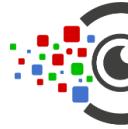


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 AI Image<sup>Lab</sup>



**DECIDER**

Improving clinical decisions in cancer

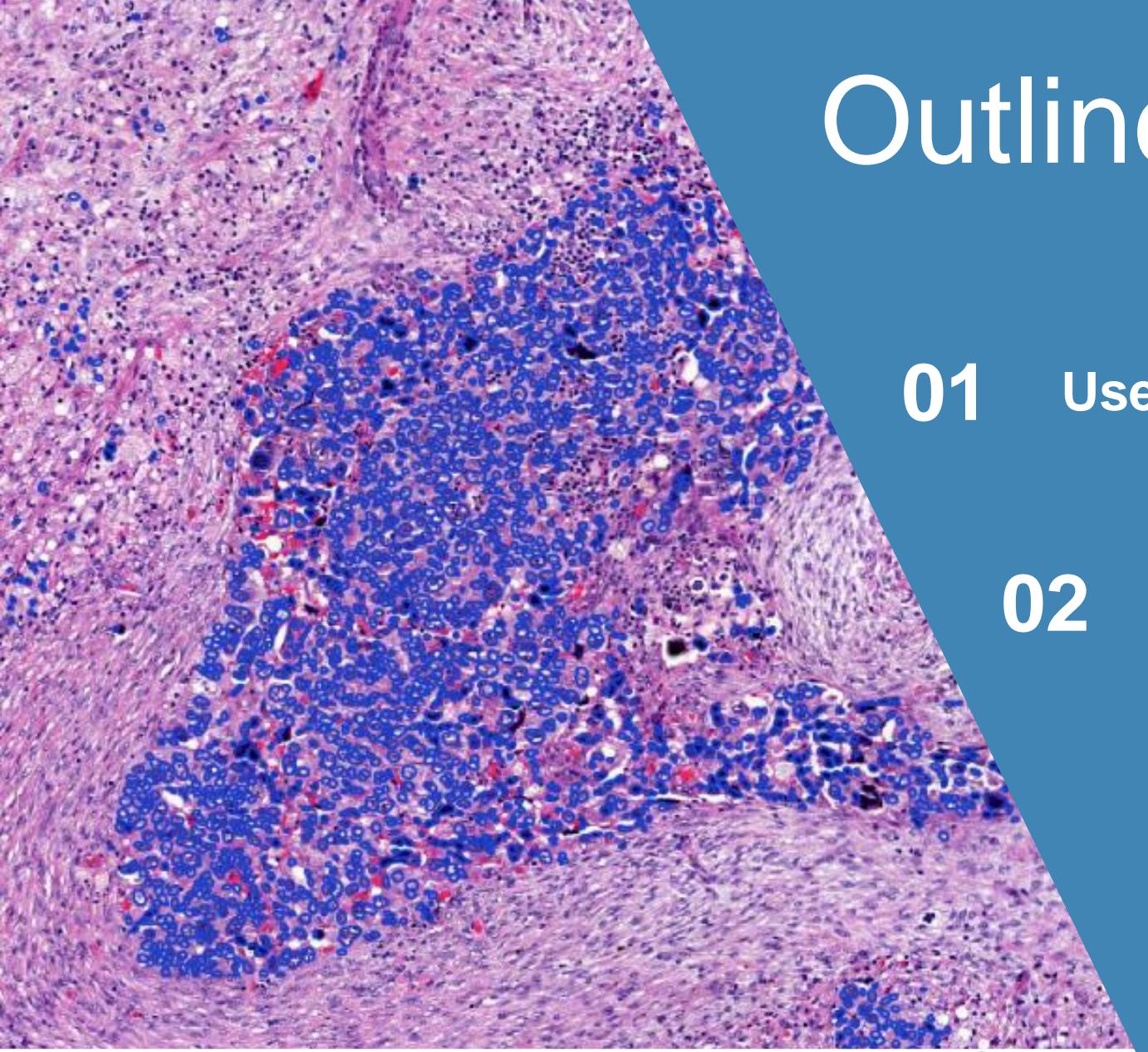
# Graph NN LAB

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# Outline

01 Useful material

02 Lab assignment

# Useful material

**Start practicing with Graph NN implementation with these notebooks:**

## **Introduction**

<https://colab.research.google.com/drive/1h3-vJGRVloF5zStxL5I0rSy4ZUPNsjy8#scrollTo=NgcpV4rjAWy->

## **Node classification**

[https://colab.research.google.com/drive/14OvFnAXggxB8vM4e8vSURUp1TaKnovzX#scrollTo=9r\\_VmGMukf5R](https://colab.research.google.com/drive/14OvFnAXggxB8vM4e8vSURUp1TaKnovzX#scrollTo=9r_VmGMukf5R)

## **Graph classification**

[https://colab.research.google.com/drive/1l8a0DfQ3fl7Njc62\\_mVXUlcA1eUclnb#scrollTo=qeORu4Zrs8Zy](https://colab.research.google.com/drive/1l8a0DfQ3fl7Njc62_mVXUlcA1eUclnb#scrollTo=qeORu4Zrs8Zy)

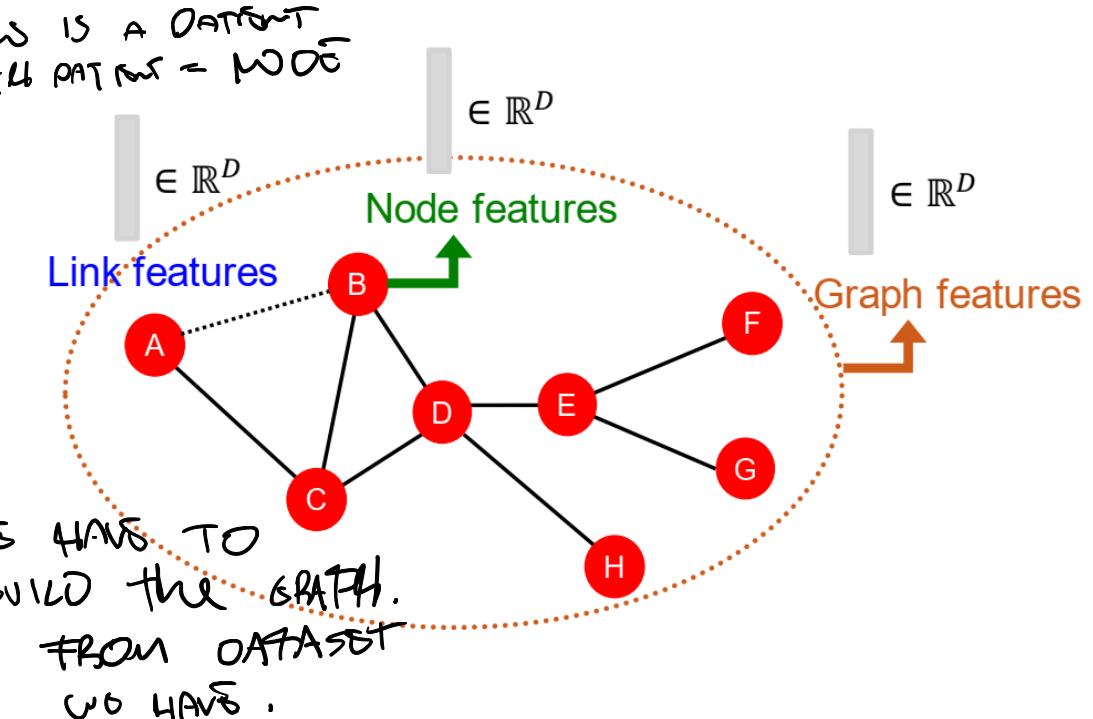
# Lab assignment

Implement a GNN to classify breast cancer patients in LUMINAL A / Luminal B

1. Each network node corresponds to a patient  $\Rightarrow$  EACH LIMS IS A DATASET EACH PATIENT = NODE
2. Node feature vector is the entire gene expression profile of the patient
3. Node label is the patient class (Luminal A/ Luminal B)
4. Edges are not provided; you must compute them using the Pearson correlation coefficient.

E.g.,  $x_{ij} = \text{corr}(\text{feat\_vector\_node}_i, \text{feat\_vector\_node}_j)$

5. Aim: predict patient node labels using a GNN
6. Are the performances better or worse compared to an MLP classifier?



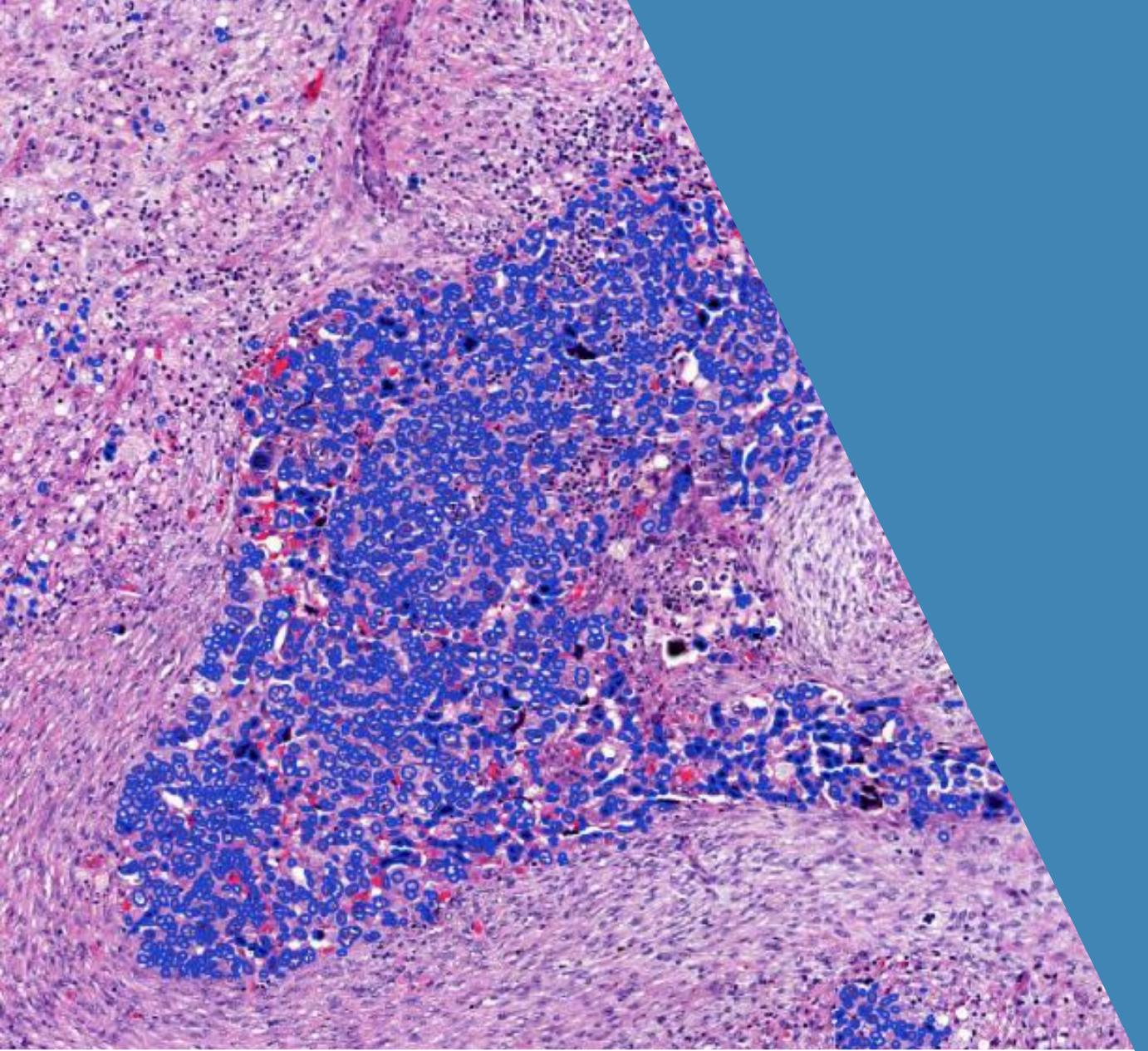
# Course Folder

You are encouraged to share your scripts on the course folder, to receive comments and feedback from colleagues and instructors.

[https://drive.google.com/drive/folders/1ynFYoc3xicaYhSi1X62w9k\\_JAj2fUrox?usp=sharing](https://drive.google.com/drive/folders/1ynFYoc3xicaYhSi1X62w9k_JAj2fUrox?usp=sharing)

Please upload your solutions with the proper naming:

e.g., LAB3\_SURNAMEName



# Questions?

*Better a stupid question in  
class than a stupid answer  
in the exam*