

# TER5 PROJECT

Learning oncology biomarkers from histological tissue slide images  
using modern AI techniques.

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# INTRODUCTION

- **Whole Slide Images (WSIs)** are large, high-resolution scans used in digital pathology.
- **Detecting meaningful biomarkers within them is a complex visual task.**
- **We compare three different model families:**

**CNN: effective locally**

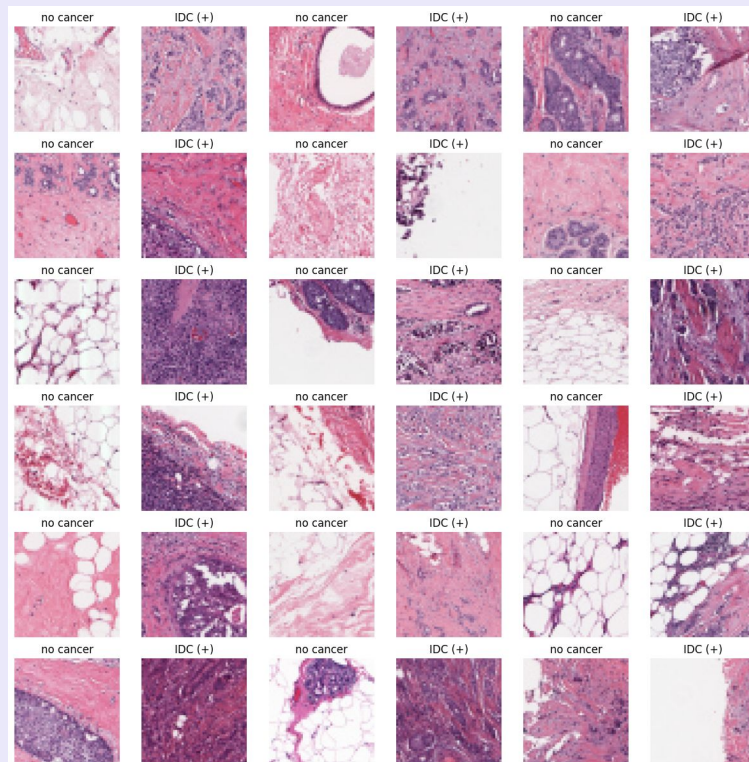
**Transformers: good for global context**

**Foundation Models: trained on millions of patches**

# Which Database did we use?

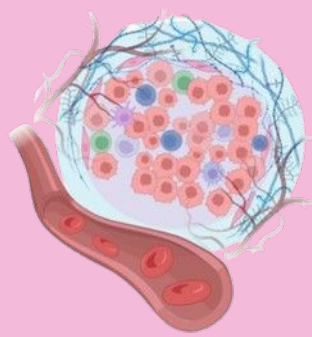
## IDC Breast Cancer Dataset (Patch-based):

- 162 WSIs
- 277,524 image patches (50 x 50 pixels)
- Binary classification: IDC-negative vs IDC-positive
- Extracted regions represent clinically relevant tumor zones.

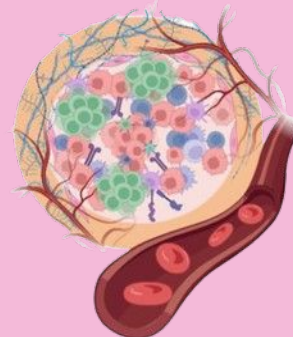


# Objective

The aim is to classify the dataset in a binary way in order to distinguish images with infiltrating ductal carcinoma (IDC) from those without.

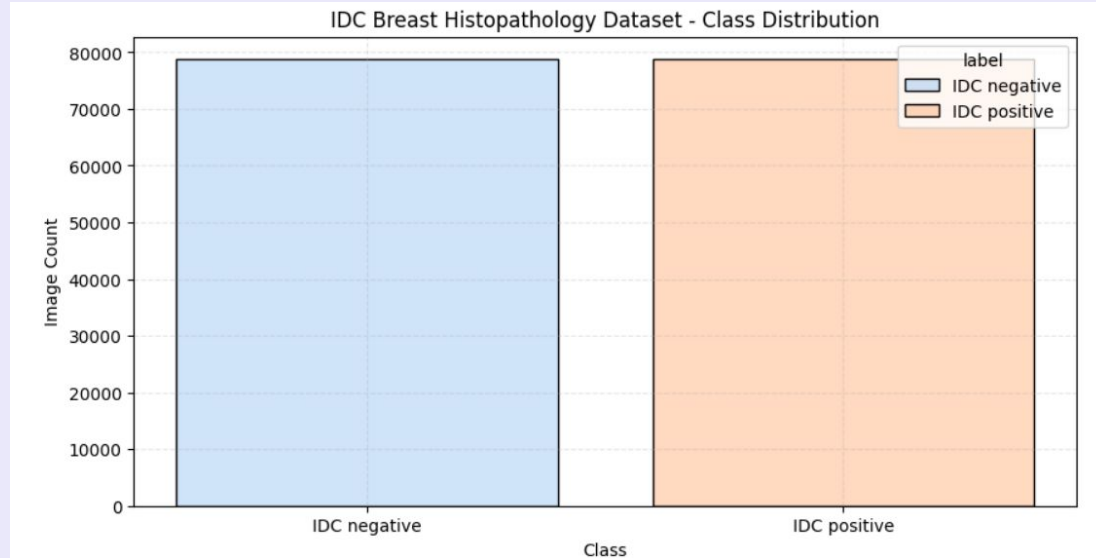


**IDC negative:**  
This class corresponds to tumor-free tissue images, i.e. cases where no infiltrating ductal carcinoma (IDC) is detected.



**IDC positive:**  
This class corresponds to tissue images that contain tumors, i.e. cases where infiltrating ductal carcinoma (IDC) is detected.

# Class Distribution

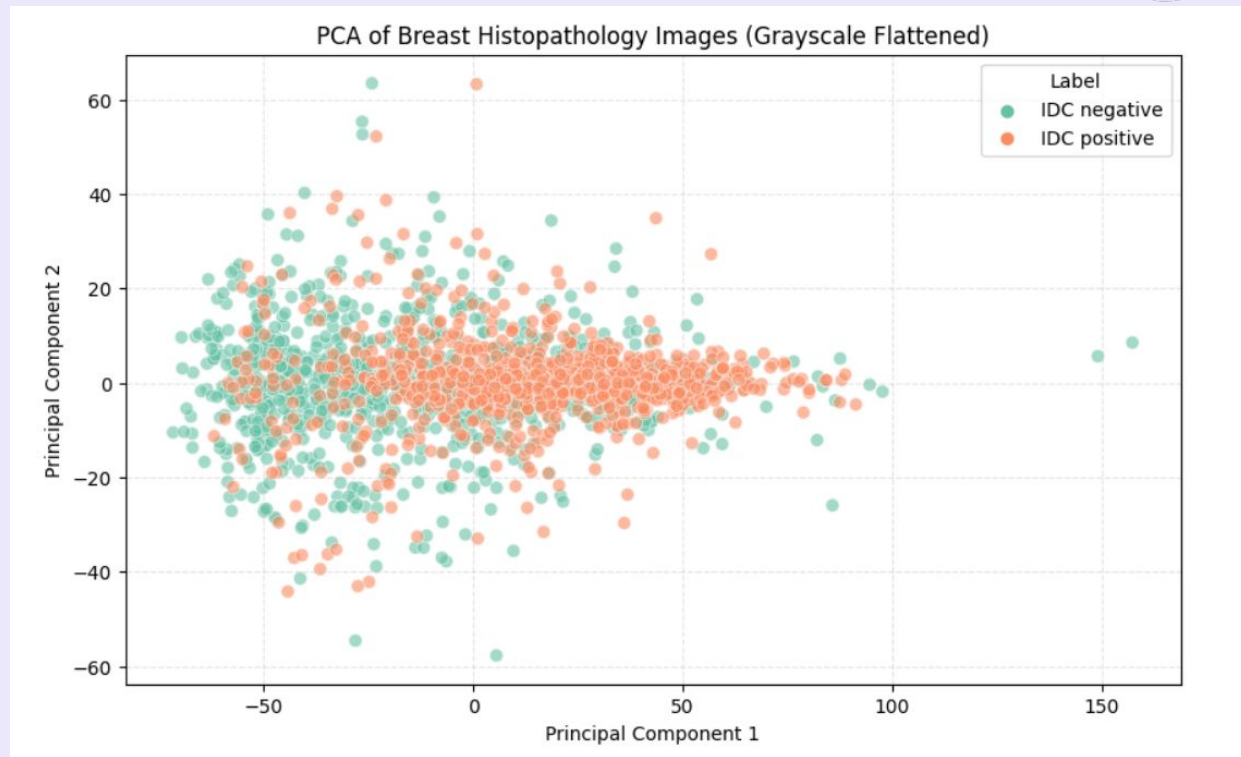


The two classes, *IDC negative* and *IDC positive*, are evenly distributed in the dataset, resulting in a well-balanced classification task.

# VISUALISATIONS : PCA

The PCA visualization shows that the two classes, *IDC negative* and *IDC positive*,

are largely intertwined in the reduced feature space.

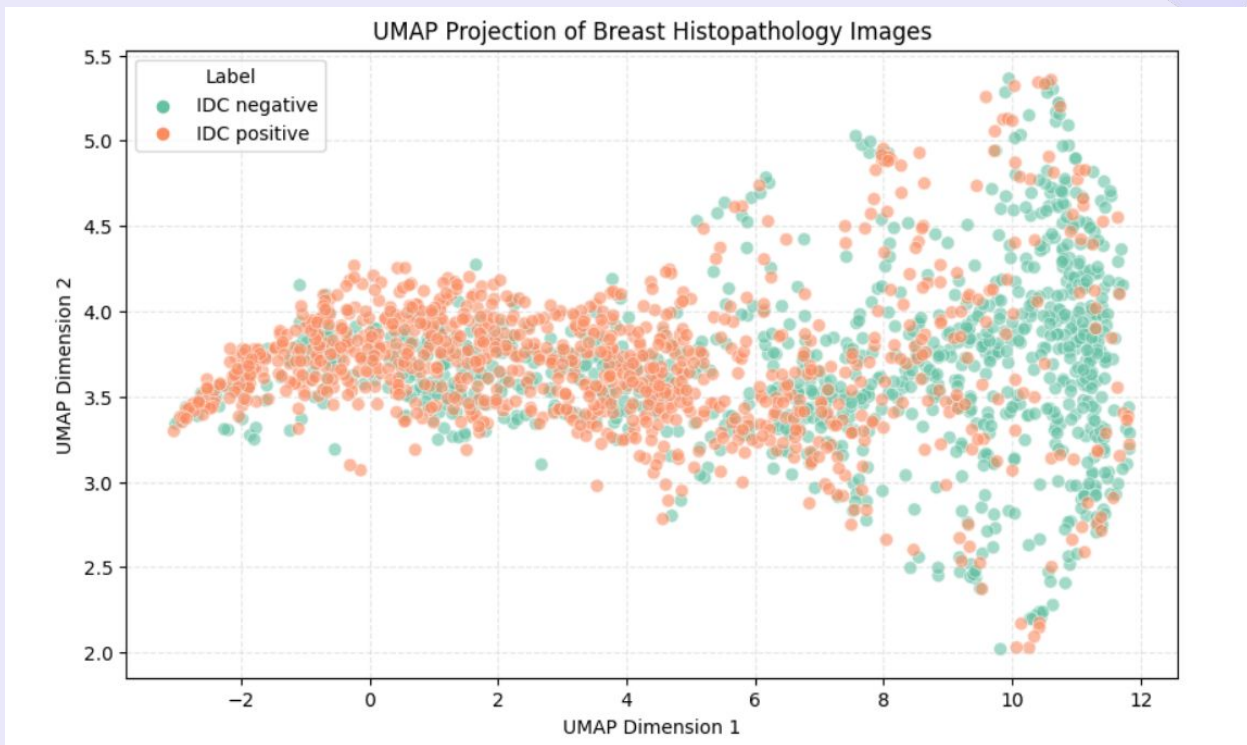




# VISUALISATIONS : U-MAP

This overlap likely reflects the subtle visual differences between tumor and non-tumor tissue,

which are difficult to capture without more advanced feature extraction methods such as CNN.







# Which Pipeline did we use?

## **1- WSI → Patch extraction**

Database contains the extracted patches

## **2- Feature embedding (via backbone model)**

The extracted embeddings can serve as digital biomarkers, capturing and summarizing key histological features relevant to diagnosis.

## **3- Classification**

Takes the extracted feature embeddings and assigns each patch to a specific category such as tumor or non-tumor based on learned patterns.



# CNN models

**EfficientNet: compound scaling, lightweight design**

- **Accuracy: train 65% / test 58%**
- **Loss: train 18.0 / test 10.0**

Poor

**ResNet: residual learning with skip connections**

- **Accuracy: train 95.2% / test 84.4%**
- **Loss: train 0.73 / test 0.65**

Good

# Transformers

**ViT: patch tokenization + self-attention**

- **Accuracy: train 94% / test 85%**
- **Loss: train 0.65 / test 0.45**

**Good**

**CTransPath: CNN + Swin Transformer hybrid**

- **Accuracy: train 98% / test 85%**
- **Loss: train 0.9 / test 0.1**

**Very Good**

# Foundation Model

**UNI: Unified Contrastive Learning for Histopathology, a general-purpose ViT encoder trained on 100M pathology patches**

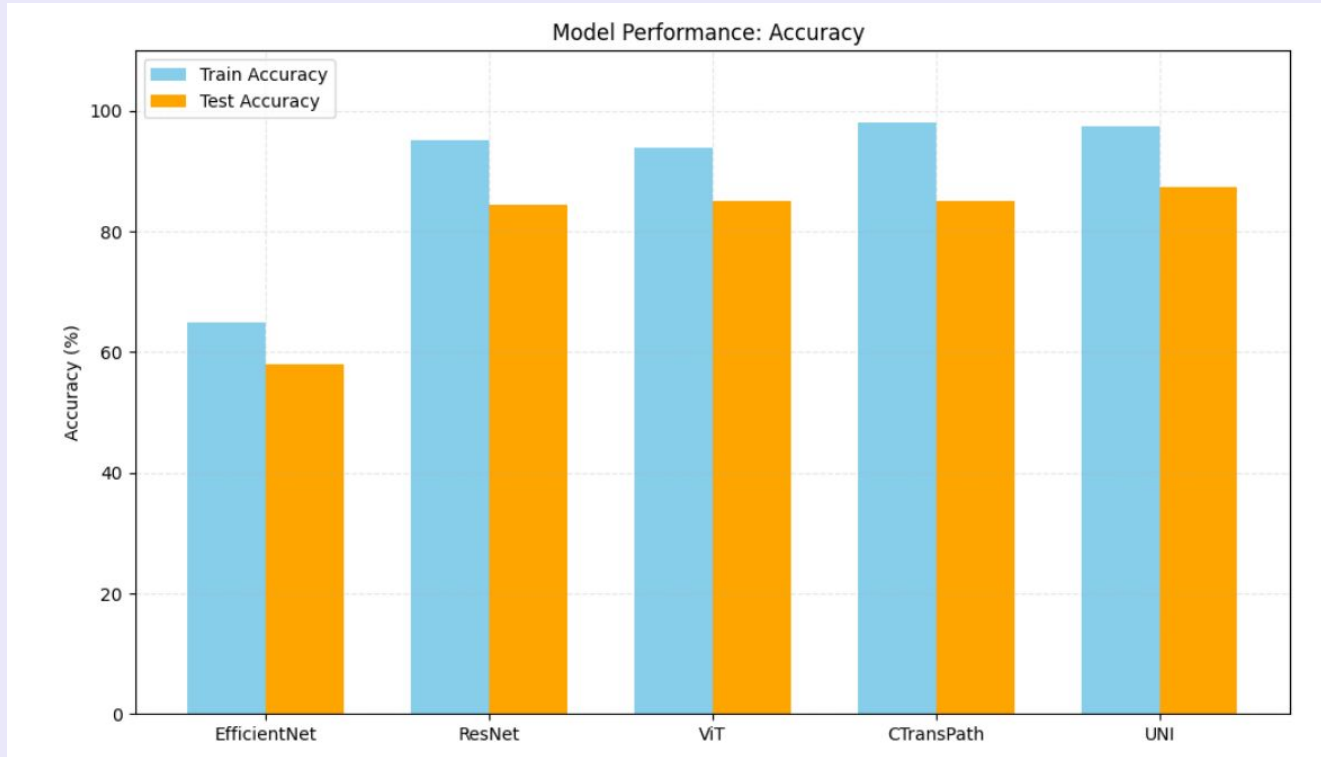
- **Accuracy: train 97.5% / test 87.5%**
- **Loss: train 0.54 / test 0.43**

**Best**

# Results

	Train Acc	Test Acc	Train Loss	Test Loss
EfficientNet	65.0%	61.36%	18.00	10.00
ResNet	95.2%	84.4%	0.73	0.65
ViT	94.5%	85.0%	0.65	0.45
CTransPath	98.0%	85.0%	0.54	0.43
UNI	97.5%	87.5%	0.54	0.43

# Results





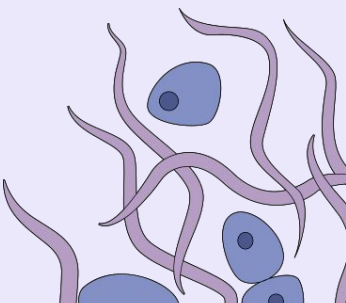
# Study Limitations

- **Limited GPUs → long training sessions (up to 20h per model)**
  - **Difficulty loading WSIs (very large images)**
  - **Lack of Model's documentation (ex: UNI)**
  - **Computational constraints, transformer based models require more compute time than it is practical.**
- 





# Conclusion

- **CNNs are fast and interpretable but lack global context**
  - **Transformers achieve better spatial reasoning**
  - **UNI outperforms others, but marginally**
  - **EfficientNet underperforms on generalization**
  - **GPU constraints limited hyperparameter tuning**
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# Perspectives



**01**

**Broader datasets (e.g., multi-organ)**

**02**

**Patch selection optimization**

**03**

**Semi- or Unsupervised learning**

The slide features a light pink background with four abstract, organic shapes in various shades of pink and magenta in the corners. The top-left shape is a simple wavy line. The top-right shape is more complex, with several small circles inside. The bottom-left shape is also complex, with several small circles and two white curved lines. The bottom-right shape is a simple wavy line.

# Thank You!

Thank you for your attention.

Questions?

# Kaggle Notebooks

**ResNet**

**EfficientNet**

**ViT**

**CTransPath**

**UNI**