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bioinformaticsdotca.github.io



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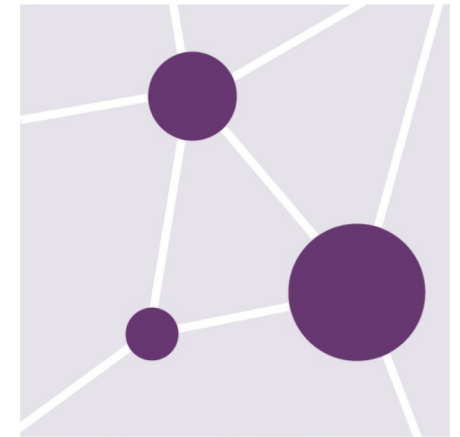


# Module 4 Lab

Constance Li

Pathway and Network Analysis

May 13, 2026





# Learning Objectives

1. Analyze **mutation data** using **ReactomeFI**
2. Use **GeneMANIA** to **integrate** findings across expression and mutation data types
3. Draw biological conclusions from multi-omics pathway analysis

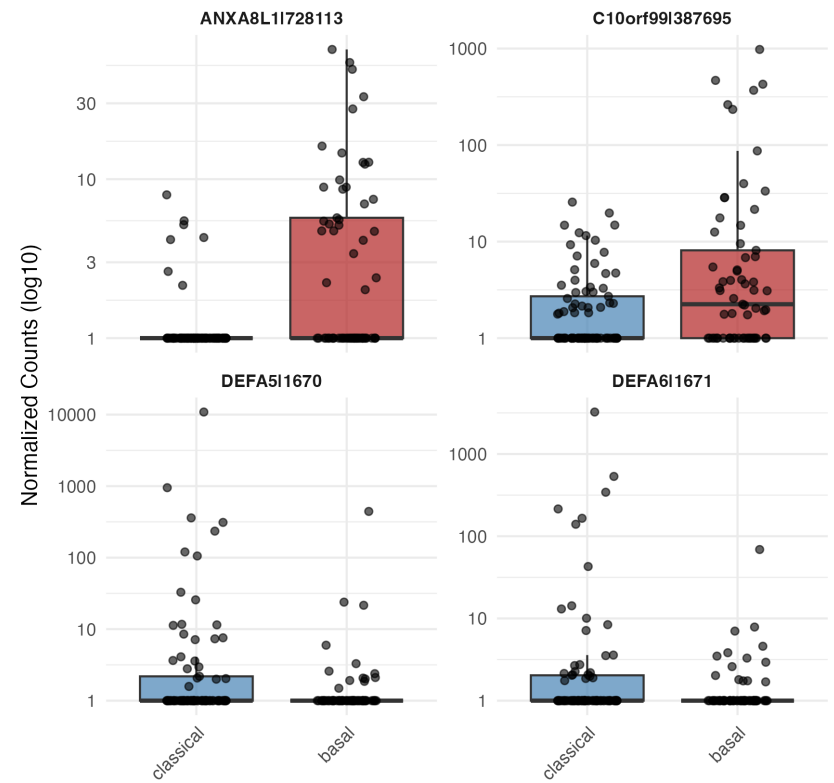


# Recall Lab 2a

Volcano Plot: Basal vs Classical  
Positive log<sub>2</sub>FC = higher in Basal



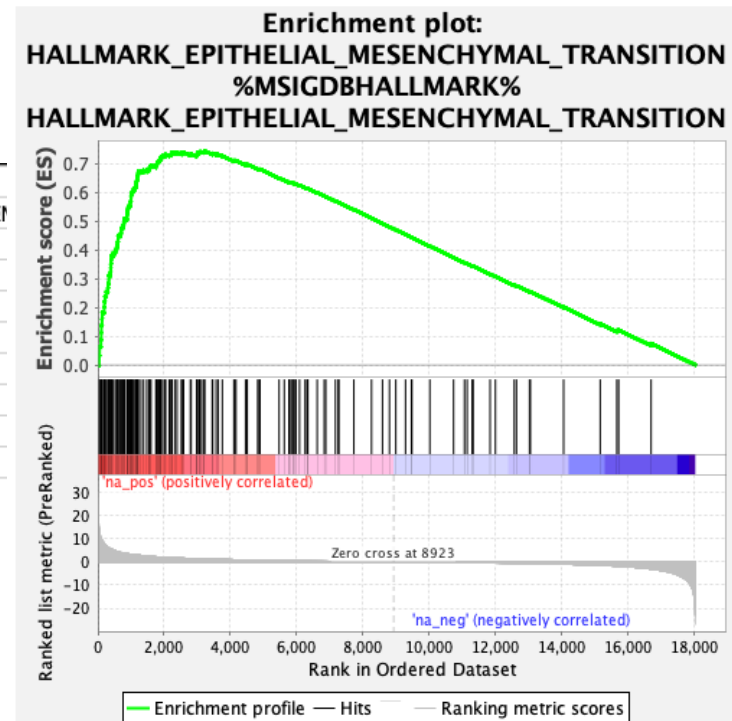
Expression of Top Differentially Expressed Genes





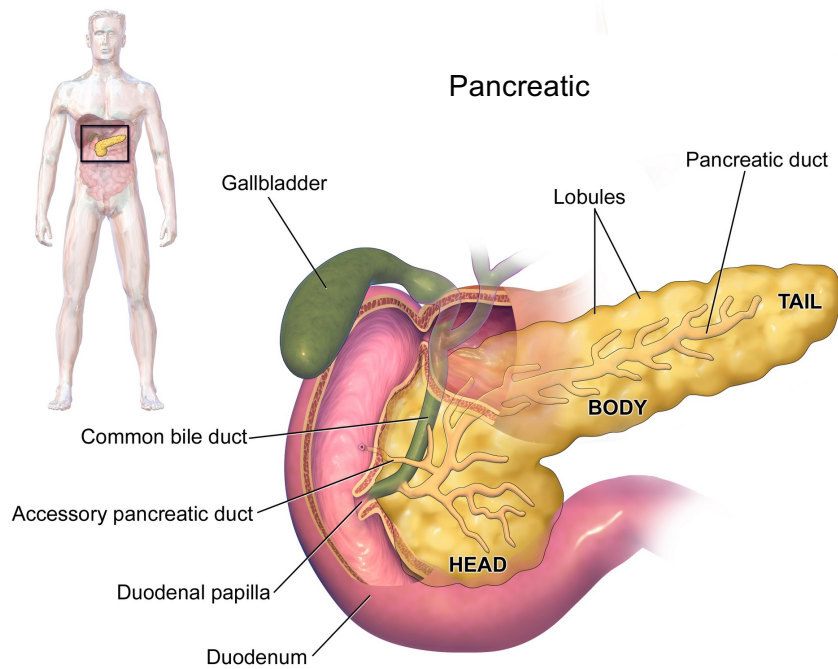
# Recall Lab 2c

NAME
HALLMARK_EPITHELIAL_MESENCHYMAL_TRANSITION%MSIGDBHALLMARK% <b>HALLMARK_EPITHELIAL_MESENCHYMAL_TRANSITION</b>
SKIN DEVELOPMENT%GOBP%GO:0043588
INTERMEDIATE FILAMENT ORGANIZATION%GOBP%GO:0045109
KERATINOCYTE DIFFERENTIATION%GOBP%GO:0030216
INTERMEDIATE FILAMENT-BASED PROCESS%GOBP%GO:0045103
INTERMEDIATE FILAMENT CYTOSKELETON ORGANIZATION%GOBP%GO:0045104
KERATINIZATION%REACTOME%R-HSA-6805567.5
KERATINIZATION%GOBP%GO:0031424
EPIDERMIS DEVELOPMENT%GOBP%GO:0008544
ELASTIC FIBRE FORMATION%REACTOME DATABASE ID RELEASE 96%1566948



	NOM p-val	FDR q-val
7	0	0
8	0	0
8	0	0
4	0	0
3	0	0
7	0	0
7	0	0
5	0	0
2	0	0
7	0	8.48E-04

# Remember PAAD?



<https://en.wikipedia.org/wiki/Pancreas>



# What is “integration”?

The process of **combining** separate parts or systems into a unified whole

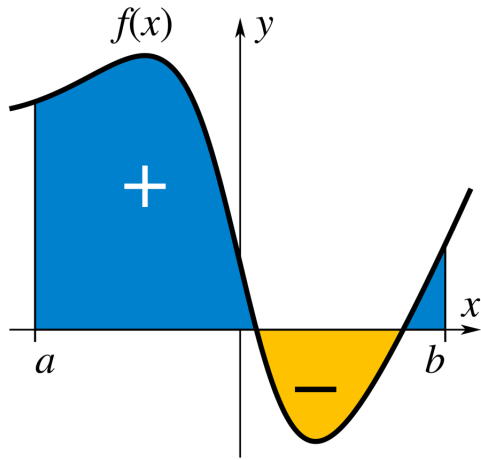
Integrate different  
molecular profiles

Integrate a collection  
of networks

Integrate everything  
you’ve learned in this  
workshop



# Let's integrate



- Download the lab materials:
  - [https://bioinformaticsdotca.github.io/PNA\\_CalSask-2605/module-4.html](https://bioinformaticsdotca.github.io/PNA_CalSask-2605/module-4.html)
  - Module 4: Instructions PDF, gene list text files
- Make sure Cytoscape is installed
  - You'll need the ReactomeFI and GeneMANIA apps



# Information and data sources

## Moffitt subtype paper

- Moffitt RA, Marayati R, Flate EL, Volmar KE, Loeza SGH, Hoadley KA, et al. Virtual microdissection identifies distinct tumor- and stroma-specific subtypes of pancreatic ductal adenocarcinoma. Nat Genet. 2015 Oct;47(10):1168–78. doi:[10.1038/ng.3398](https://doi.org/10.1038/ng.3398)

## Mutations data

- Raphael BJ, Hruban RH, Aguirre AJ, Moffitt RA, Yeh JJ, Stewart C, et al. Integrated Genomic Characterization of Pancreatic Ductal Adenocarcinoma. Cancer Cell. 2017 Aug;32(2):185-203.e13. doi:[10.1016/j.ccell.2017.07.007](https://doi.org/10.1016/j.ccell.2017.07.007)

## RNA-seq counts data

- <https://gdac.broadinstitute.org/>