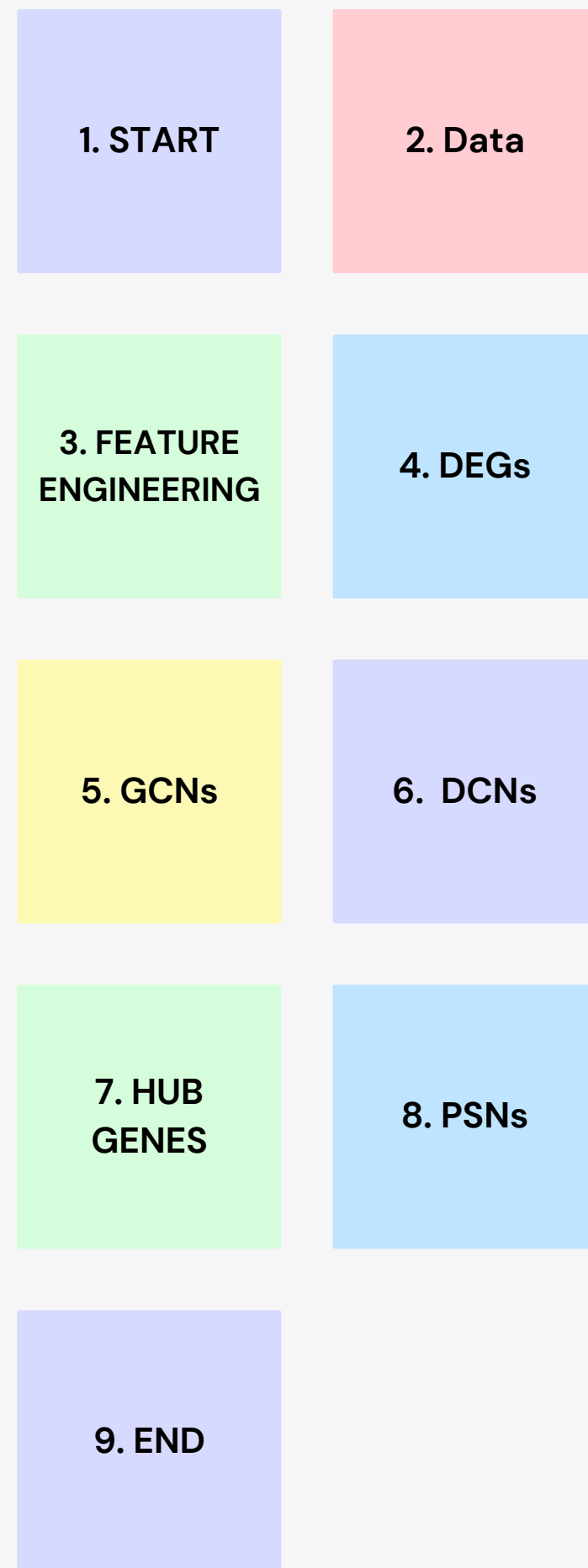
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DCNs: TUMOR vs NORMAL







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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 those hub genes were present.