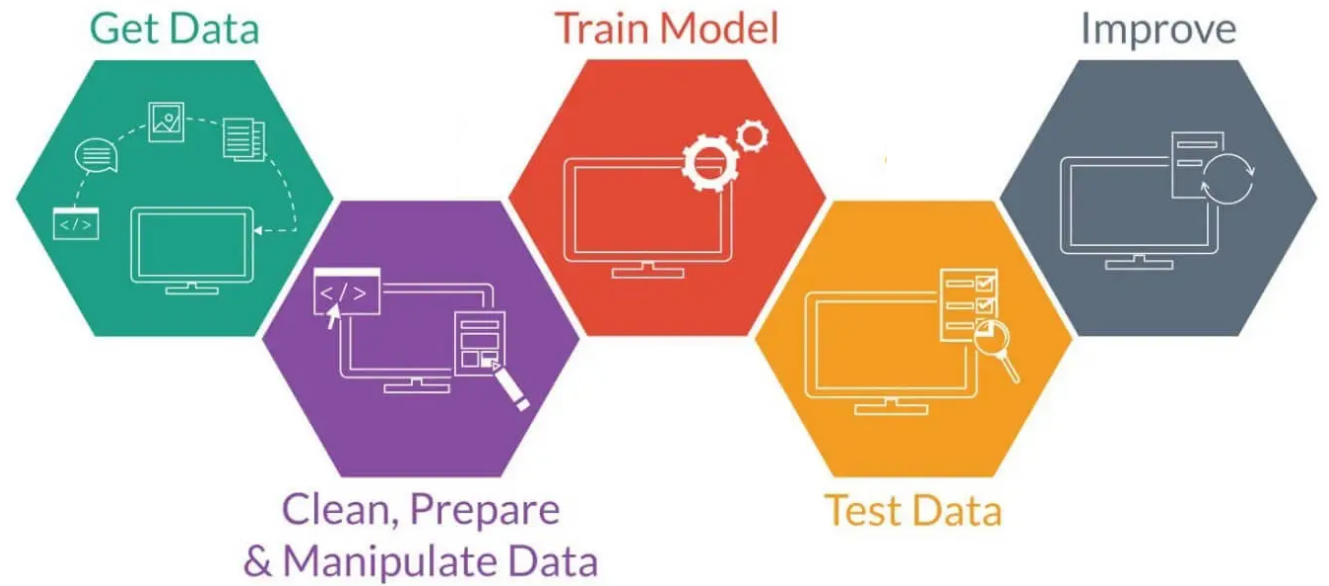


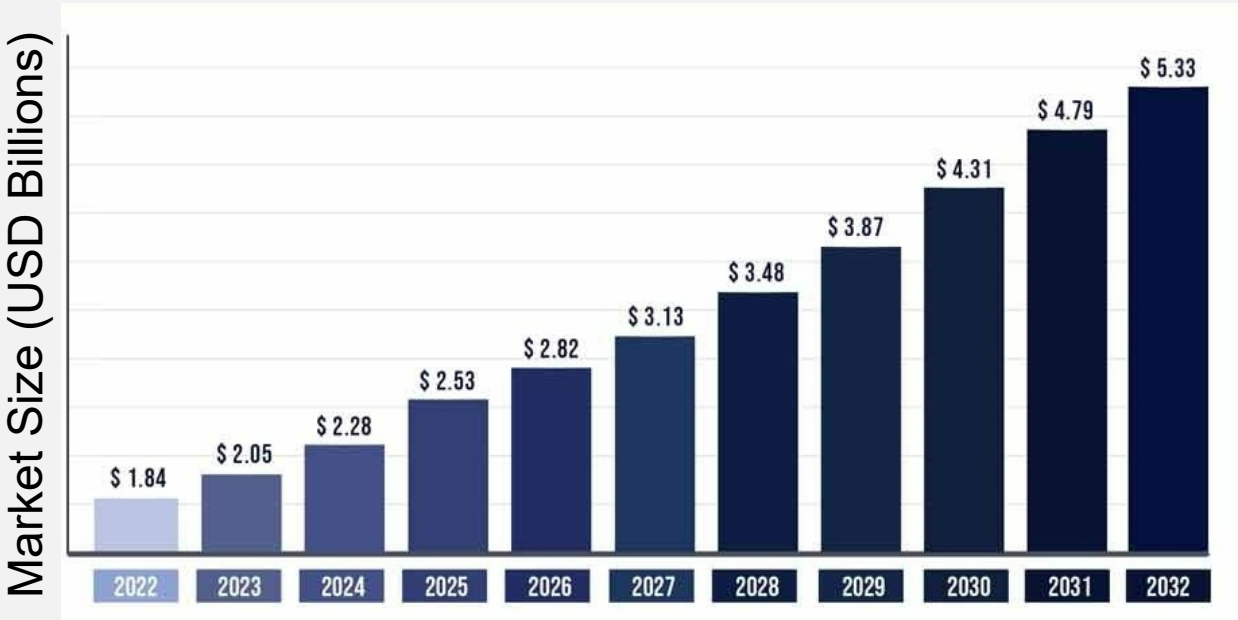
# ML IN DIGITAL PATHOLOGY

Yuan Nghiem  
Leo Grant Berman  
Albert Bulik



# PROBLEM STATEMENT

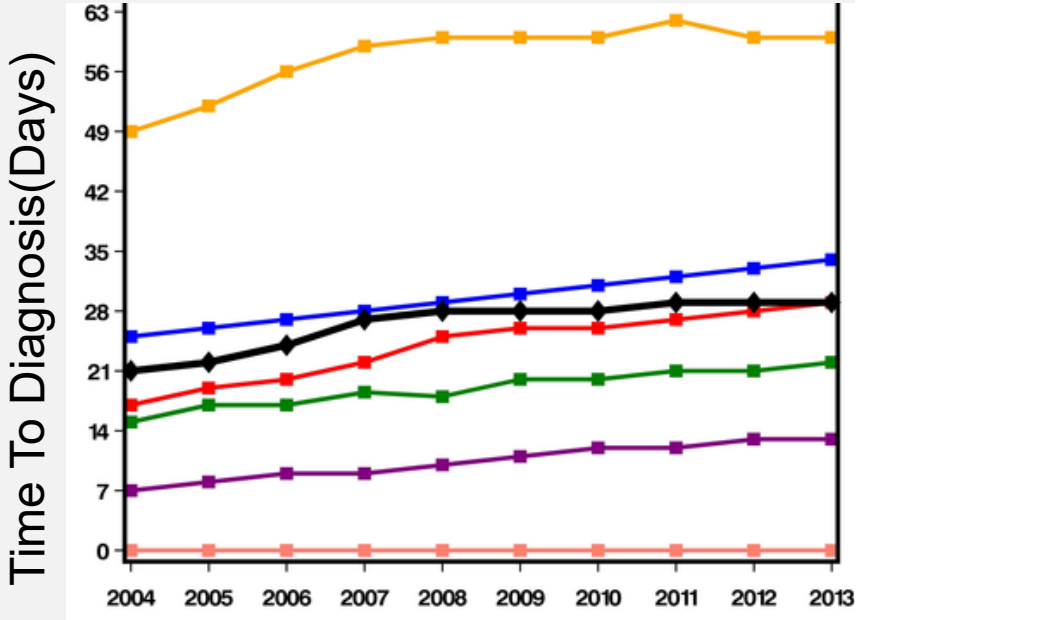
The demand for cancer diagnosis is increasing.



YEAR (2022 to 2032)

[SOURCE](#)

As a result, time to treatment is increasing.

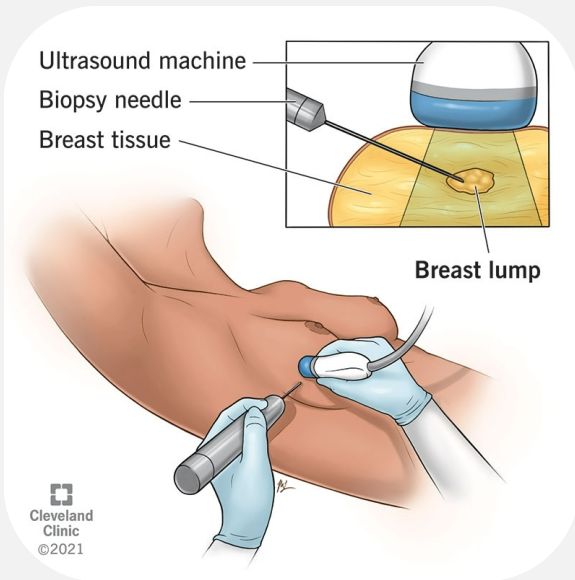


DIAGNOSIS YEAR (2004 to 2013)

[SOURCE](#)

# TEMPLE UNIVERSITY HEALTH DIGITAL PATHOLOGY (TUHDP) CORPUS

## Biopsy



## Staining

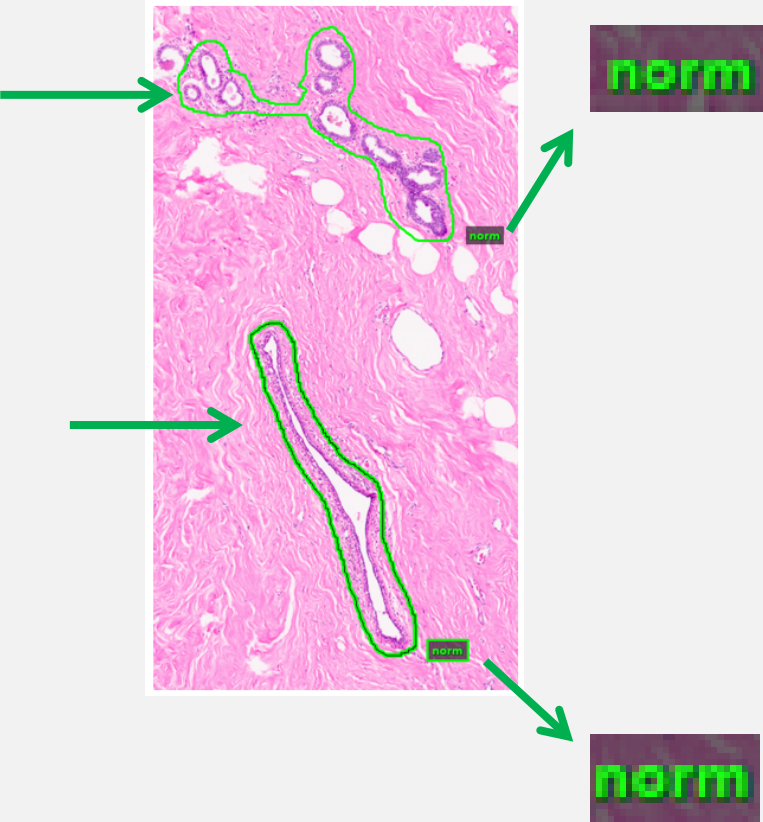


## Scanning

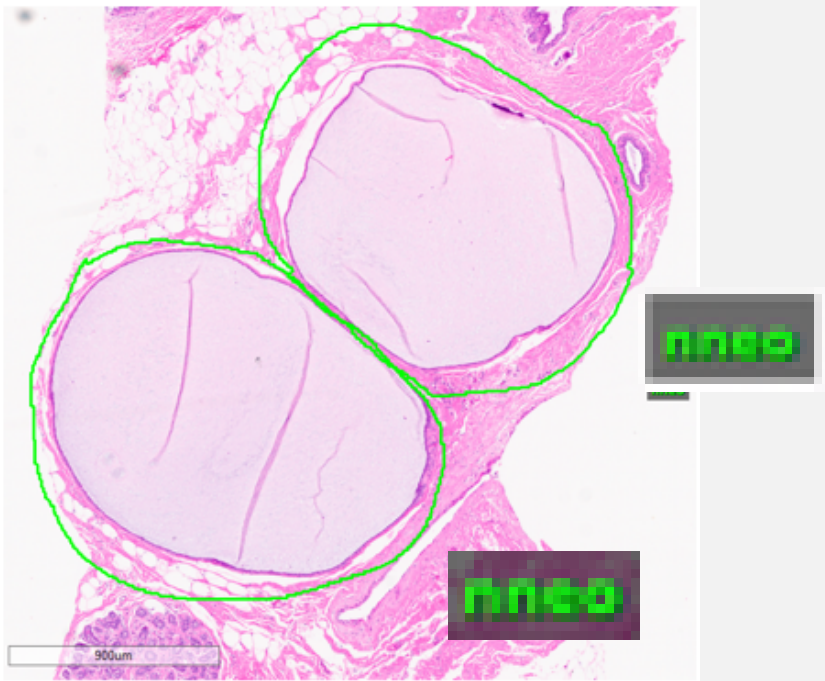


# LABELED DATA TYPES

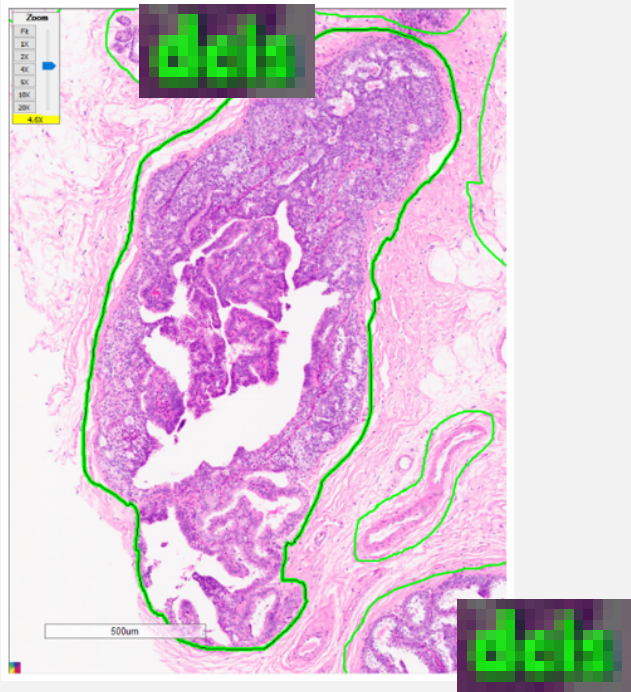
Non-Cancerous Types



Carcinogenic-Signs Type



Cancerous Types



# OUR PRODUCT

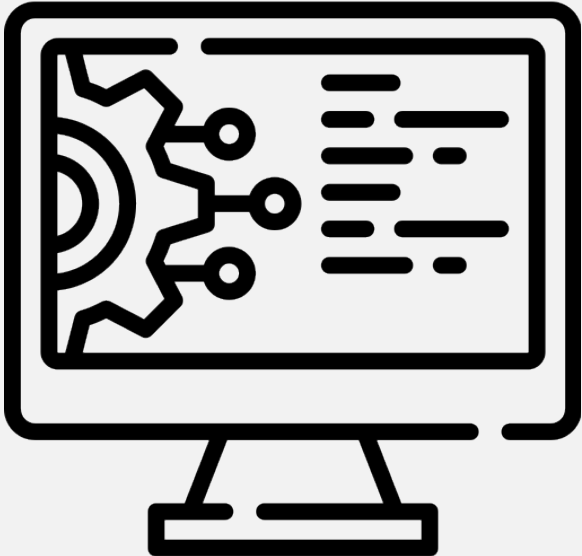
Scanning



(Data)



System



(Model)



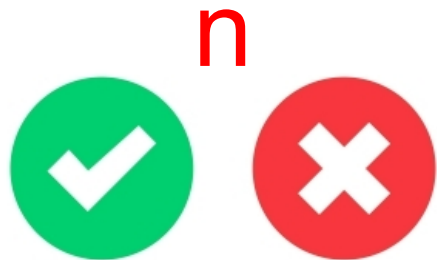
Prediction Report



(Output)

# DESIGN CRITERIA / REQUIREMENTS

## Image Classification



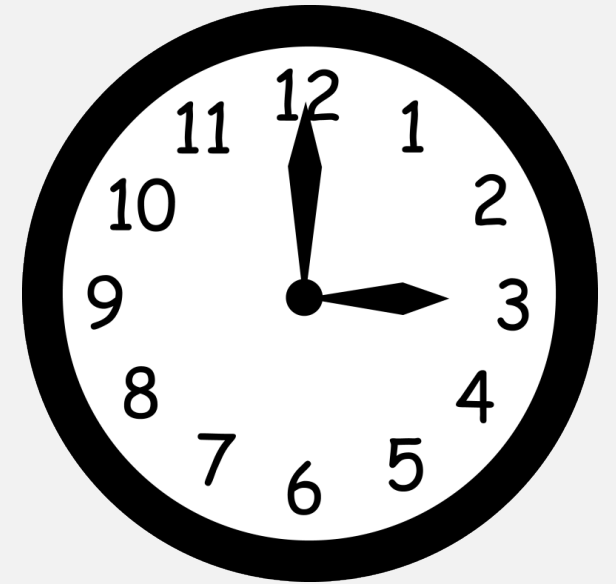
Whole Slide Image  
Classification

[Source](#)



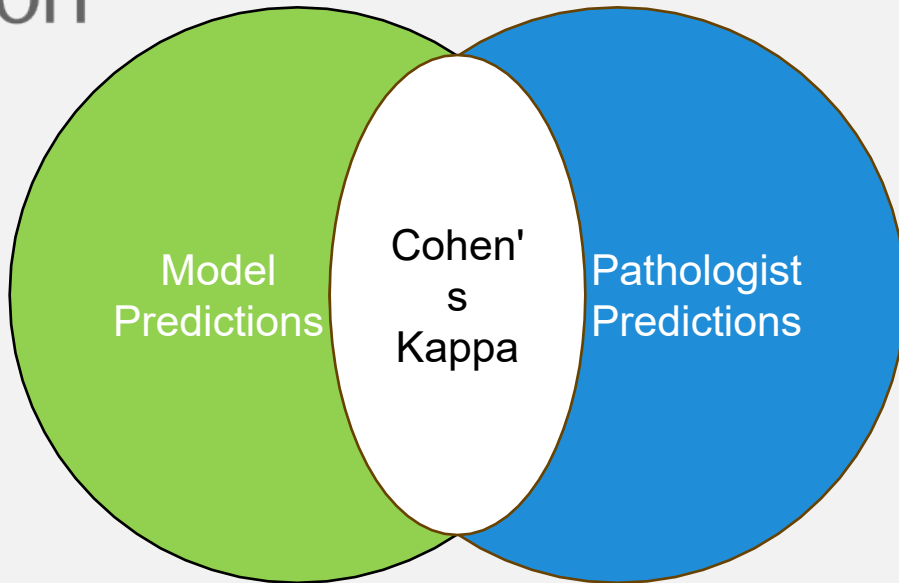
Functional GUI

- Show location of areas
- Show area's probability of malignancy

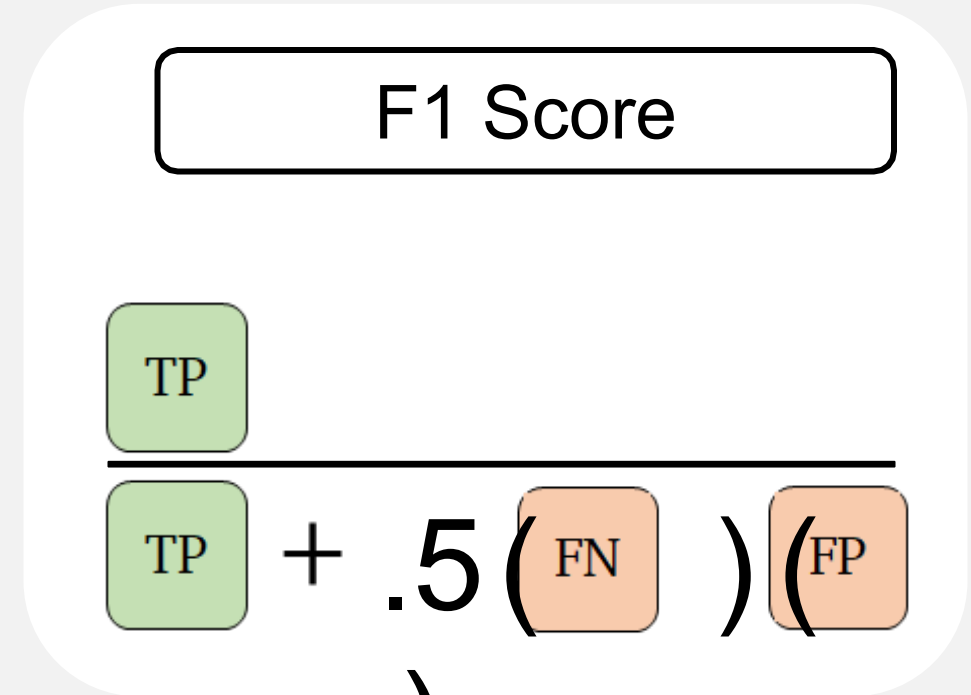


Train Time < 3 Days

# DESIGN CRITERIA / REQUIREMENTS



Cohen's Kappa > 60%  
[Source](#)



F1 Score > 90%  
[Source](#)

# DEEPER LOOK AT F1 SCORE

Actual

Classes



Predicted

Classes



F1 Score

TP

TP

+

.5

( FN )

( FP )

)

Apples

True Positive = 4

False Negative = 0

False Positive = 1

F1 Score = 89%

Bananas

True Positive = 0

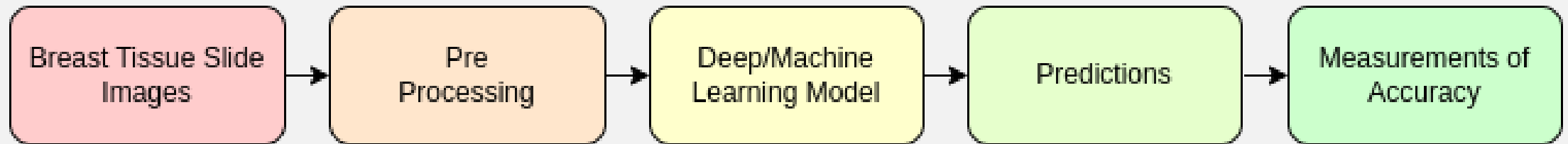
False Negative = 1

False Positive = 5

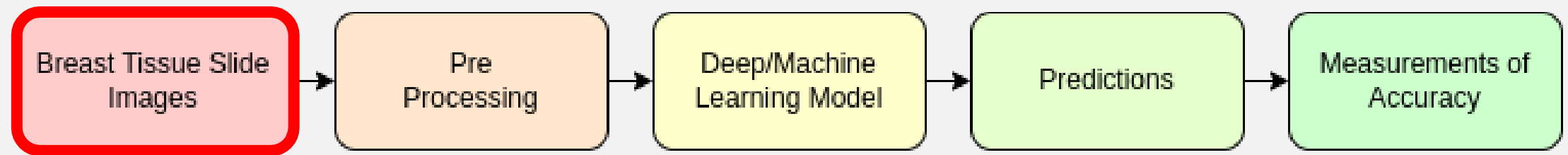
F1 Score = 0%



# PRELIMINARY DESIGN



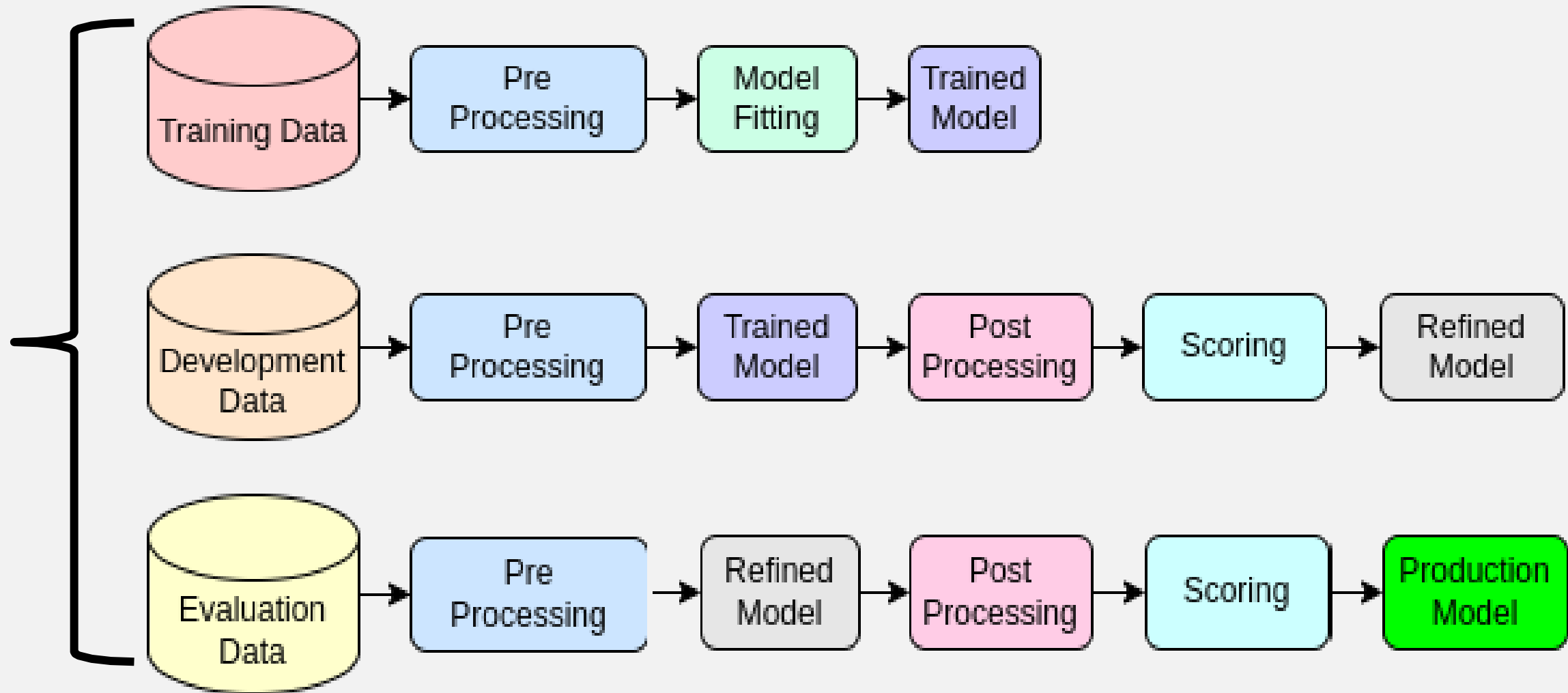
# PRELIMINARY DESIGN



# TRAINING, TUNING, EVALUATION PIPELINES

3,505 Tissue Images

1.23 Terabytes



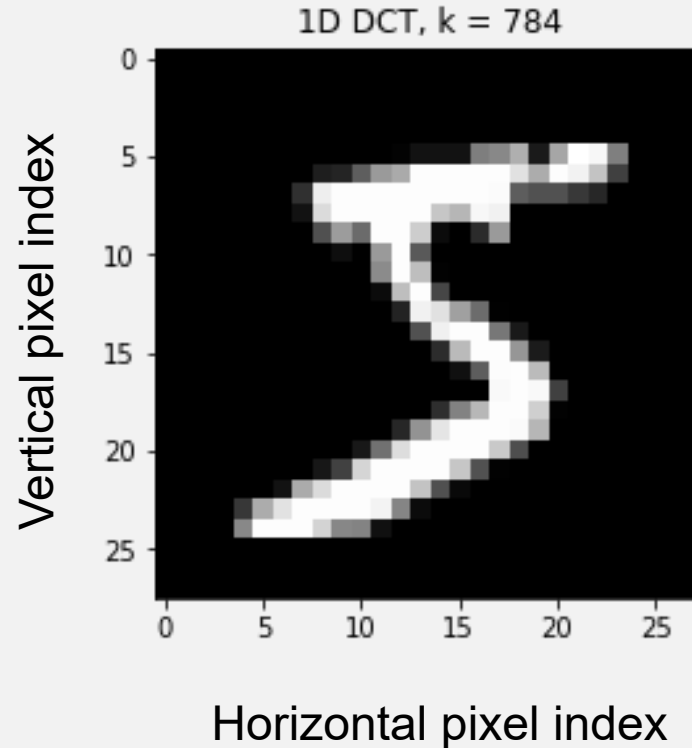
# PRELIMINARY DESIGN



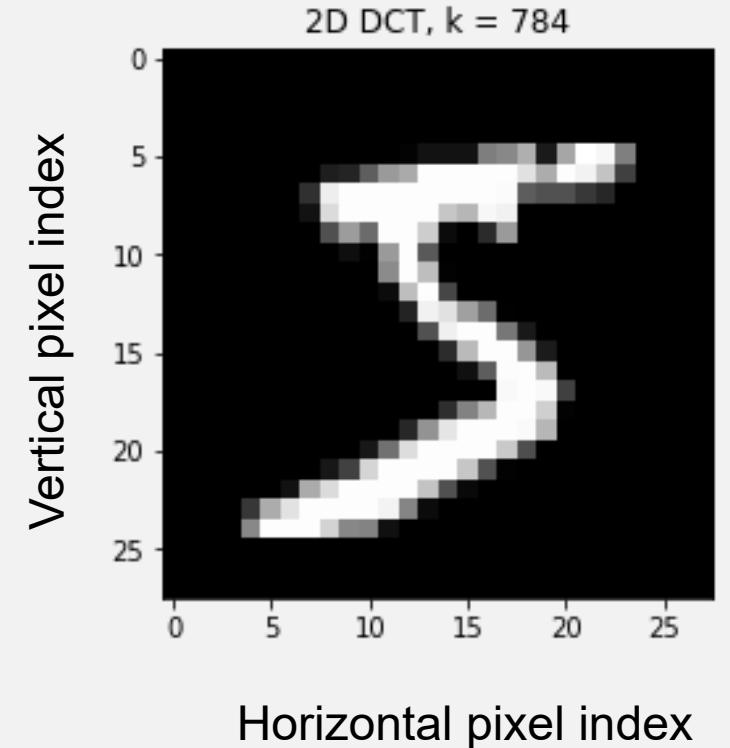
# 2-DIMENSIONAL DCT

- Belongs in the **pre-processor** stage
- Converts colors to frequencies
- Allows us to retain **fewer features**

One-dimensional DCT



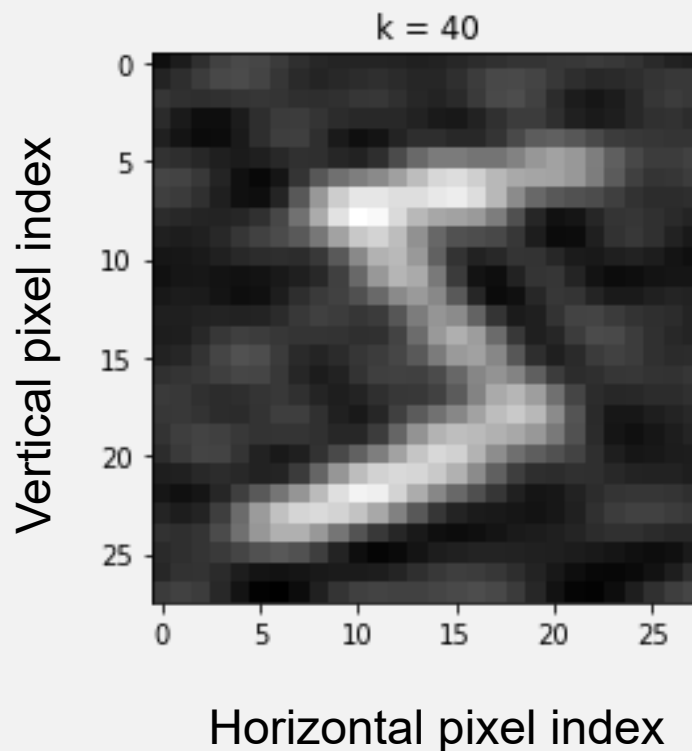
Two-dimensional DCT



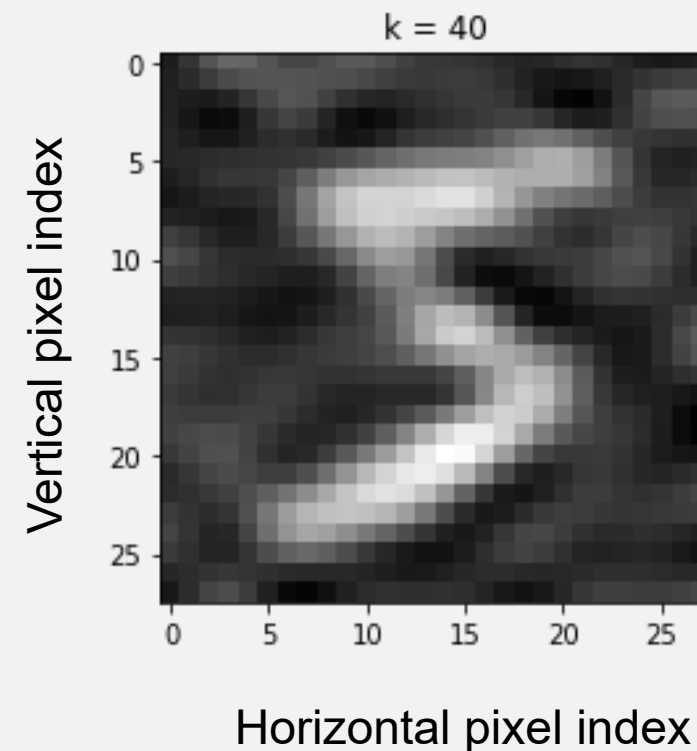
## 2-DIMENSIONAL DCT

- Belongs in the **pre-processor** stage
- Converts colors to frequencies
- Allows us to retain **fewer features**

One-dimensional DCT



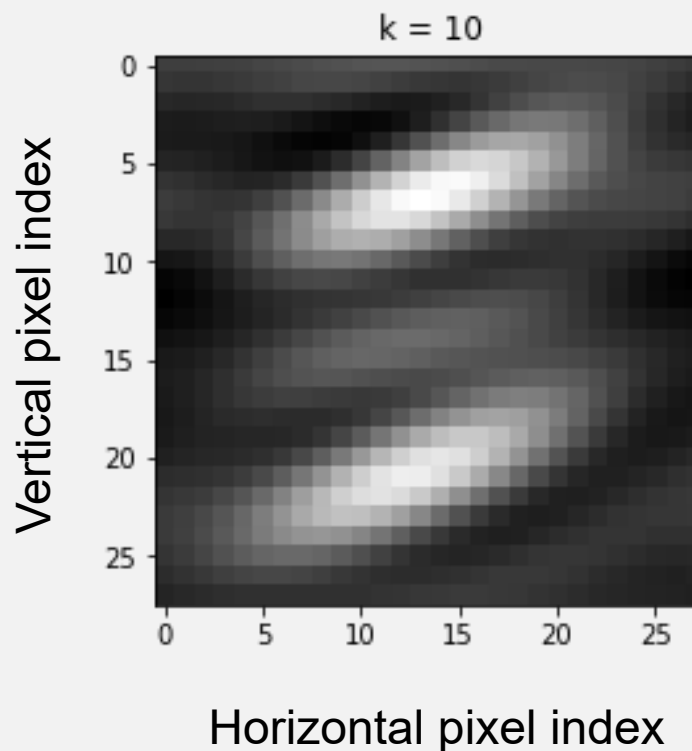
Two-dimensional DCT



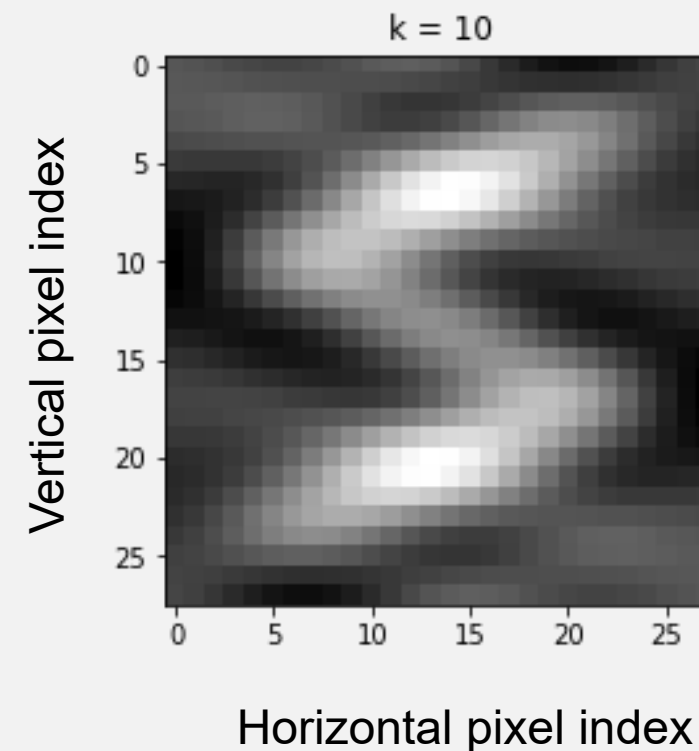
## 2-DIMENSIONAL DCT

- Belongs in the **pre-processor** stage
- Converts colors to frequencies
- Allows us to retain **fewer features**

One-dimensional DCT



Two-dimensional DCT



# PRINCIPAL COMPONENT ANALYSIS (PCA)

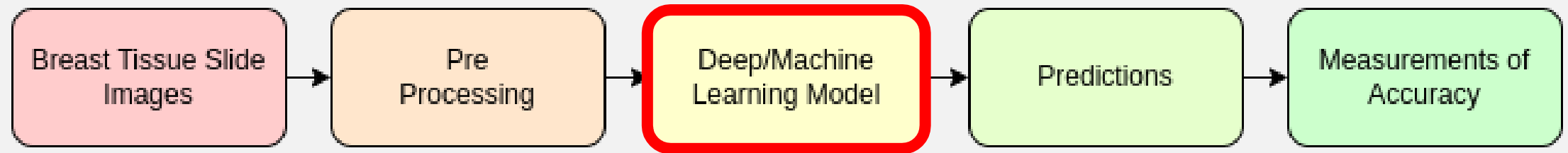
f1	f2	f4	f7
f3	f5	f8	f11
f6	f9	f12	f14
f10	f13	f15	f16



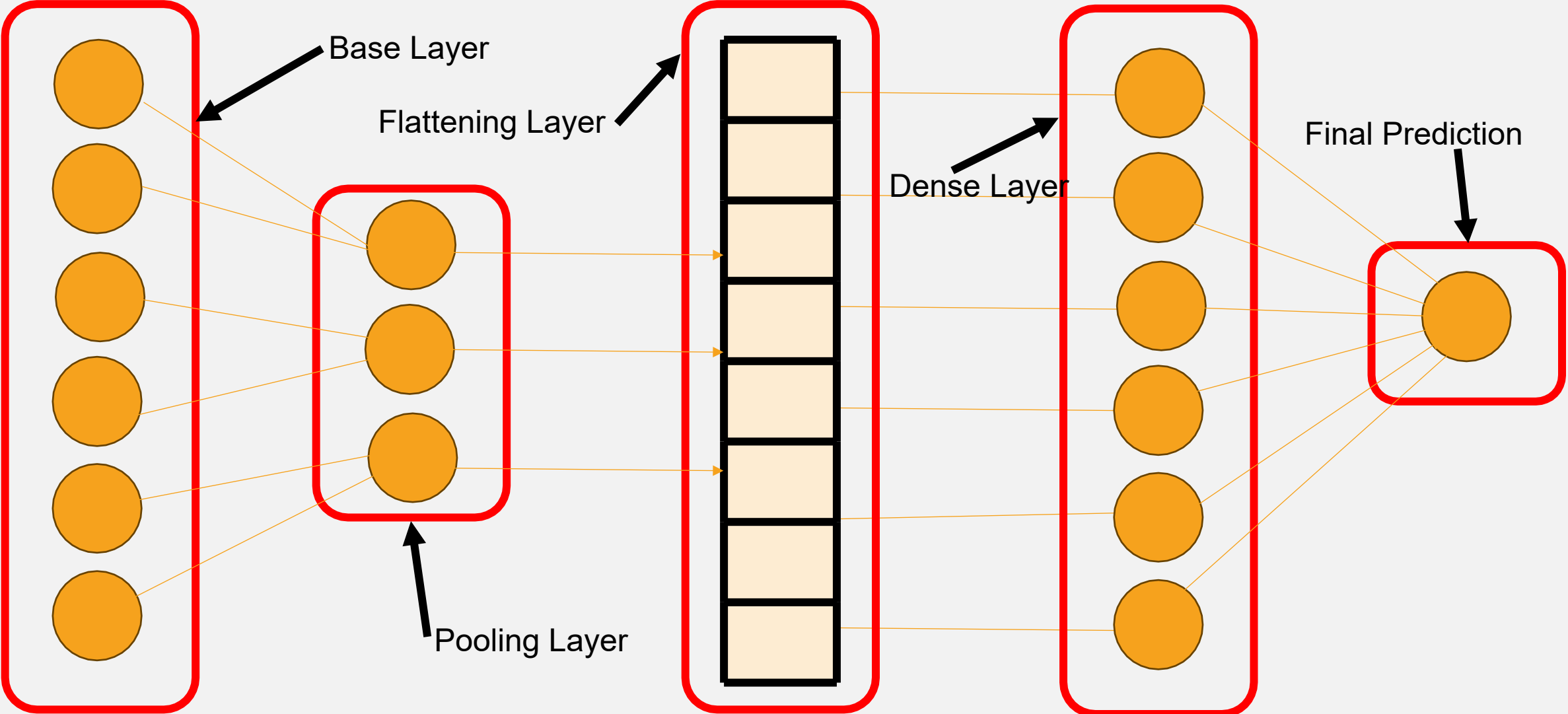
PC1	PC2	PC4	X
PC3	PC5	X	X
PC6	X	X	X
X	X	X	X



# PRELIMINARY DESIGN

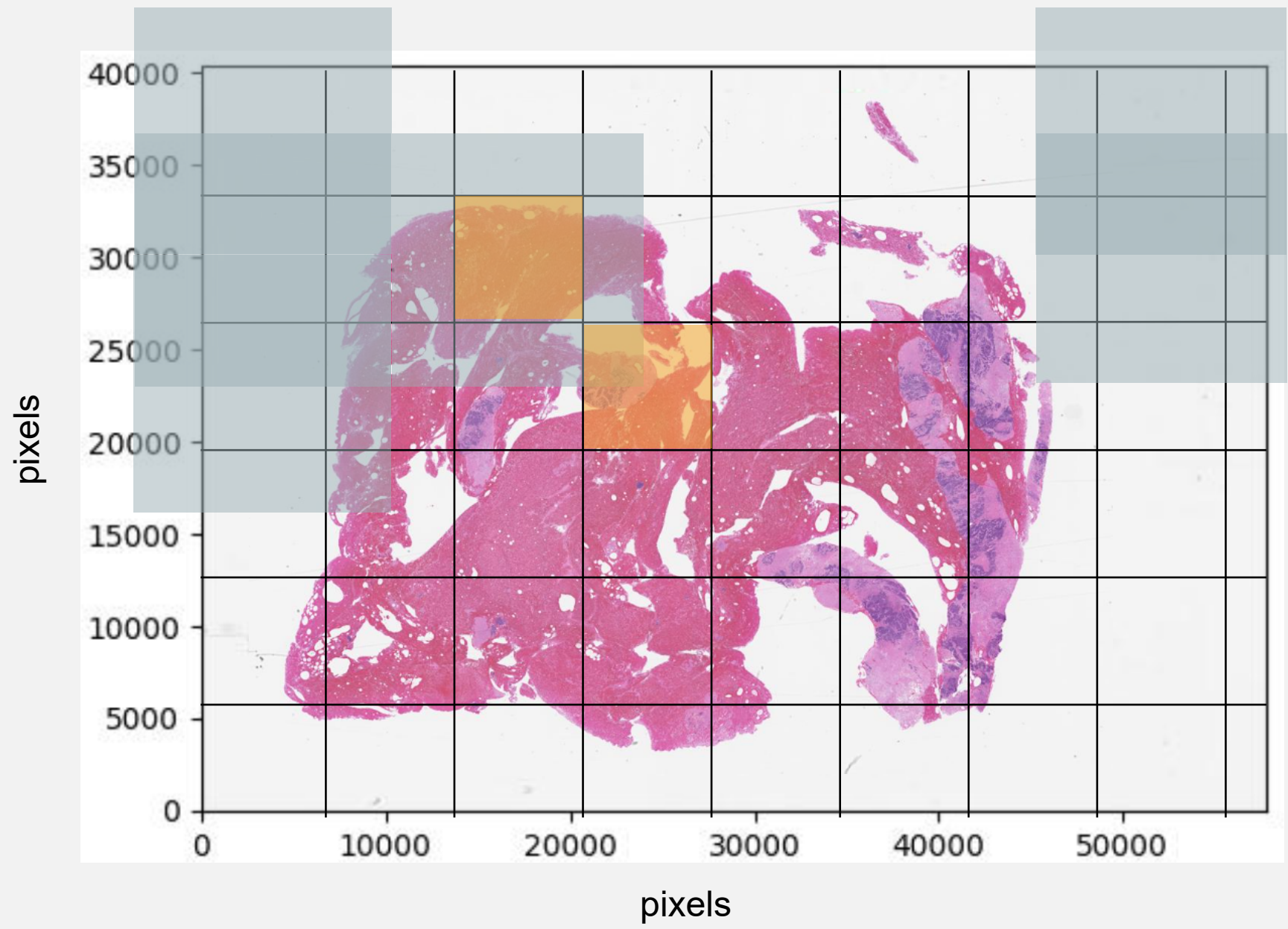


# CONVOLUTIONAL NEURAL NETWORK

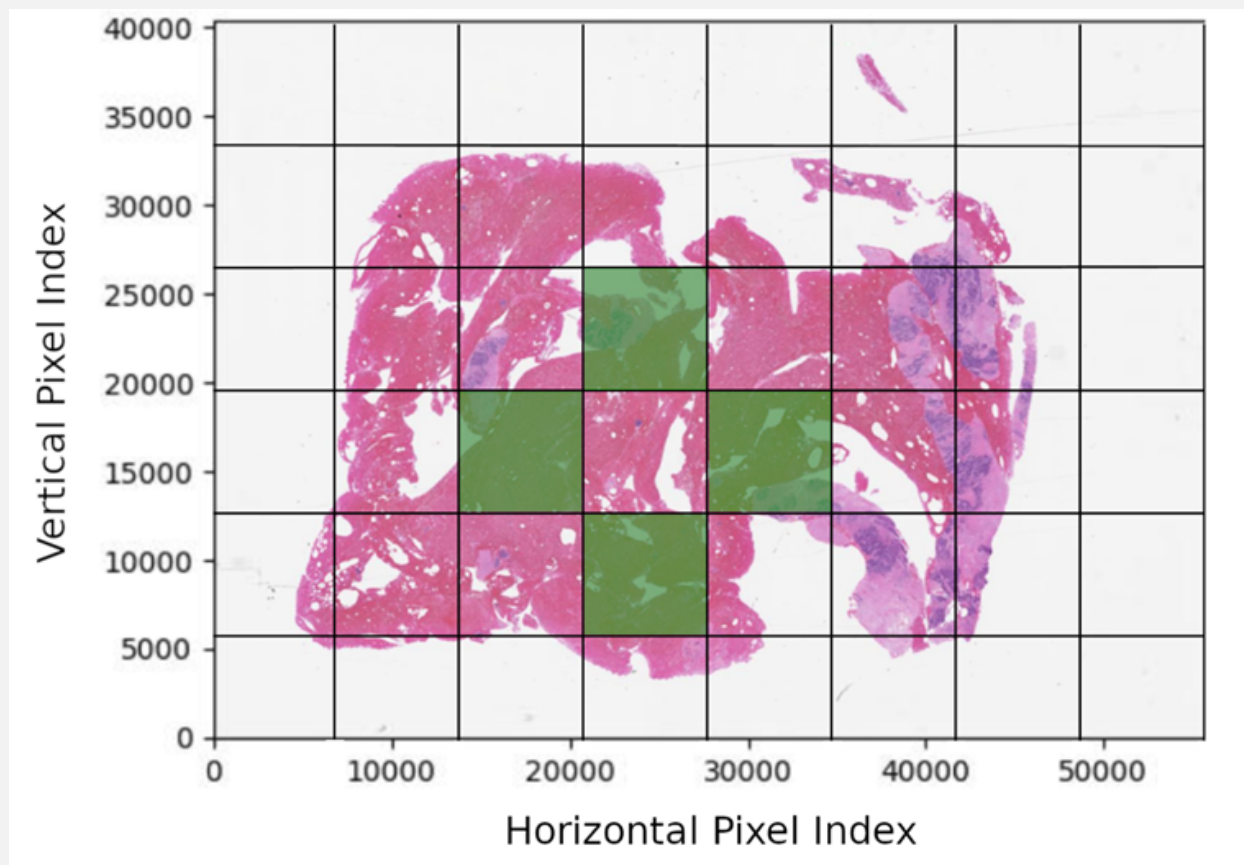


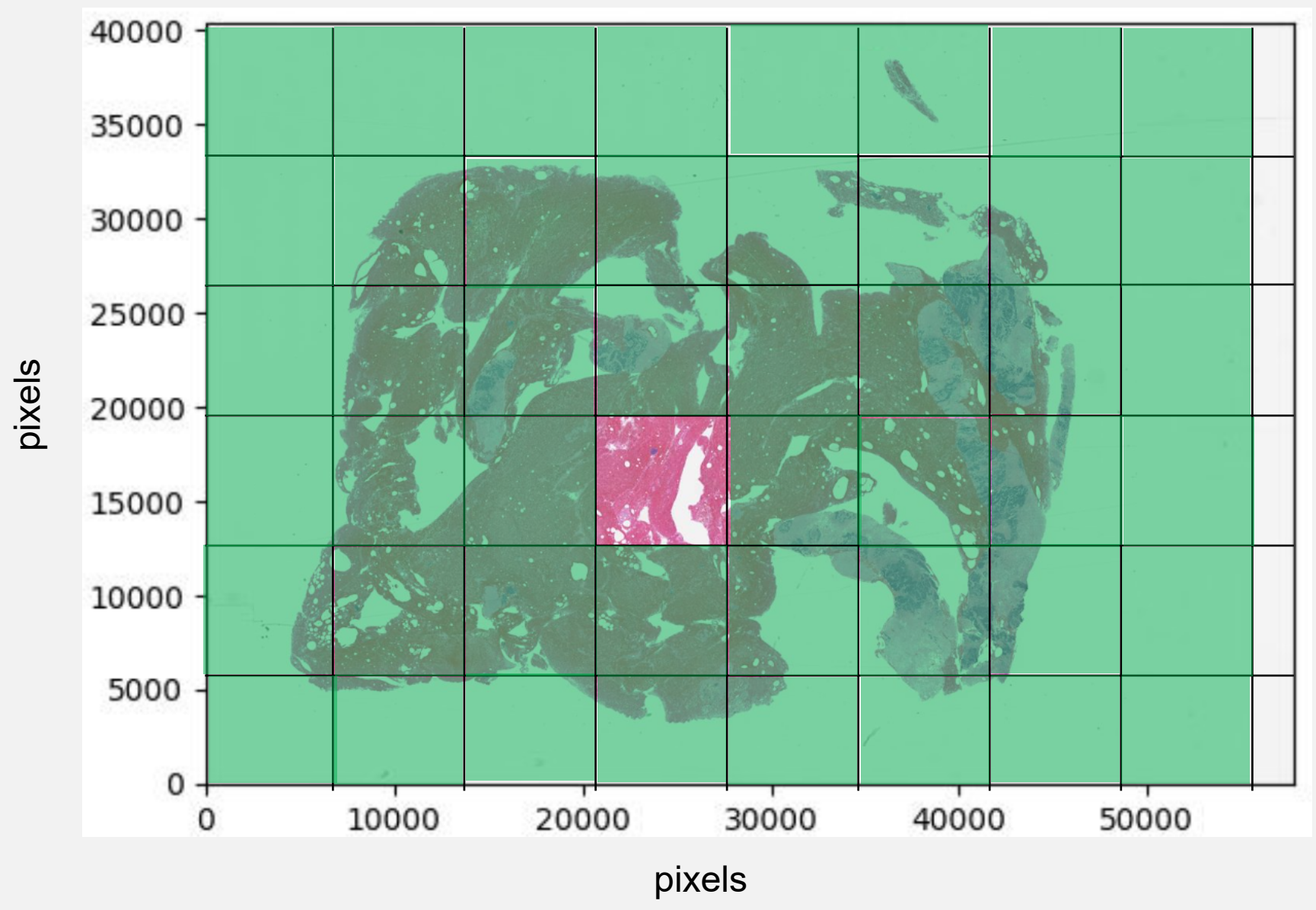
# PRELIMINARY DESIGN



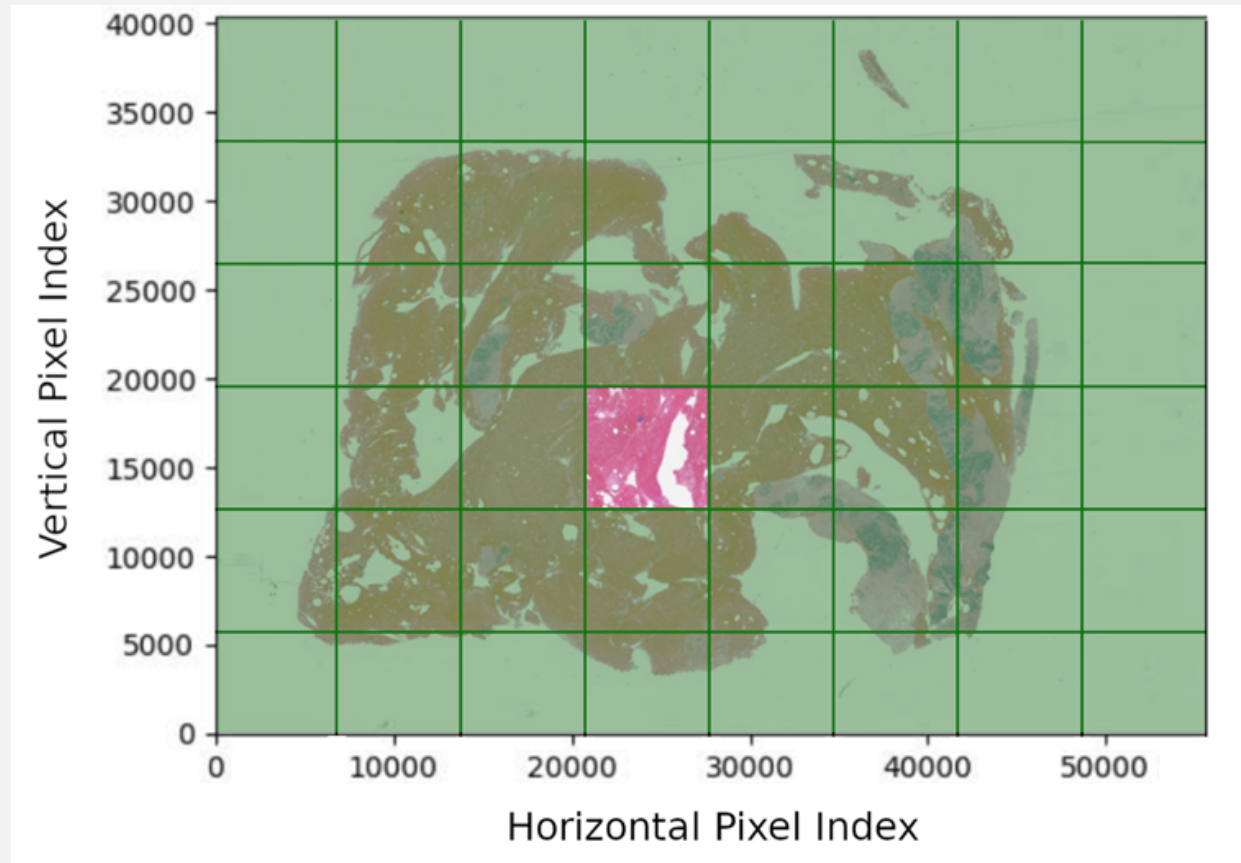


# FRAME TO PATCH

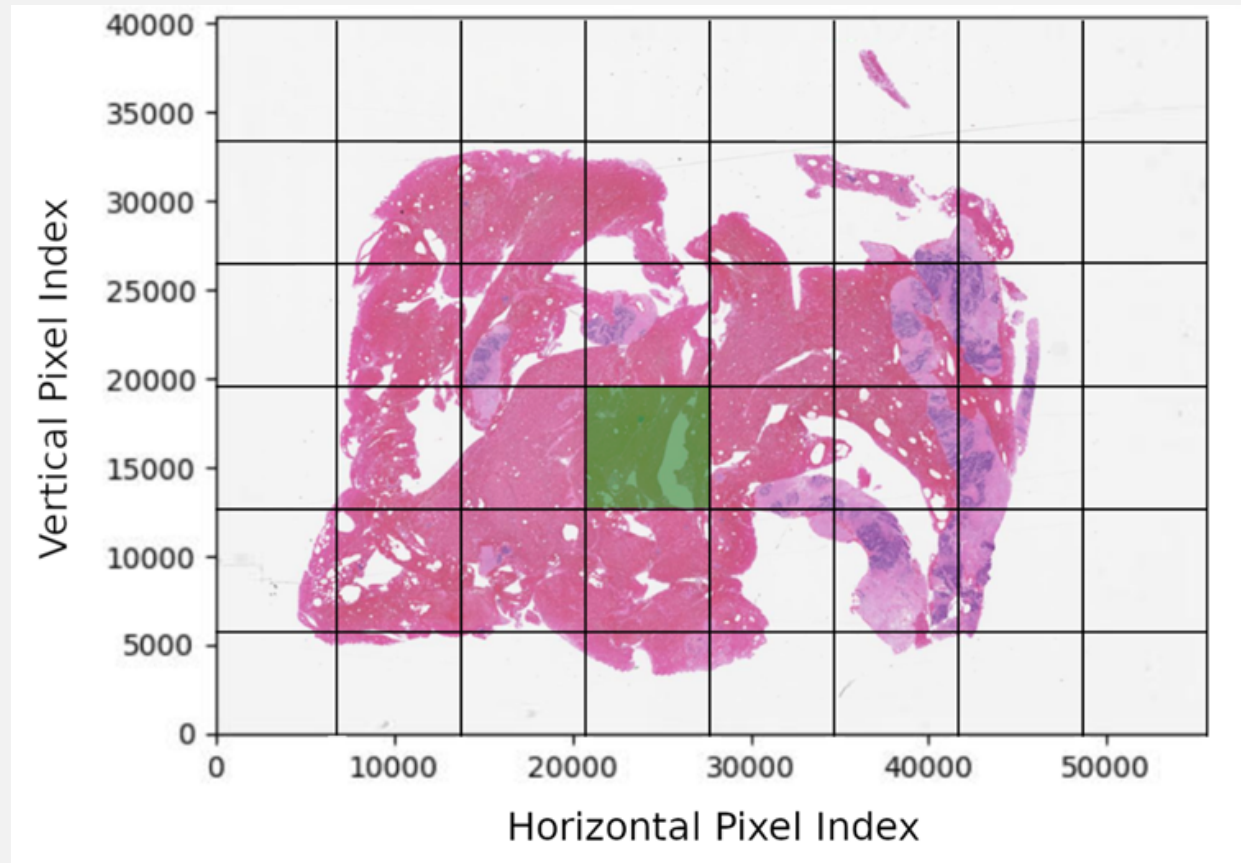




# WINDOW TO PATCH

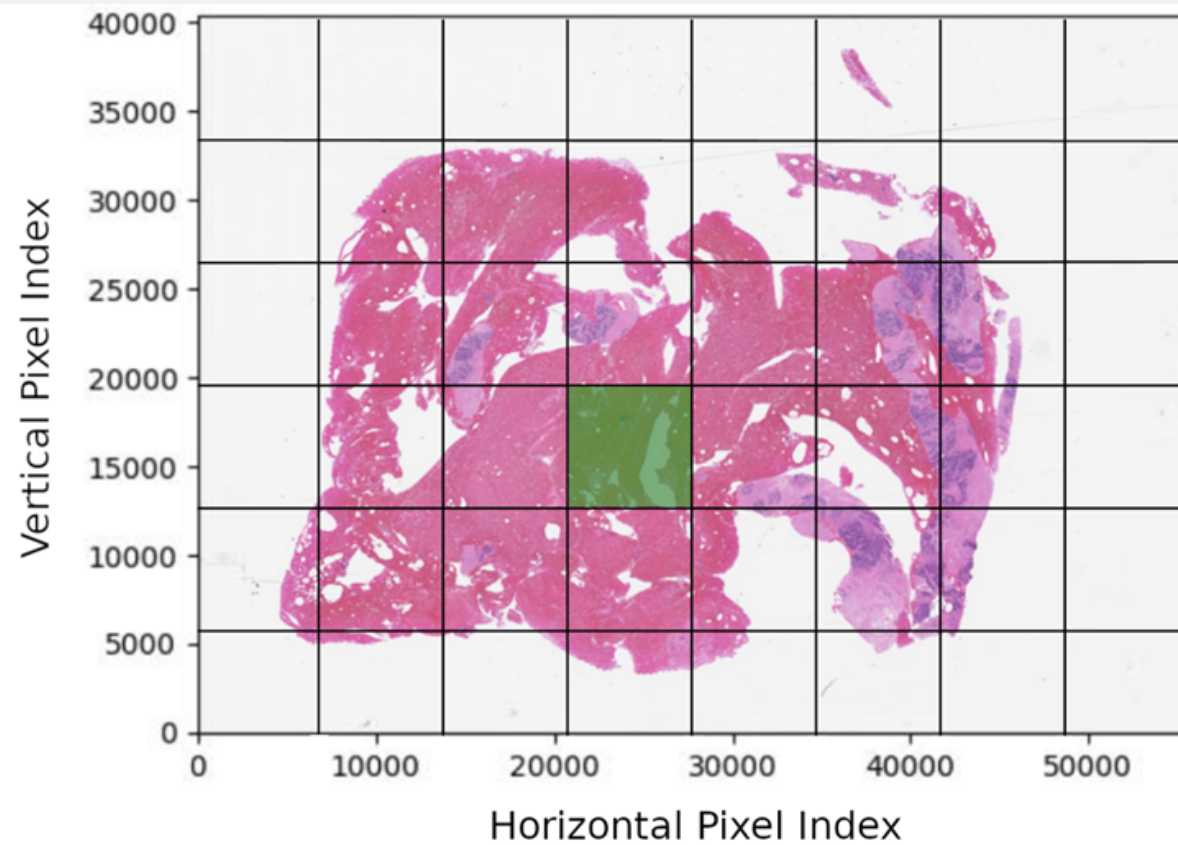
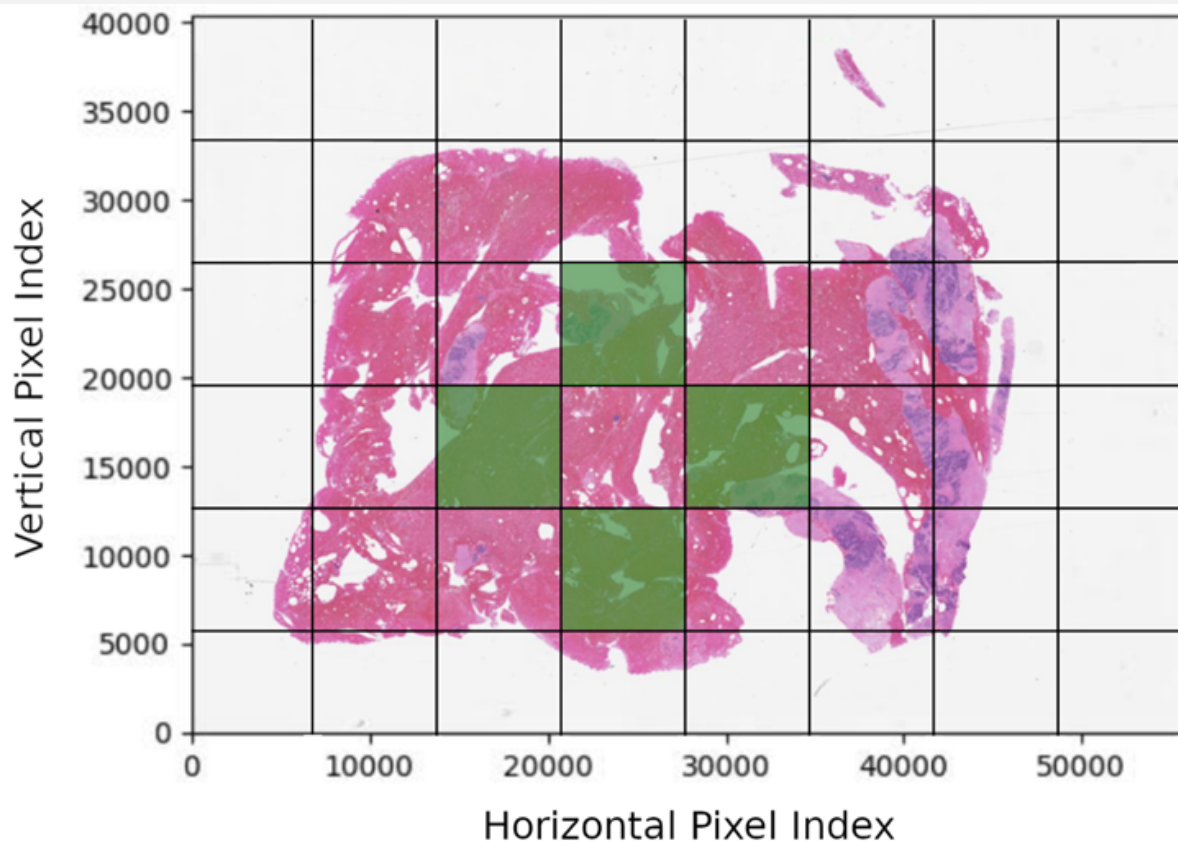


# FRAME TO PATCH

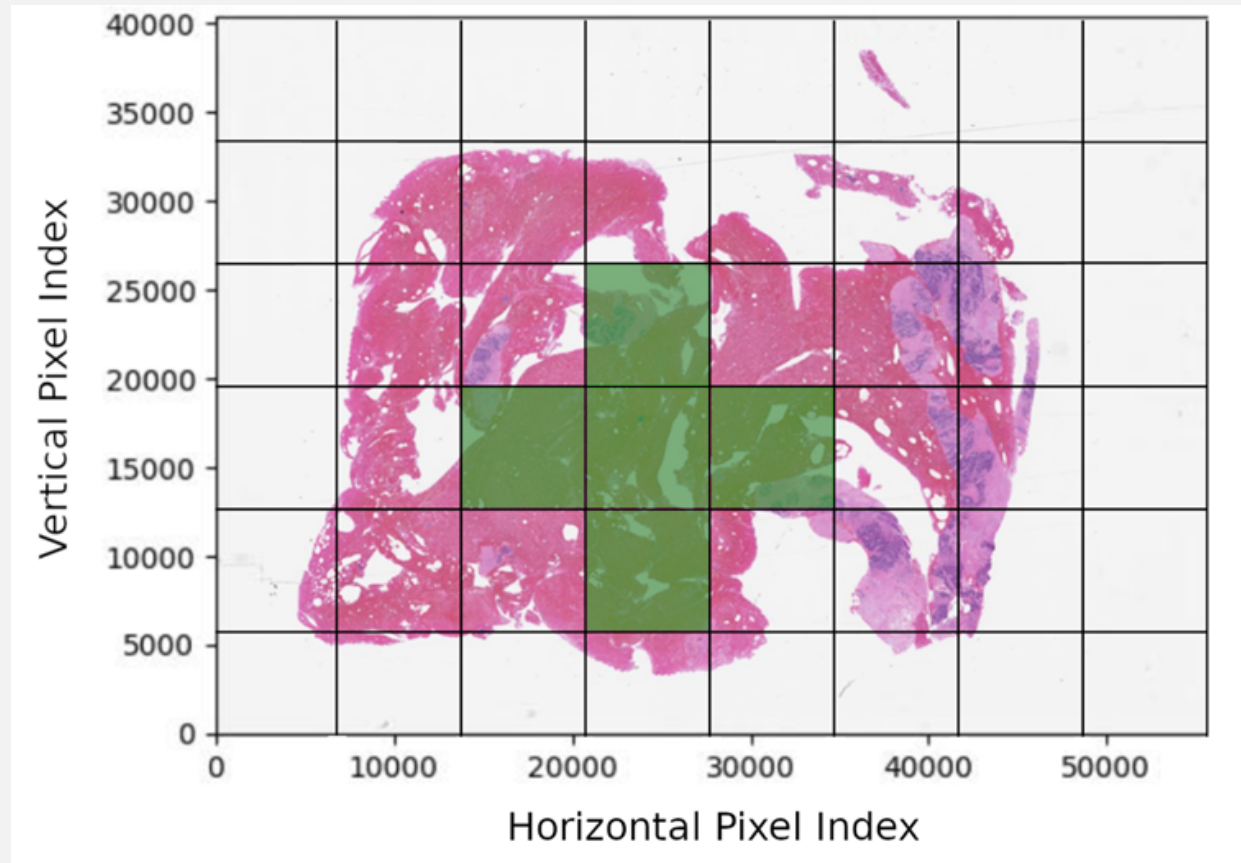




# FRAME TO PATCH

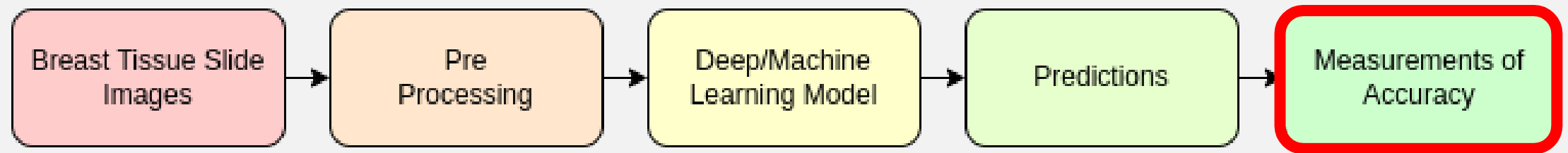


# FRAME TO PATCH

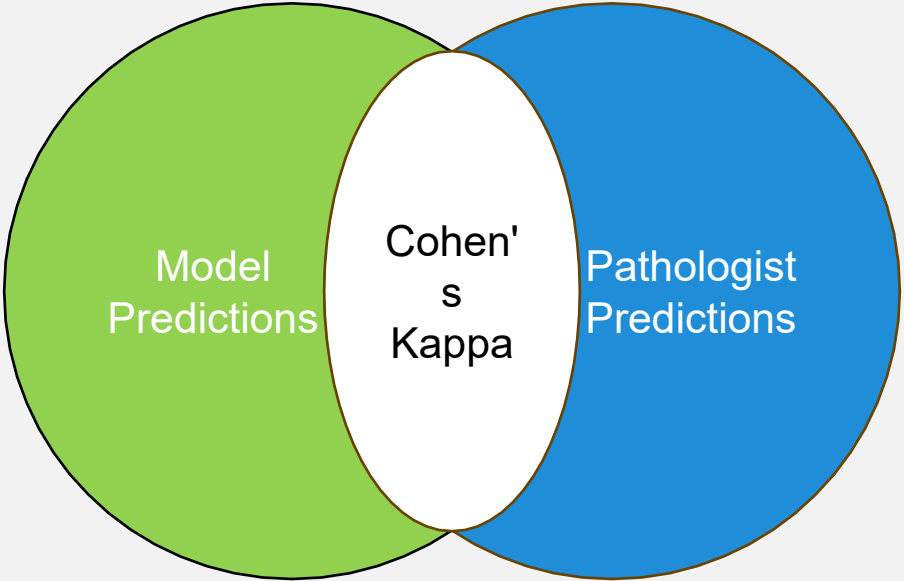


PATCH TO IMAGE LEVEL

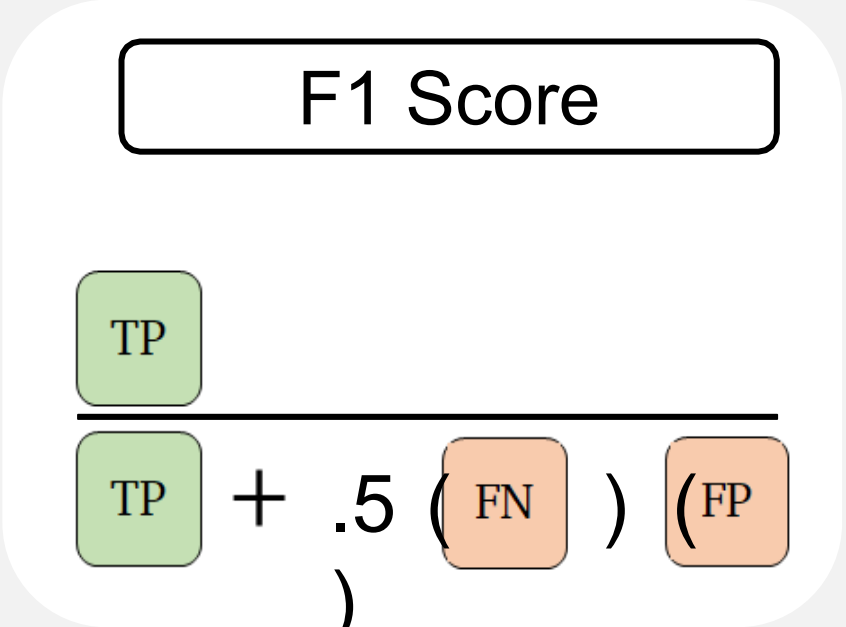
# PRELIMINARY DESIGN



# FUTURE WORK



Cohen's Kappa > 60%  
[Source](#)



F1 Score > 90%  
[Source](#)



No Sobel/Laplacian Filters

# QUESTIONS?

*Acknowledgements:*

*Dr. Joseph Picone – Mentor & Data Coordinator*

*Claudia Dumitrescu – AI Expert*

*Phuykong Meng – GUI Planning & Scoring*

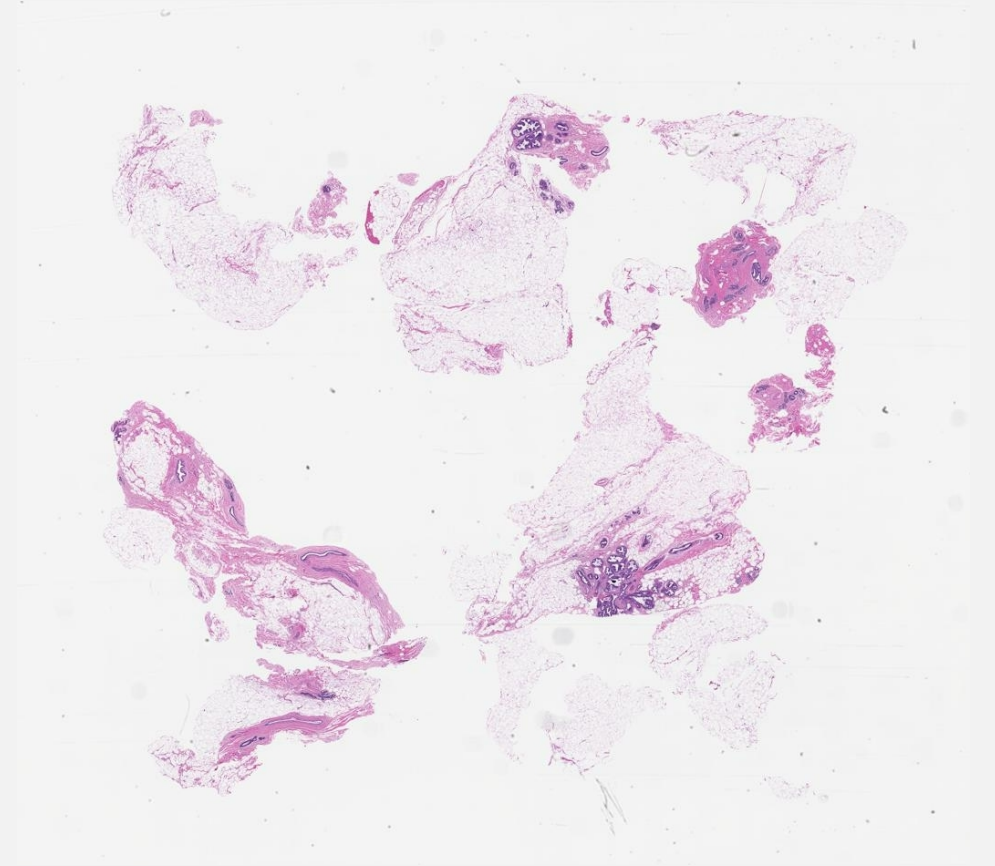
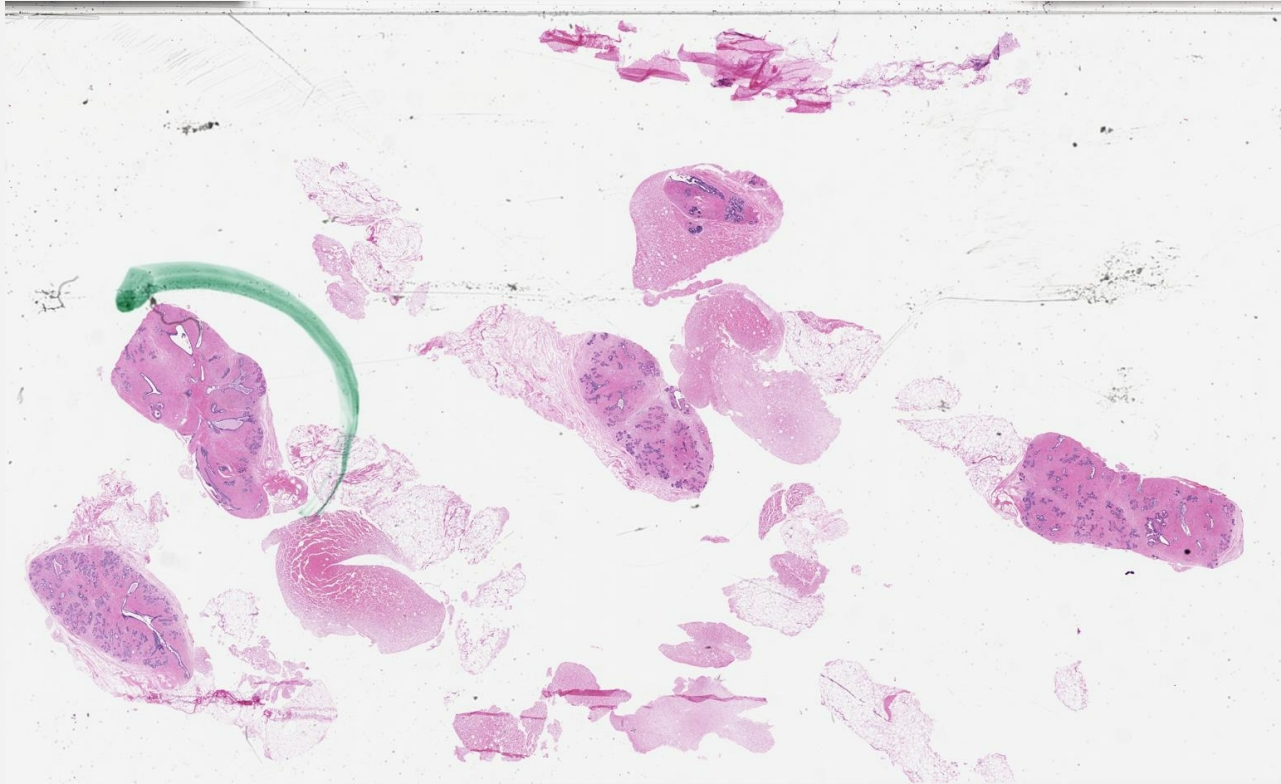
*For the curious*

- [\*Benefits of Machine Learning in Healthcare\*](#)
- [\*Machine Learning in Healthcare\*](#)
- [\*What is Machine Learning in Healthcare?\*](#)
- [\*Significance of Machine Learning in Healthcare\*](#)
- [\*The Potential for Artificial Intelligence in Healthcare\*](#)

## PROGRESS SINCE LAST PRESENTATION

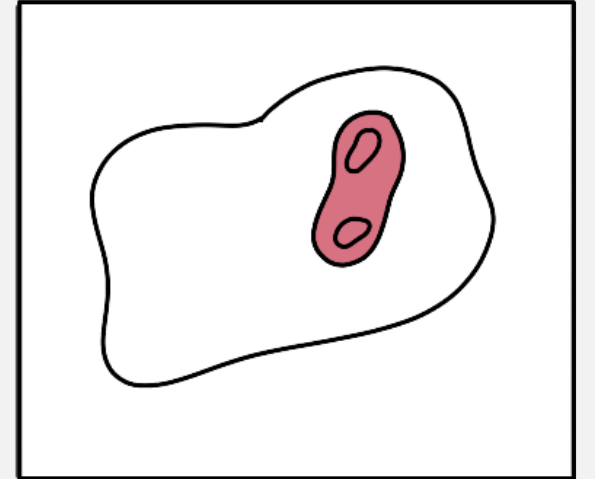
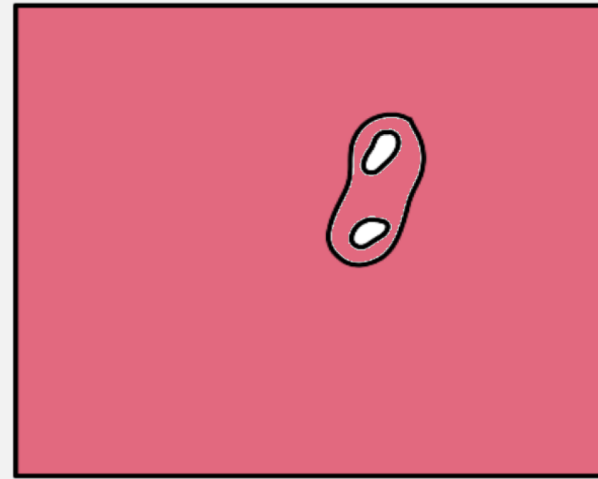
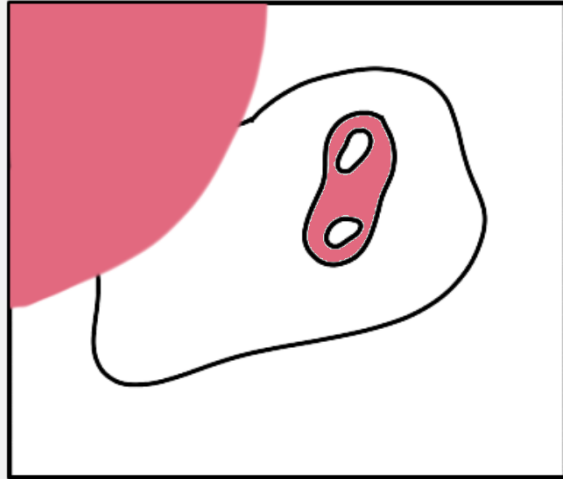
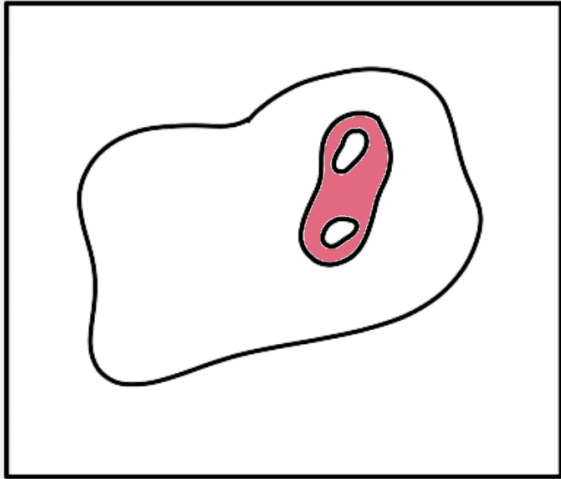
- 2D Discrete Cosine Transform (DCT)
- Principal Component Analysis (PCA)
- Convolutional Neural Network (CNN)

# TUHDP BIOPSY SLIDE SAMPLE

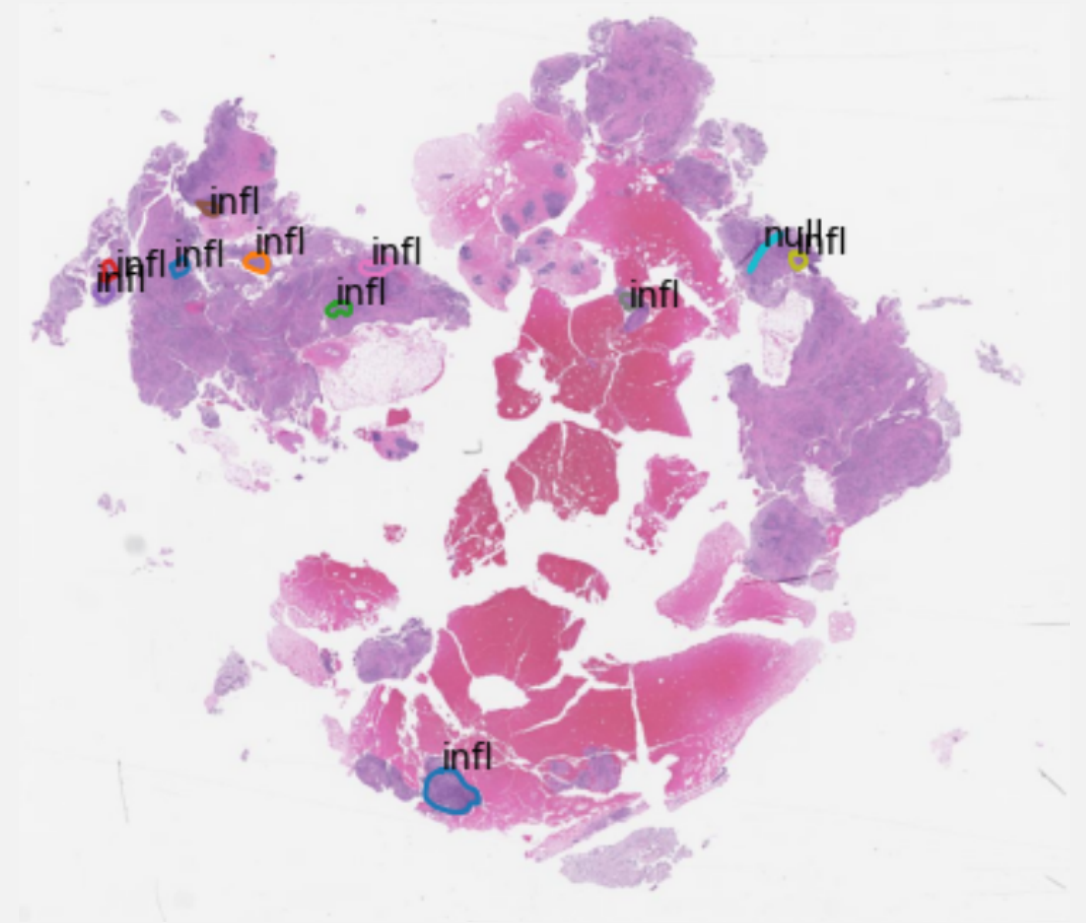
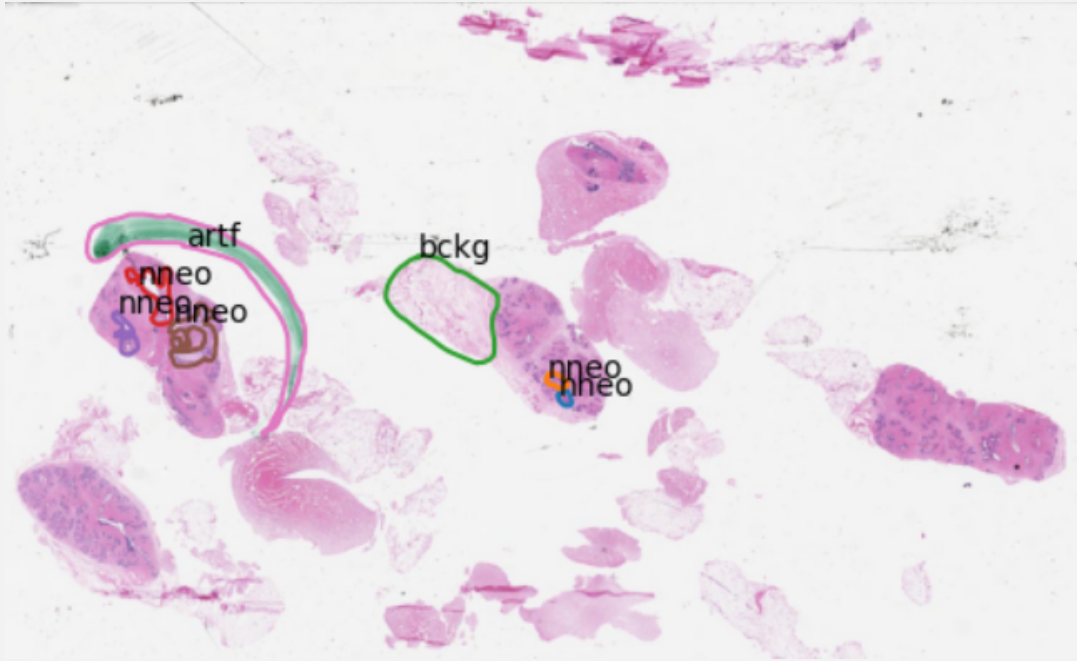




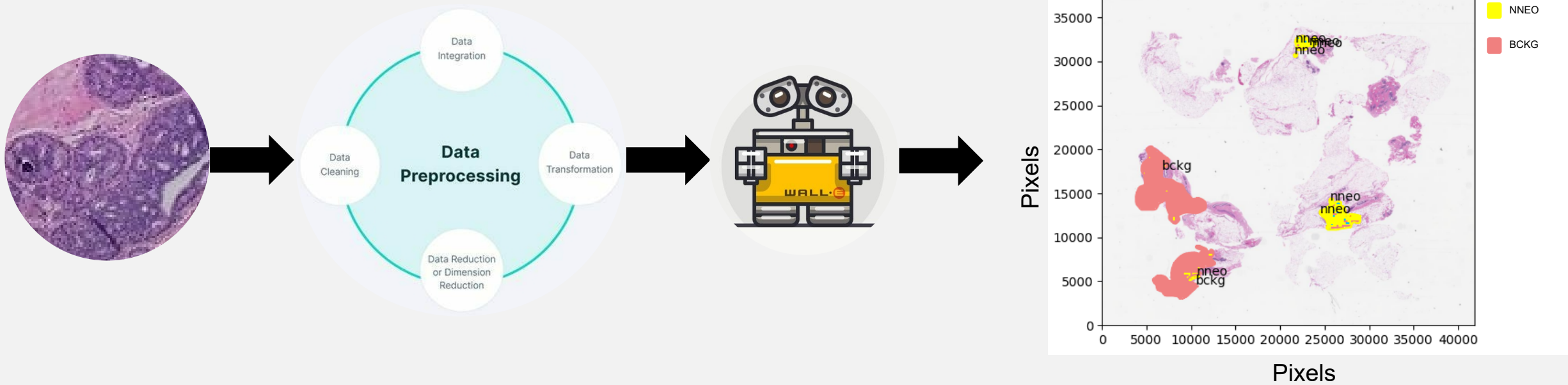
# WINDOW TO PATCH



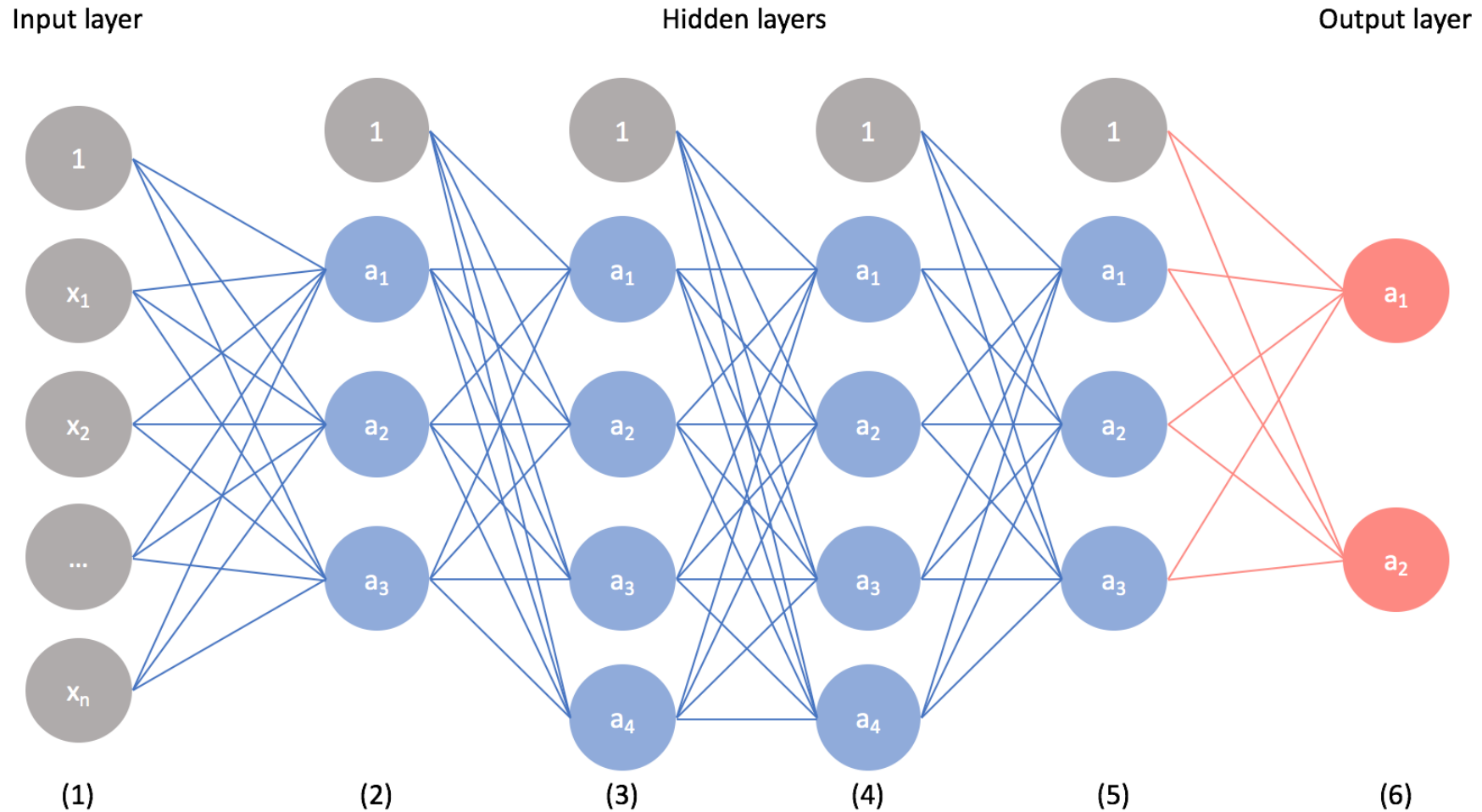
# TUHDP BIOPSY SLIDE SAMPLE



# DEEPER LOOK AT MODEL I/O



# CONVOLUTIONAL NEURAL NETWORK

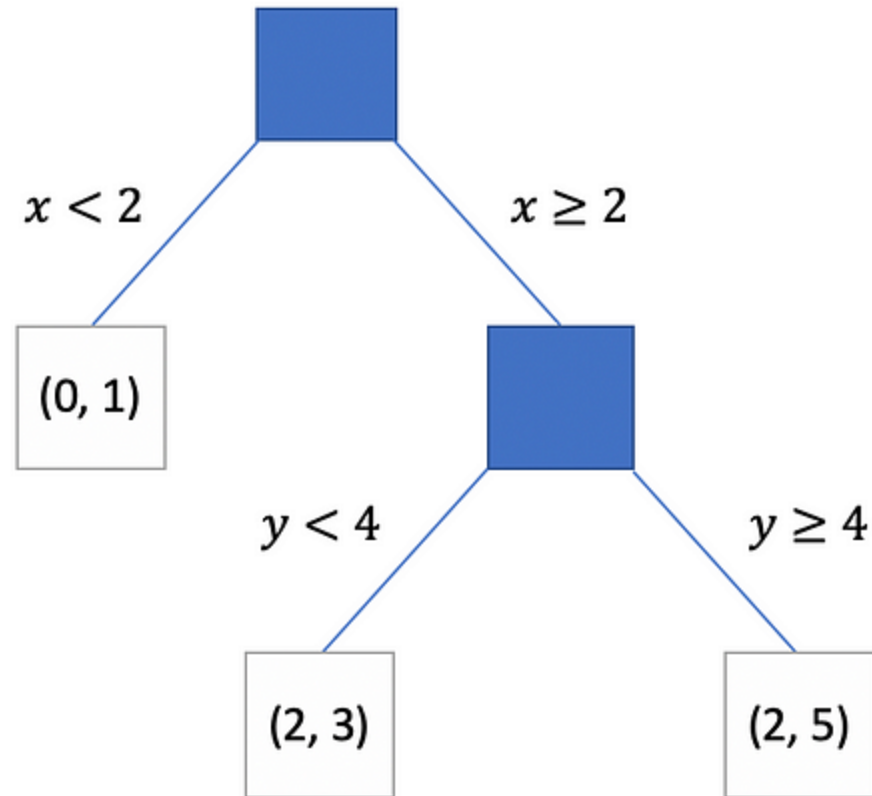


# RANDOM FOREST

Dataset

x	y
0	1
2	3
2	5

Isolation Tree



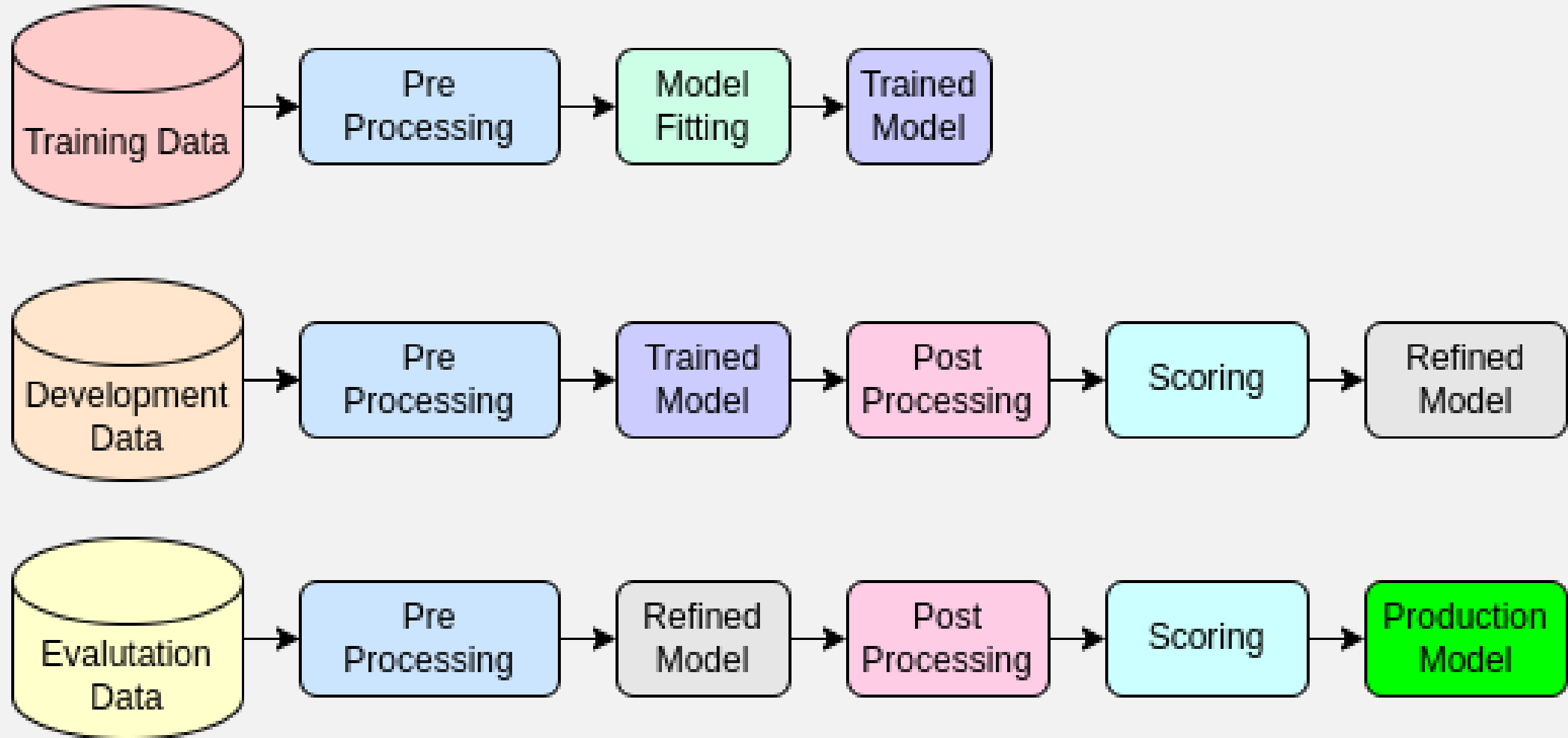
# WHAT IS A MODEL?

- Three essential components for machine learning
  - Data
  - Model
  - Training algorithm
- Model
  - **Mathematical representation** of relationships in the data
- Training algorithm
  - Uses data
  - Adjusts variables *in the model* until **output matches input**

# TEMPLE UNIVERSITY HEALTH DIGITAL PATHOLOGY CORPUS

3,505 Tissue Images

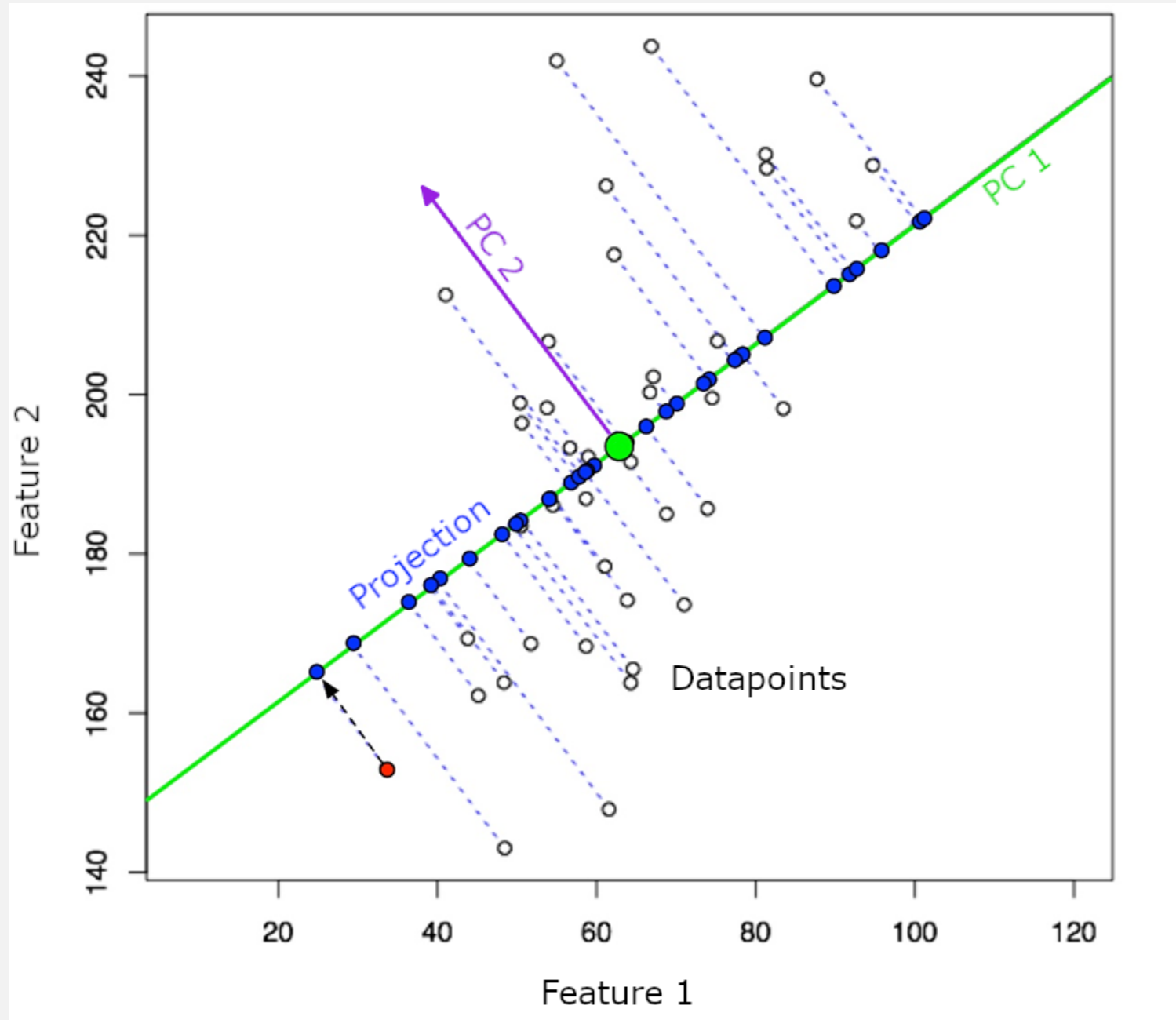
1.23 Terabytes



## PRINCIPAL COMPONENT ANALYSIS (DIMENSION REDUCTION)

- *After feature generation*
- Also belongs in the **pre-processor** stage after digitizing/segmenting slides and before training
- Reduces the number of features while minimizing information lost from feature reduction
- **Fewer features** compared to no PCA or dimension reduction





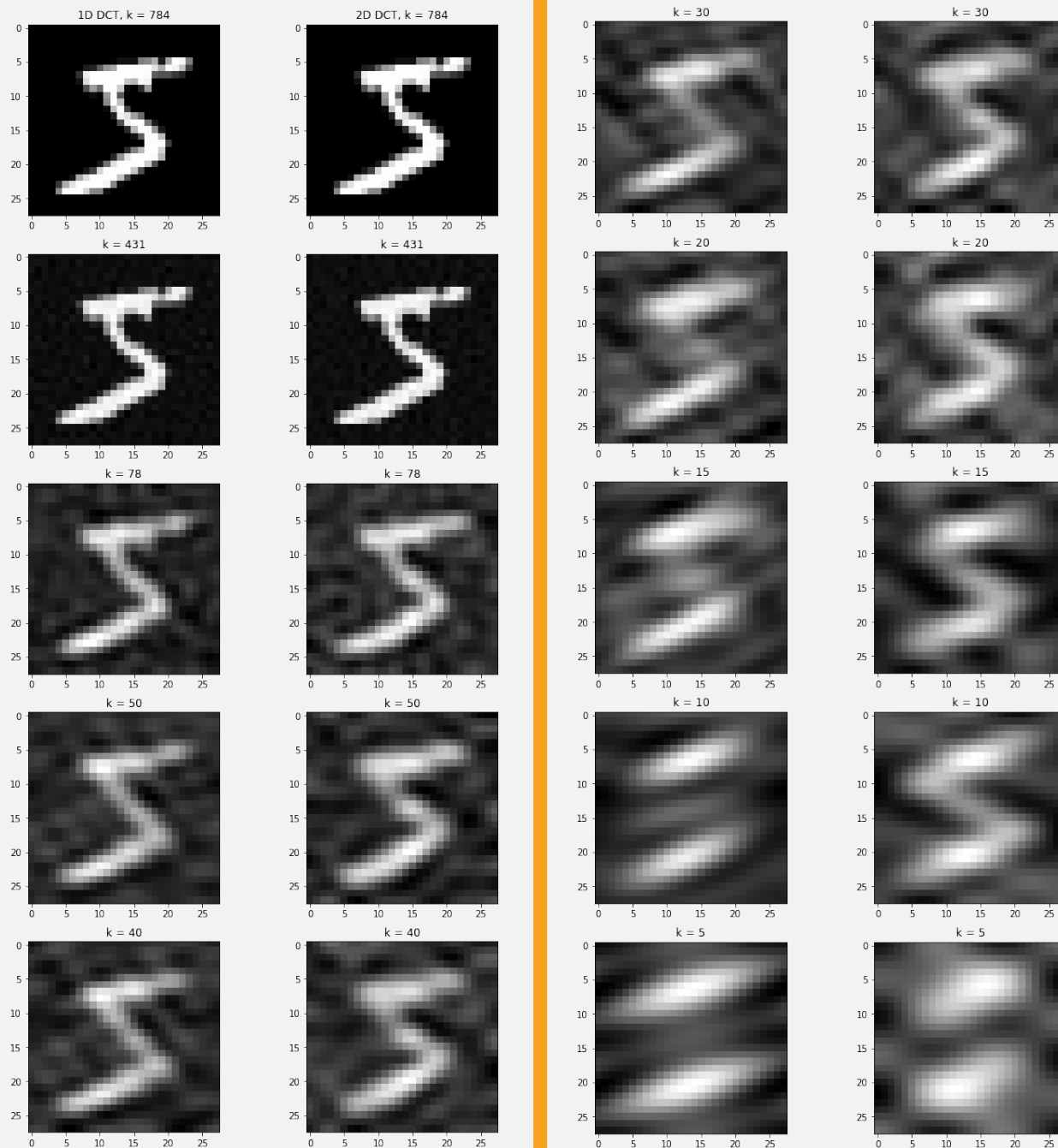
# TESTING GOALS

## 2-DIMENSIONAL DCT

- Part of **feature generation**
- Belongs in the **pre-processor** stage after digitizing/segmenting slides and before training
- Greater spectral/energy density than one-dimensional DCT
- Therefore, **fewer features** compared to one-dimensional DCT

# 2-DIMENSIONAL DCT

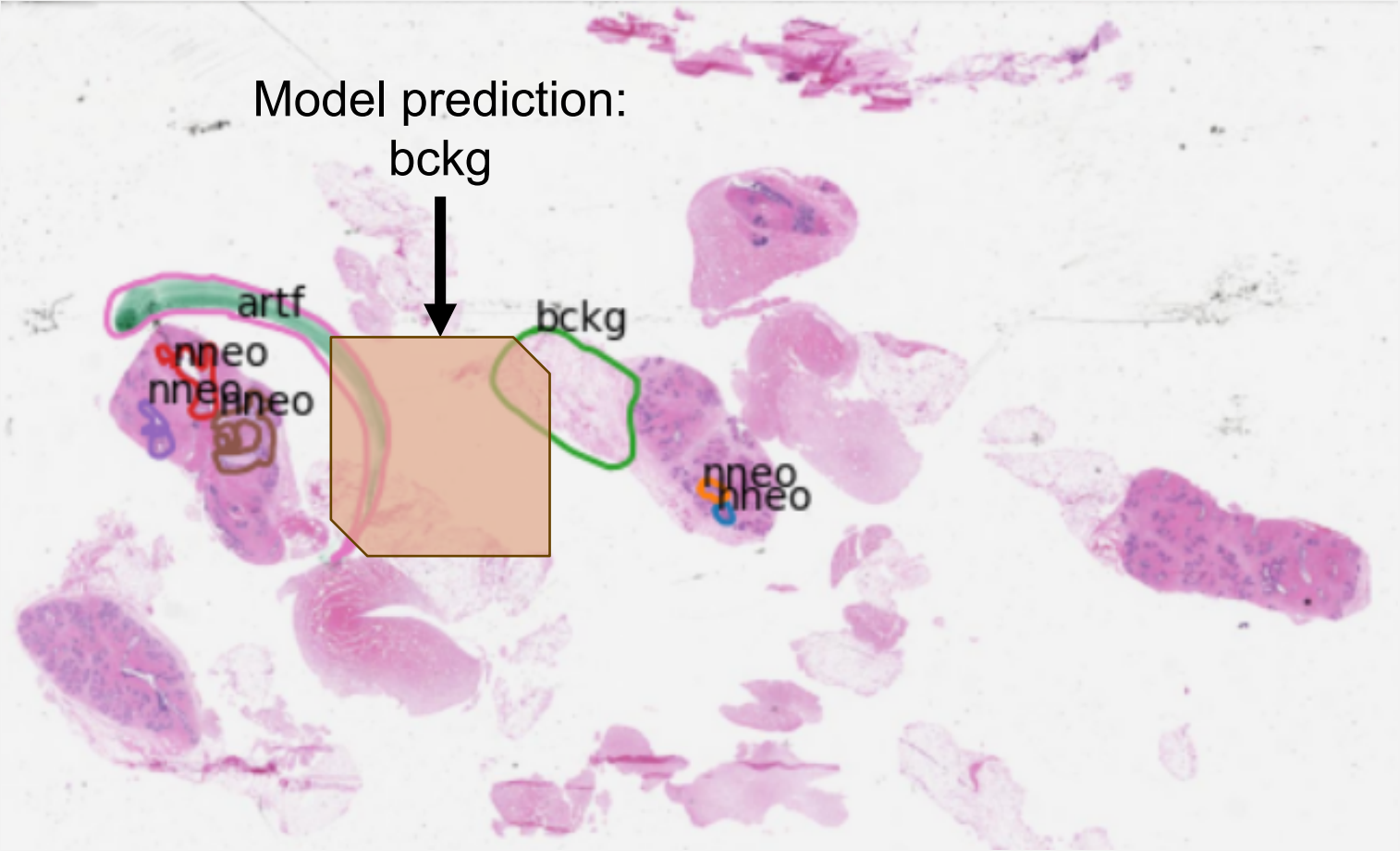
- Belongs in the **pre-processor** stage
- Converts colors to frequencies
- Allows us to retain **fewer features**



## SCORING

- Find **overlapping area** between the model's predictions and the human pathologist's predictions
- Use overlapping area to construct a **confusion matrix**
- Use confusion matrix to generate  **$F_1$  scores and Cohen's kappa coefficients**

# OVERLAPPING AREA



Frame Labels	Pathologist (True)										Sums across rows
	Unlab	Bckg	Norm	Null	Artf	Nneo	Infl	Susp	Indc	Dcis	
Unlab	20	1	4	1	7	6	7	0	4	6	56
Bckg	6	30	1	3	5	2	4	7	3	3	64
Norm	3	2	29	6	0	1	1	3	1	2	48
Null	3	0	4	24	3	4	6	3	5	0	52
Artf	5	6	4	7	29	3	6	2	3	3	68
Nneo	1	6	6	1	3	21	5	3	3	5	54
Infl	6	3	7	4	0	6	26	7	1	2	62
Susp	6	4	6	4	5	3	6	22	0	6	62
Indc	5	5	2	6	0	3	6	0	28	0	55
Dcis	5	0	6	6	7	7	4	7	5	24	71
Sums across columns	60	57	69	62	59	56	71	54	53	51	592

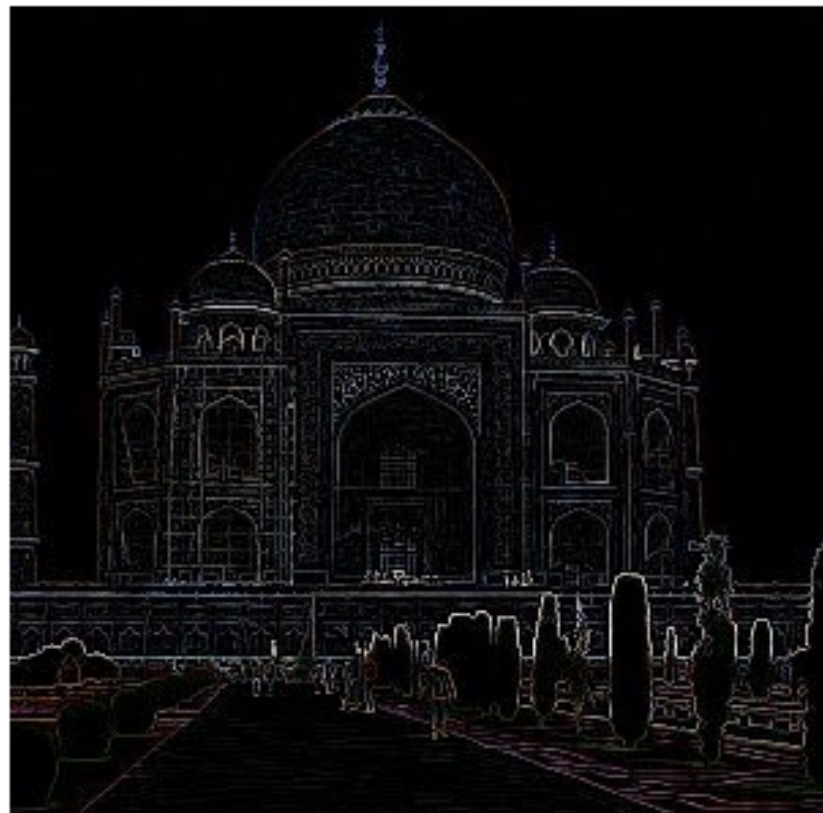


Frame Labels	Pathologist (True)		Sums across rows
	NNeo	Not NNeo	
NNeo	21	33	TP+FP = 54
Not NNeo	35	503	FN+TN = 538
Sums across columns	TP+FN = 56	FP+TN = 536	592

# FILTERS



Original image



Filter "Laplace" applied

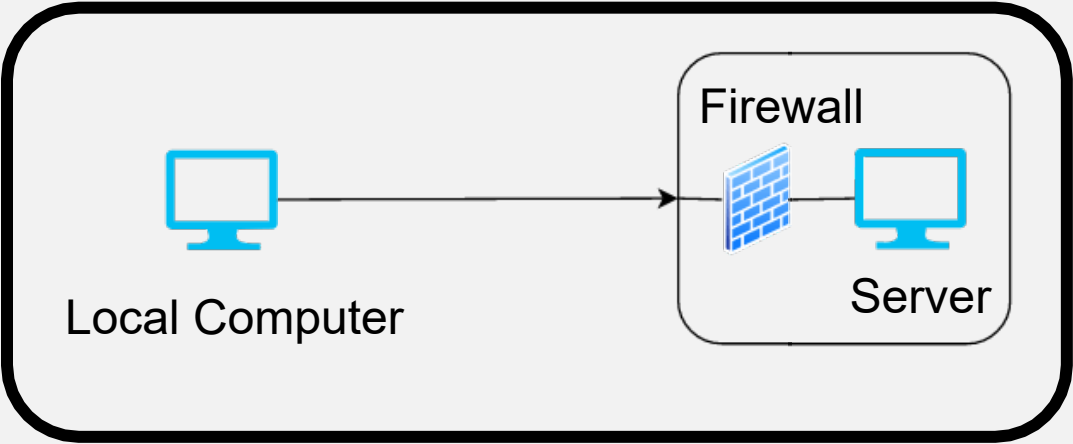


# FRAME LEVEL EVALUATION RESULTS

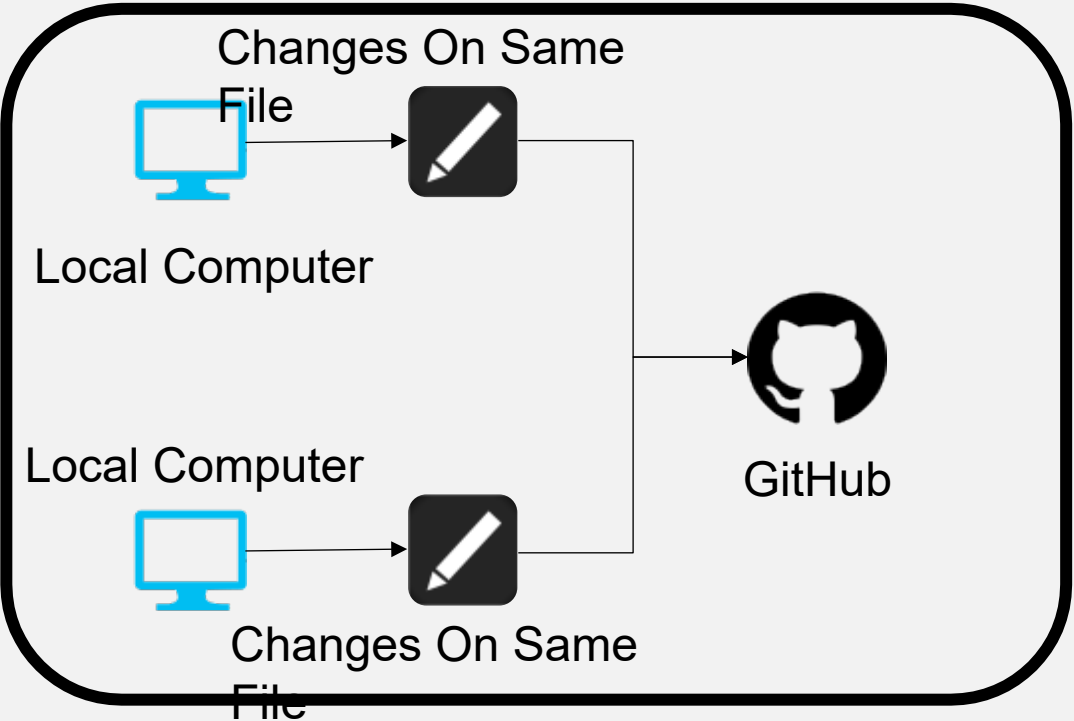
	Dataset	Accuracy Rate [%]
Random Forest	TRAIN	100.00
	DEV	86.33
	EVAL	85.87

# GITHUB AND THE SERVER

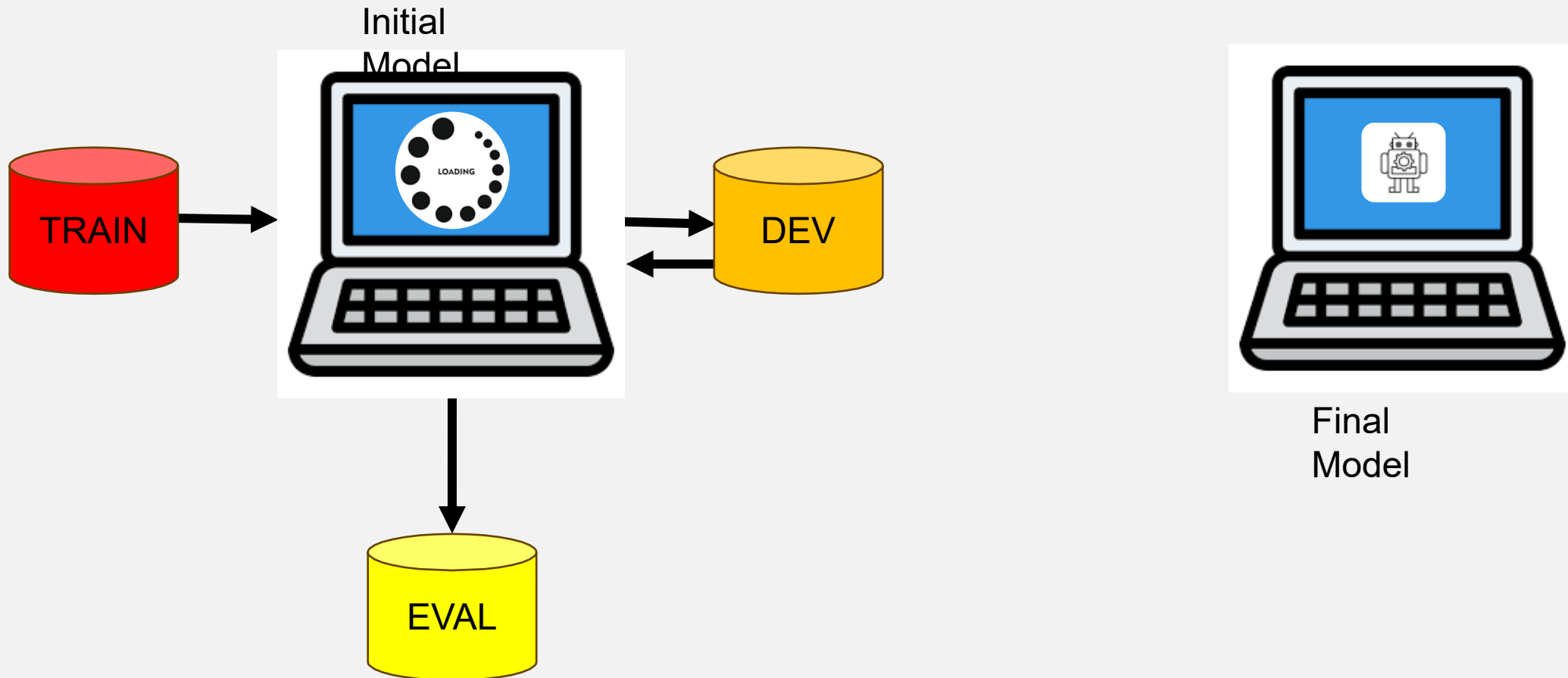
The Server



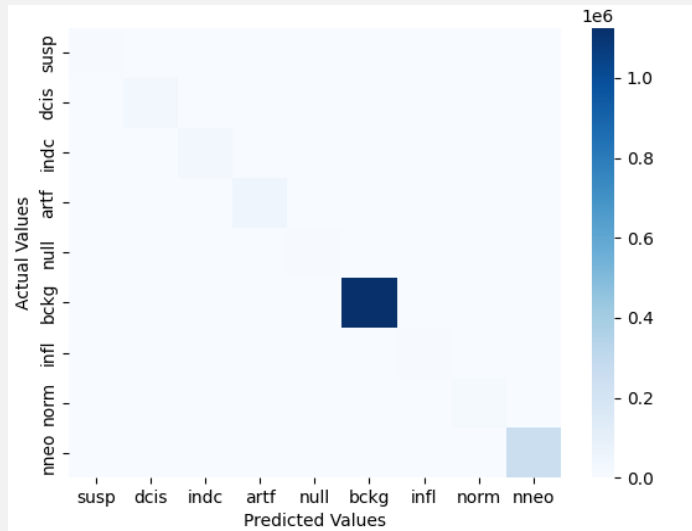
GitHub



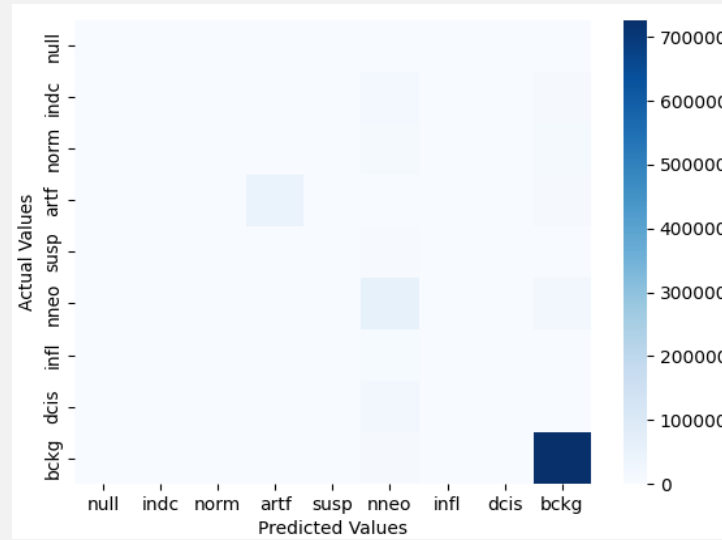
# EXPLAIN TRAIN, DEV, AND EVAL IN THE PROCESS



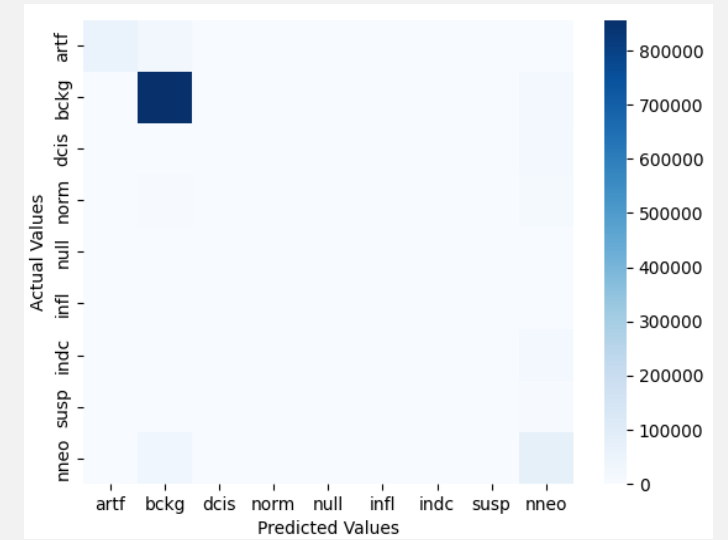
# RNF CONFUSION MATRIX



TRAIN

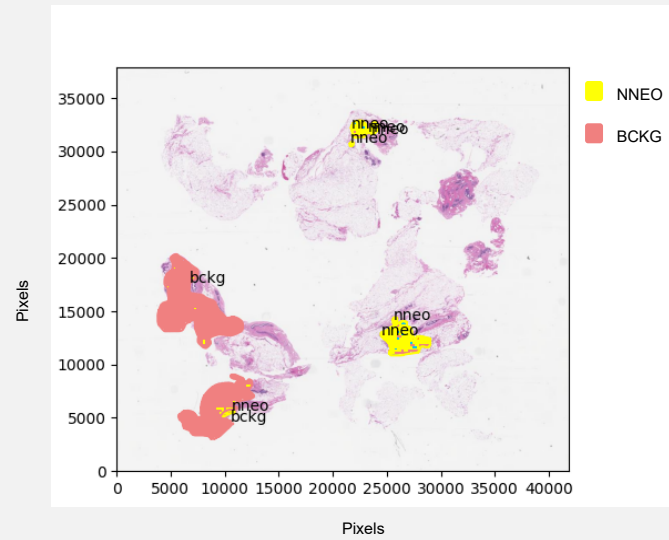


DEV



EVAL

# RNF DECISION SURFACES



# HOW THIS FITS

