

BINF 5003: Data Mining, Modeling, and Biostatistics

Week 14 - Module 6

Review

Gene Set Enrichment Analysis (GSEA)

What are gene sets?

- A **gene set** is a collection of genes grouped together based on shared characteristics, such as:
 - Biological pathways (e.g., glycolysis, immune response).
 - Chromosomal location.
 - Functional annotations (e.g., transcription factors, kinase activity).
 - Gene expression patterns under certain conditions.
- Examples of widely used gene sets:
 - **KEGG (Kyoto Encyclopedia of Genes and Genomes)**: Pathway-based groupings.
 - **Gene Ontology (GO)**: Grouped by molecular function, cellular component, and biological process.
 - **Hallmark gene sets**: Broad biological themes (e.g., hypoxia, apoptosis).

Why Gene Sets Matter?

- Biological systems often function through groups of genes working together in pathways or networks
- Analyzing individual genes may miss the bigger picture, but focusing on gene sets can reveal system-level changes

Gene Set Enrichment Analysis (GSEA) – example

Gene
ATM
MAPK3
MTOR
RAF1
PIK3CA
AKT1
MAPK1

Sort the genes based on fold change or statistical significance
(e.g., p-value)



Gene	Rank
MAPK1	1
MAPK3	2
RAF1	3
...	...
PIK3CA	18
AKT1	19
MTOR	20

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MAPK Pathway

PI3K-Akt Pathway

Gene Set Enrichment Analysis (GSEA) – example

Our genes of interest

Gene	Rank*
MAPK1	1
MAPK3	2
RAF1	3
...	...
PIK3CA	18
AKT1	19
MTOR	20

MAPK pathway is **upregulated** because its genes are at the top of the ranking

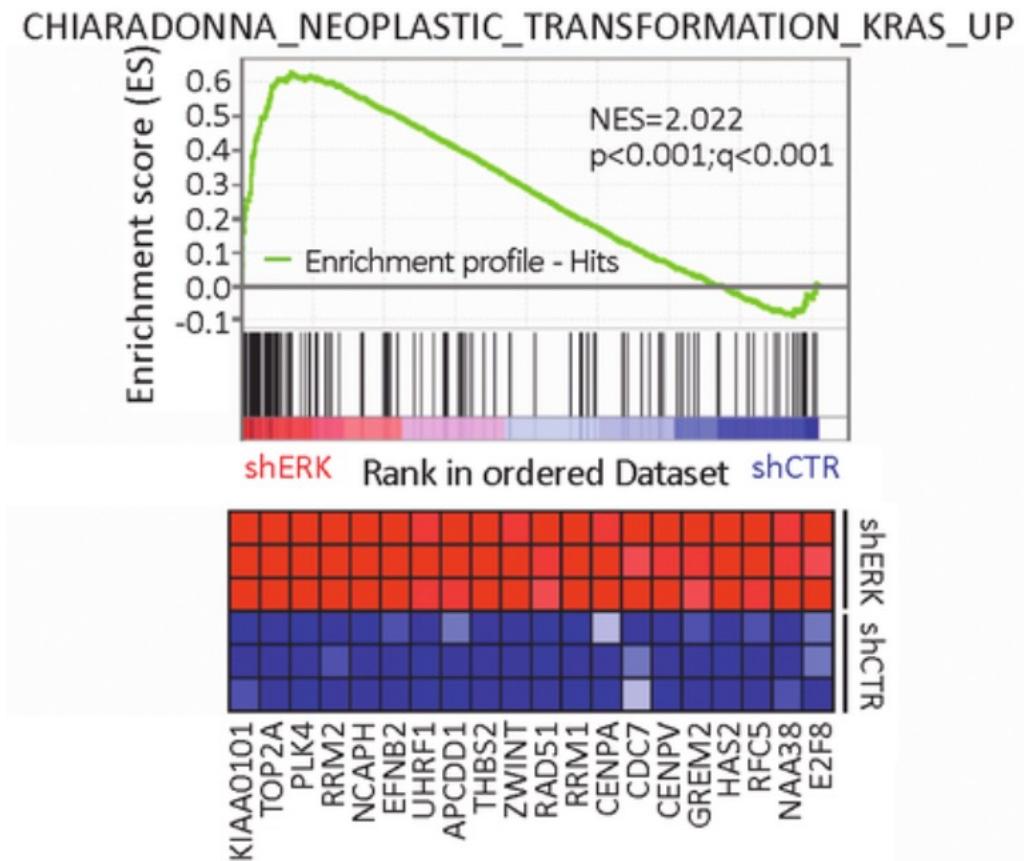
PI3K-Akt pathway is **down regulated** because its genes are at the bottom of the ranking

MAPK Pathway Genes :
MAPK1 , MAPK3, RAF1 ,BRAF, MEK1, KRAS,...

PI3K-Akt Pathway Genes:
PIK3CA, AKT1, MTOR, MYC, FOXO1, HIF1A, ...

Visualizing GSEA Results - Enrichment Plots

- Show the distribution of a gene set across a ranked list
- Peaks indicate where gene sets are enriched; steep peaks suggest strong enrichment



Let's finish the GSEA tutorial