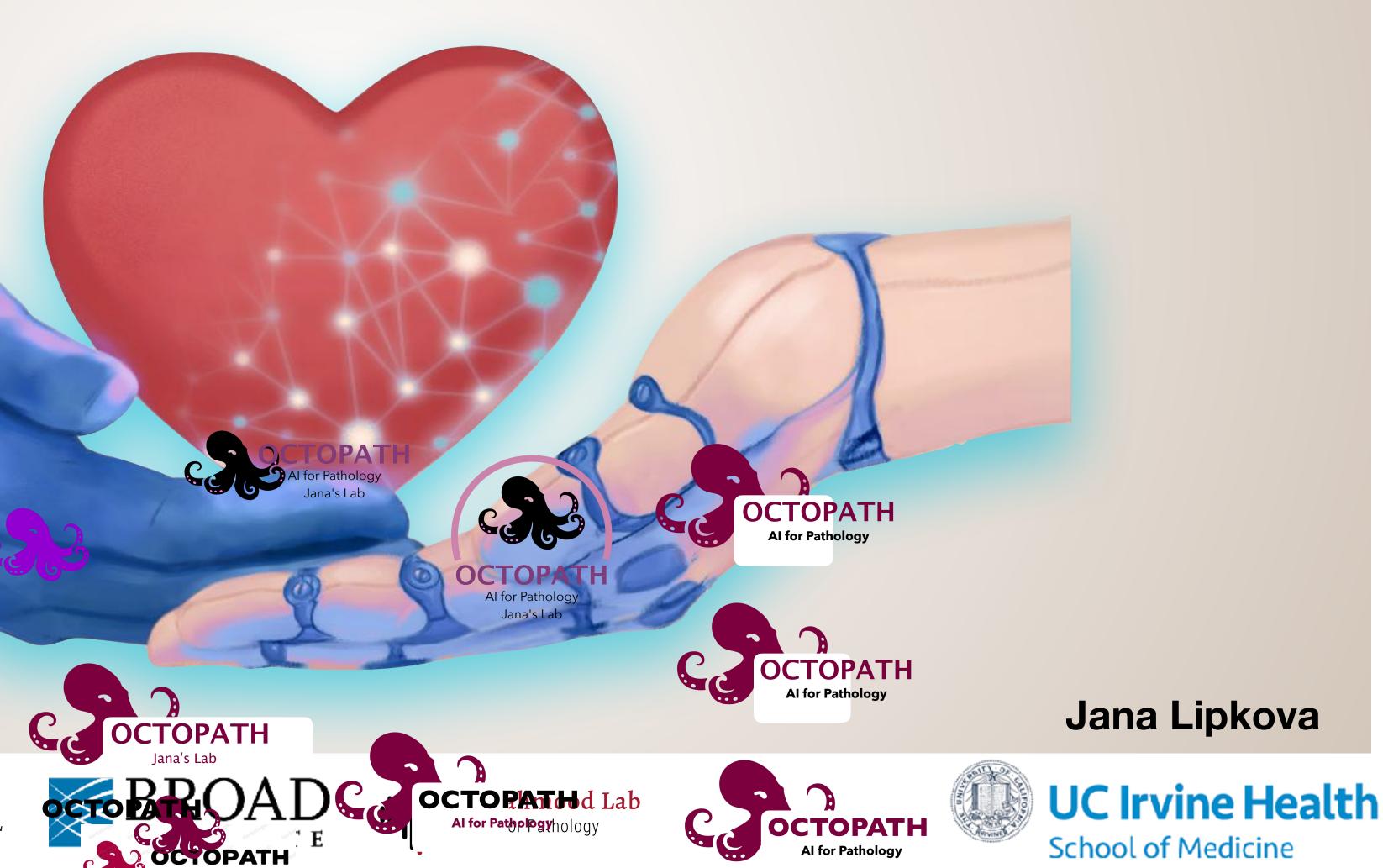
Al-based assessment of cardiac allograft rejections





BRIGHAM AND WOMEN'S HOSPITAL





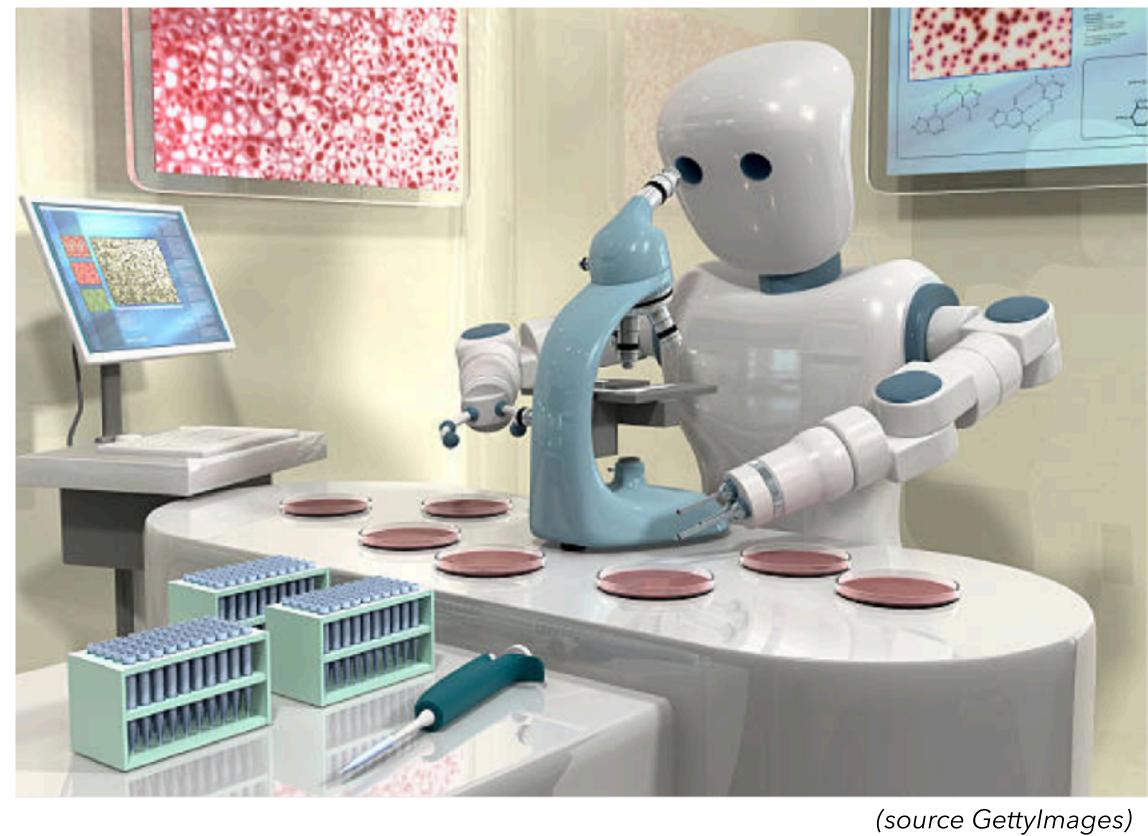
Background:

- Histopathology data _
- Cardiac Allograft Rejections _

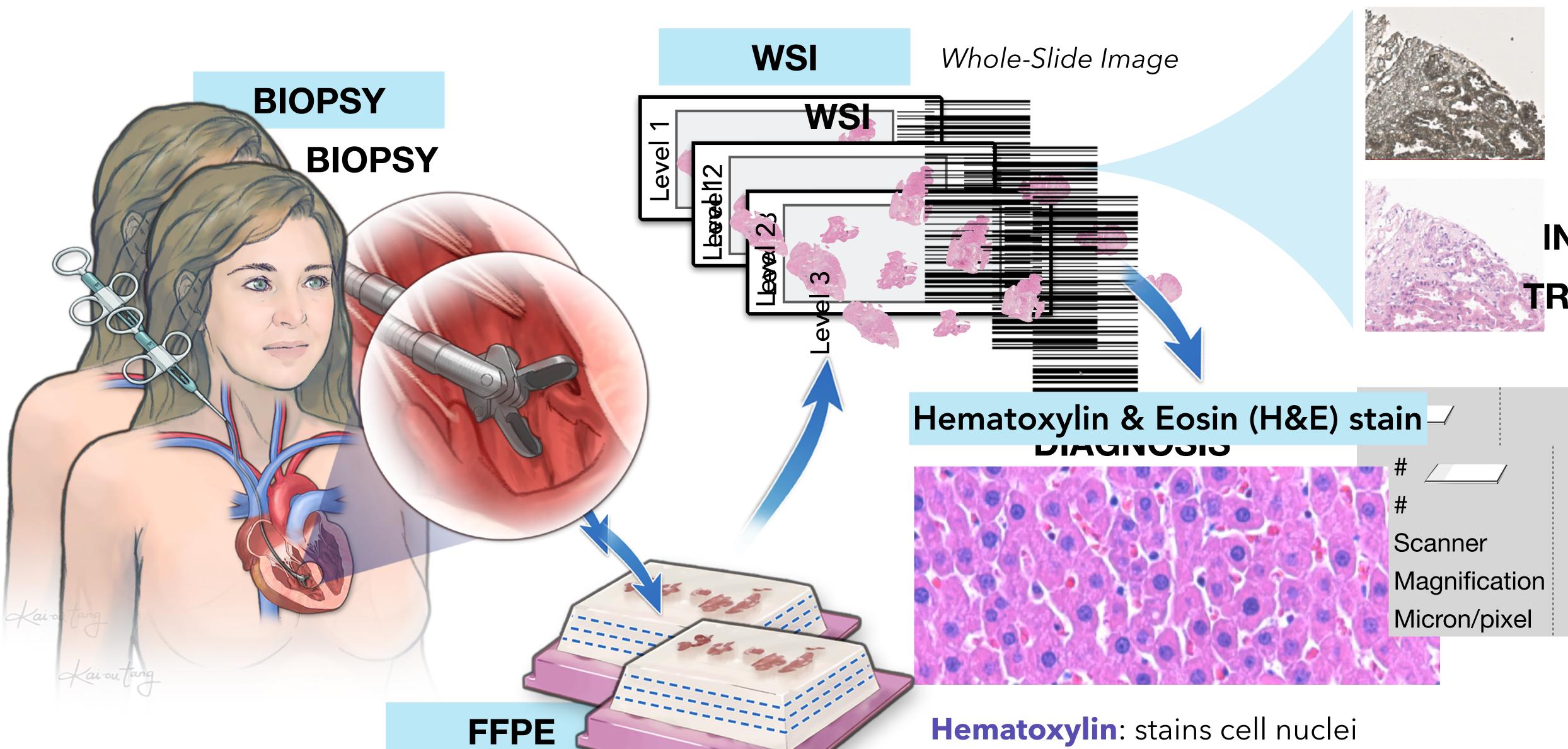
Al-based assessment of allograft rejections







Histology 101

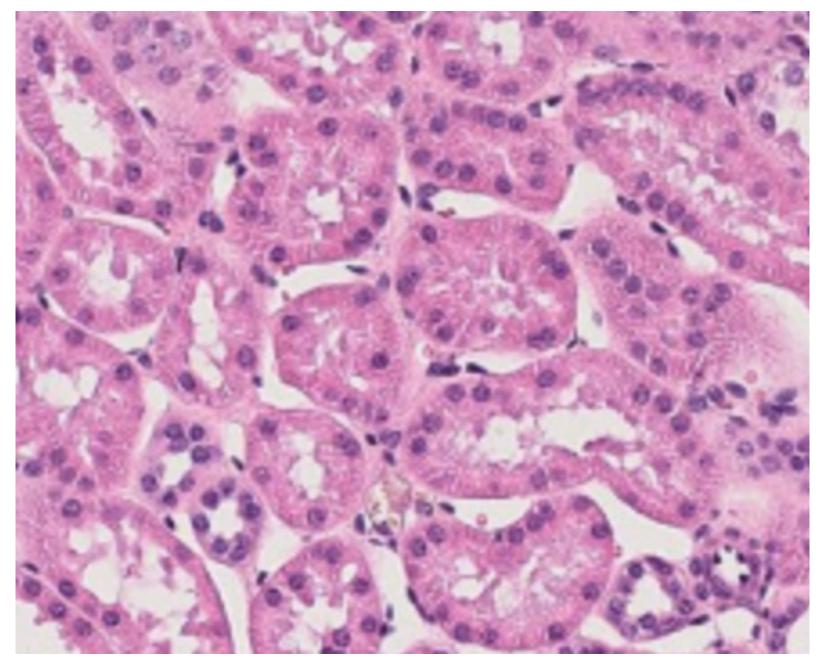


(Formalin-fixated, parafie embedded,

Eosin: the extracellular matrix and cytoplasm

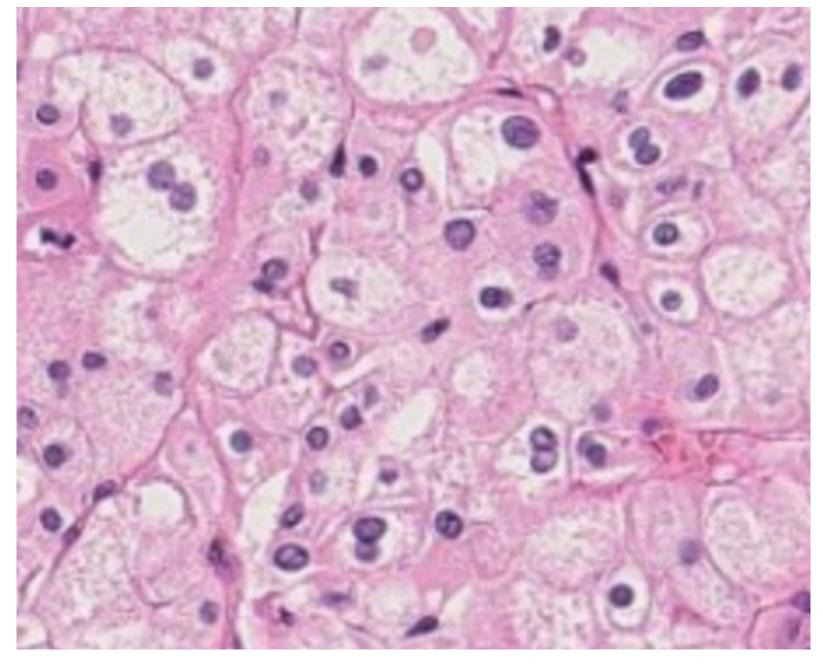
Cancer detection/classification 101

Normal tissue (kidney)



- Symmetric regular structure
- One nuclei per cell
- Cell/nuclei regular shape

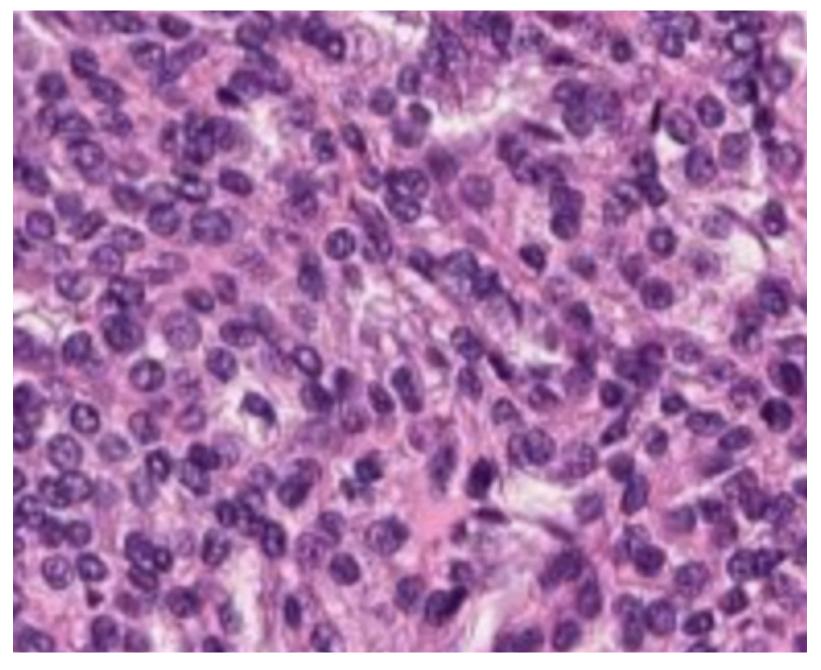
Chromophobe renal carcinoma



- Enlarged nuclei
- Double nuclei per cell
- Irregular shape

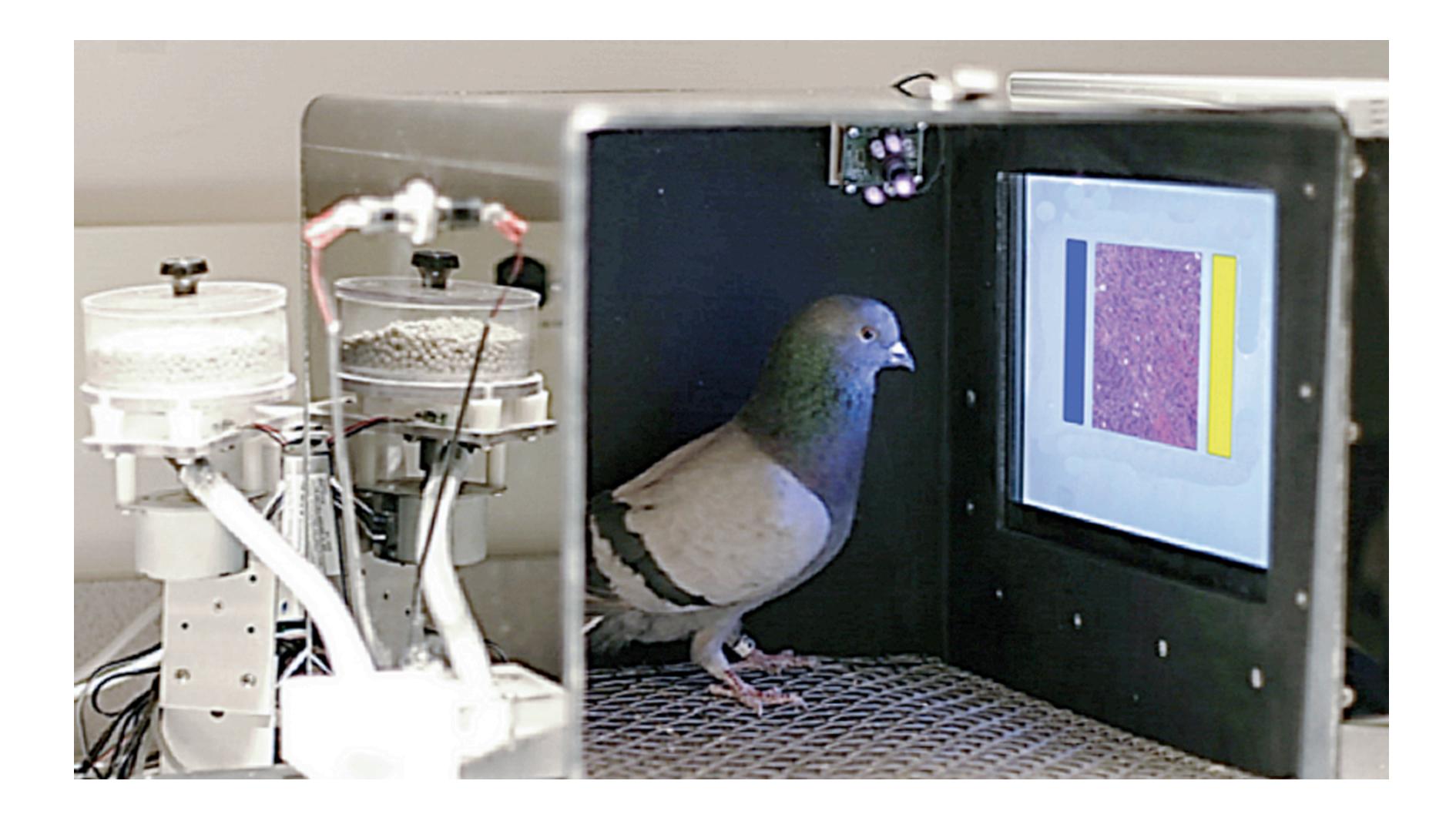
iei ei per cell pe

Papillary renal carcinoma



- Papillary cores lined by neoplastic cells
- Tubulopapillary architecture

Fun fact: Also pigeons can detect cancer

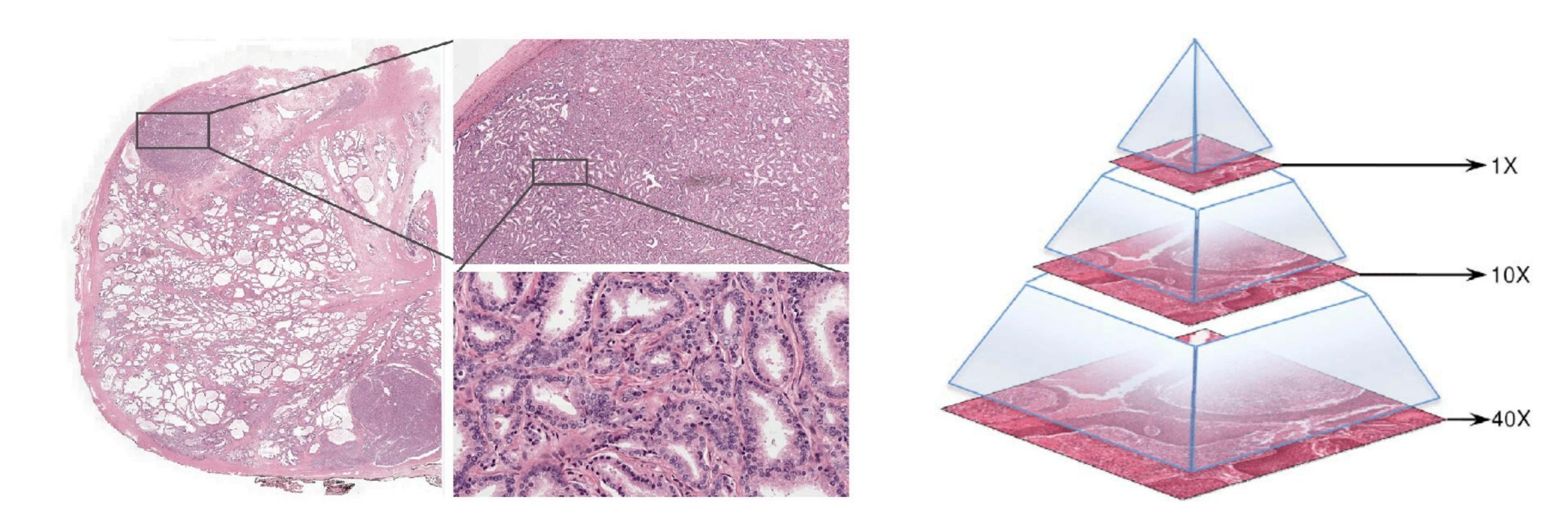


R. Levenson et al: Pigeons (Columba livia) as Trainable Observers of Pathology and Radiology Breast Cancer Images, PloS one, (2015)



Digital Pathology: Whole Slide Images (WSIs)

- High resolution scan of an entire tissue section (0.25 0.5 microns per pixel)
- Gigapixel image: 100,000 x 100,000 pixels
- 100 WSI have cca same amount of pixels as whole ImageNet
- Different Stains: H&E, IHC

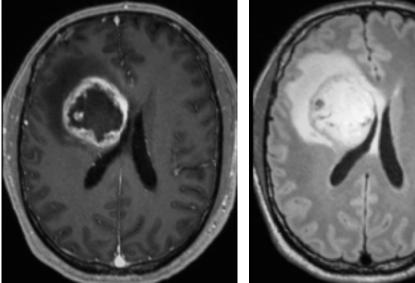




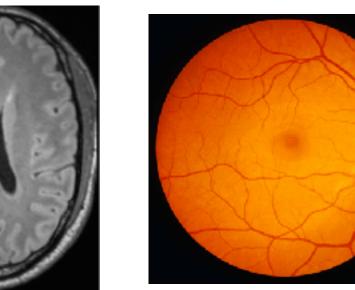
Medical Data

Radiology

Photography



(MRI head scan)





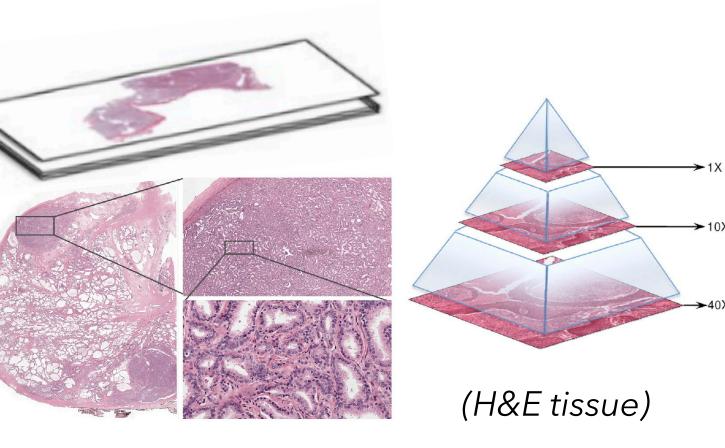
(Fundus / skin photography)

- 3D images
- gray-scale images
- resolution: ~1 mm
- size: 256x256x256 voxels

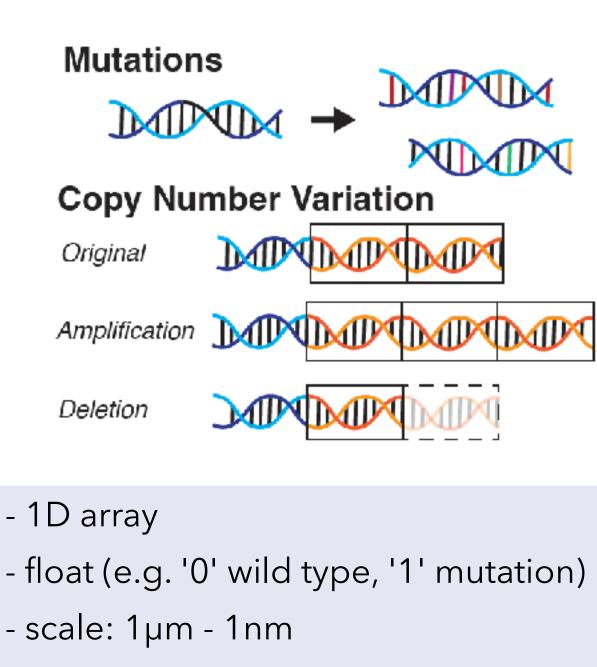
- 2D images
- RGB
- 10 µm 1 mm
- size: ~1,700x1,700 pixels

- 2D images
- RGB
- scale: ~0.1µm
- 100,000x100,000 pixels
- (varies with magnification, tissue size etc)

Histology

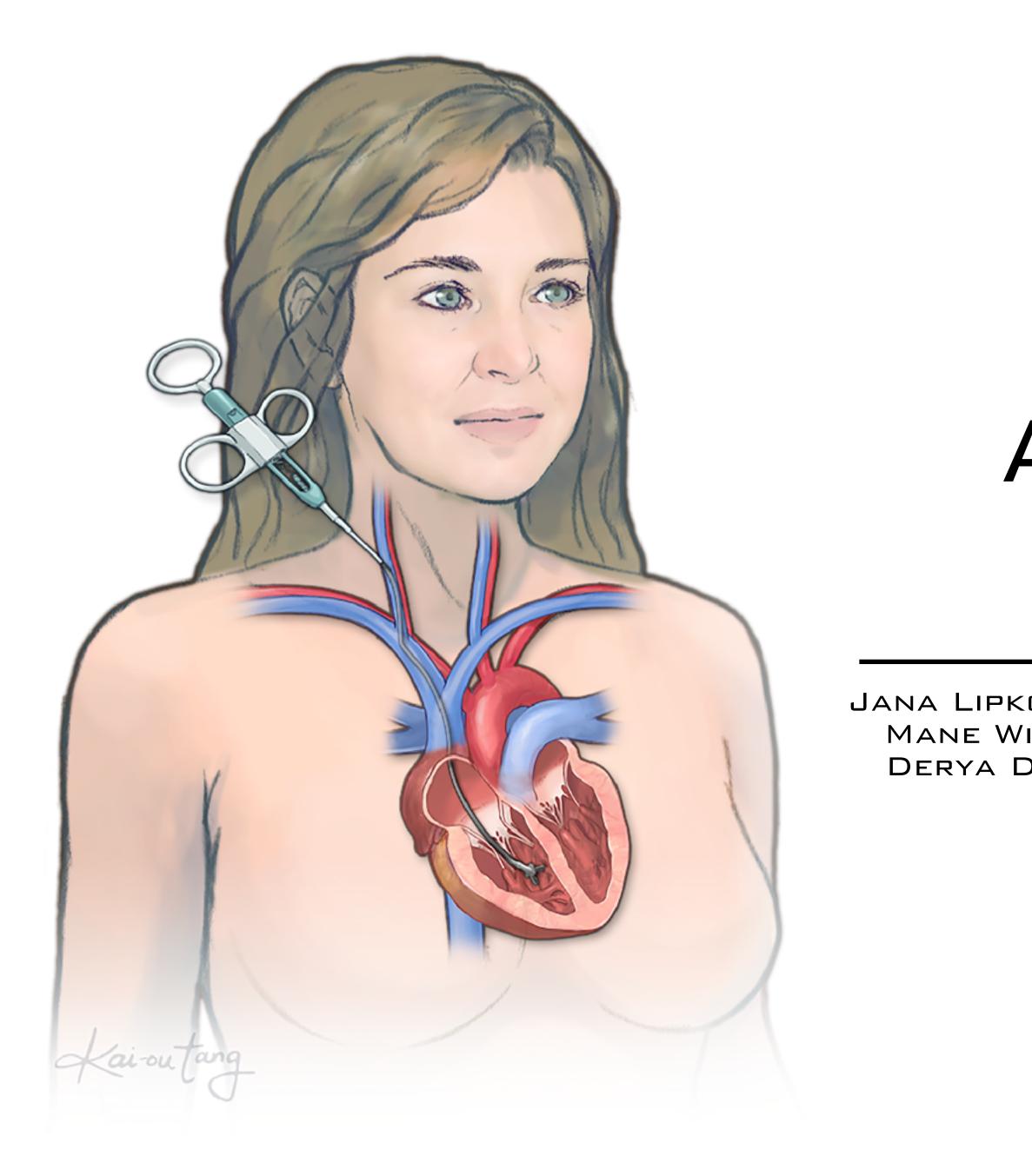


Genomics



- 1D array
- scale: 1µm 1nm
- ~20,000 protein-coding genes





Al-Assessment of Cardiac Allograft Rejections

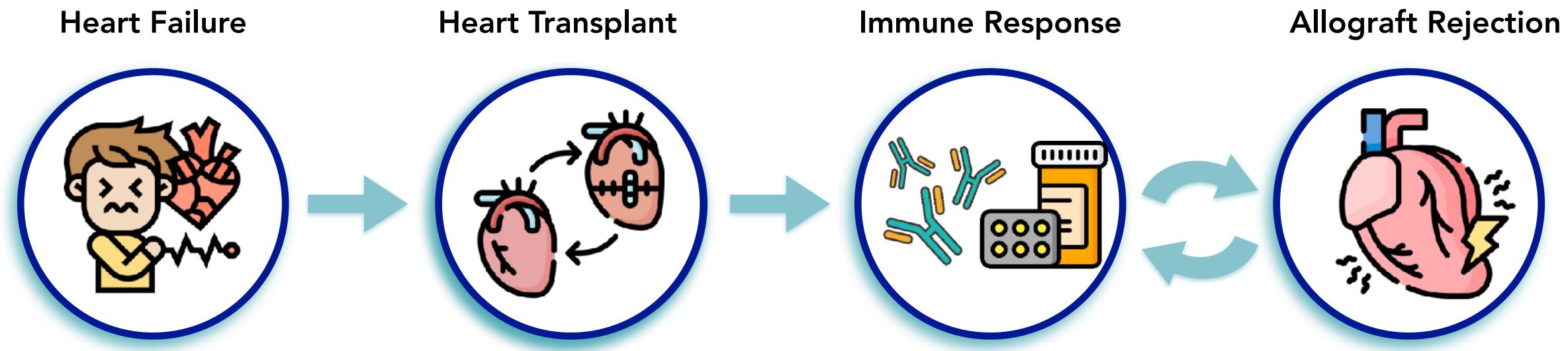
JANA LIPKOVA, TIFFANY Y CHEN, MING Y LU, JINGWEN WANG, MAHA SHADY, MANE WILLIAMS, RICHARD MITCHELL, MEHMET TURAN, GULFIZE COSKUN, Derya Demir, Deniz Nart, Funda Y Barbet, Katja E Odening, Yara BANZ, FAISAL MAHMOOD

Lipkova et al. Nature Medicine (2022)





BACKGROUND



- Leading cause of hospitalization in USA/EU
- 26 million cases / year

- Patients with end-stage failure
- 5000 transplants / year

- Immunosuppressives
- Patient-specific set-up

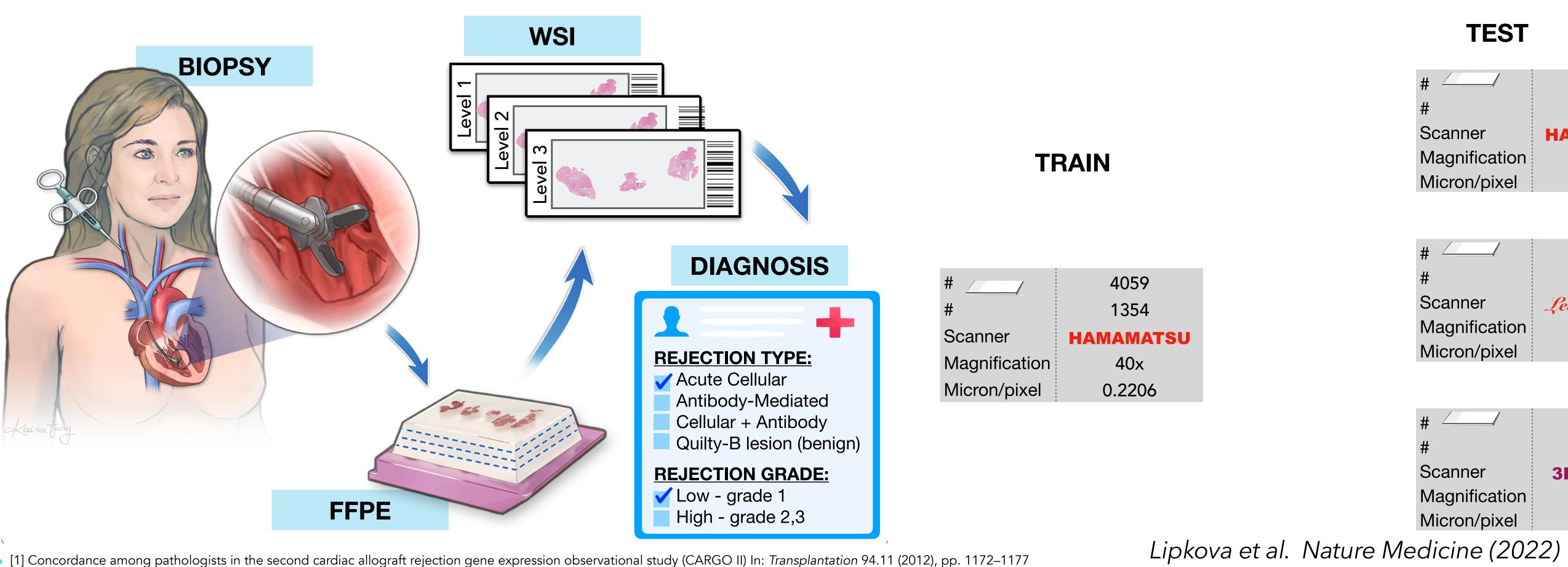
- Main complication & main cause of death
- 40% recipients



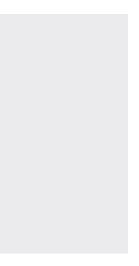
MOTIVATION

APPLICATION:

- Early stages of rejections are **asymptomatic** → surveillance **Endomyocardial biopsy** (EMB)
- Gold-standard: manual assessment H&E-stained biopsies:
- Rejection type & grade determines the immunosuppressive treatment regime

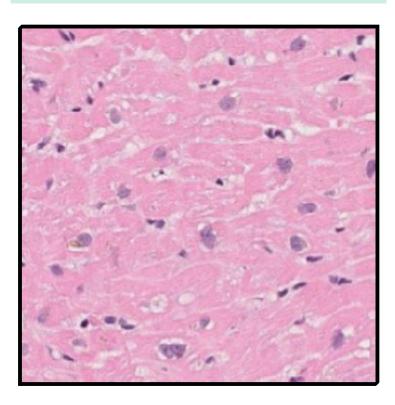


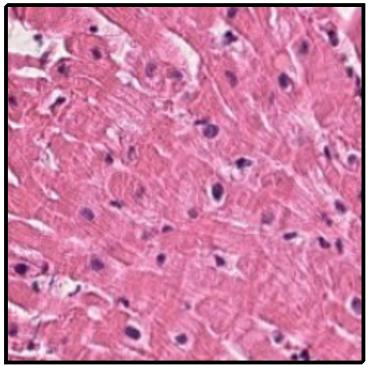
detection and subtyping of rejections (acute cellular, antibody-mediate, benign mimickers) and grading (I-III)

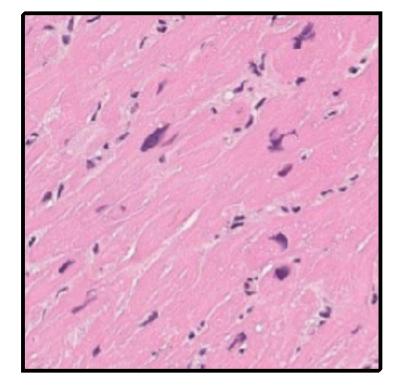


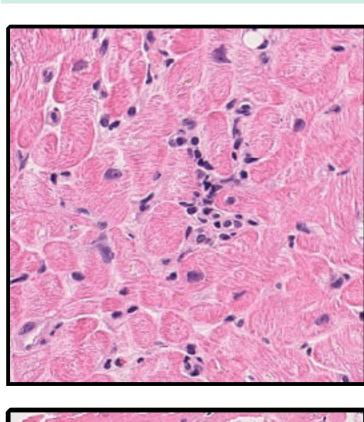
101: Rejection Types

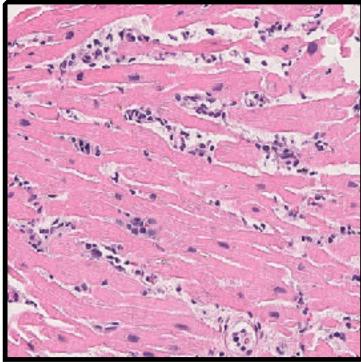
Normal tissue

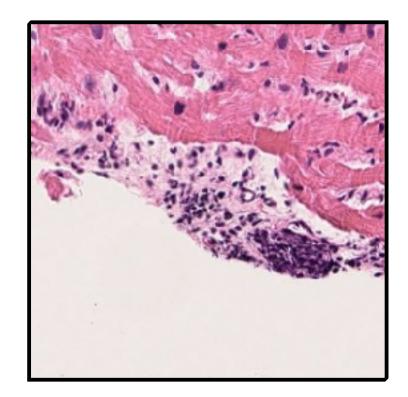




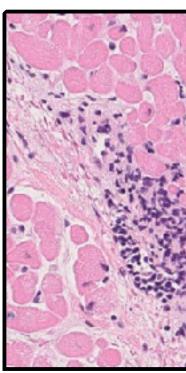


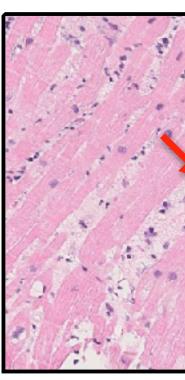


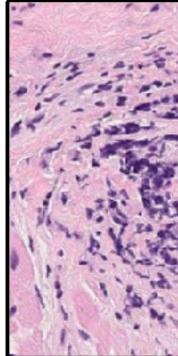


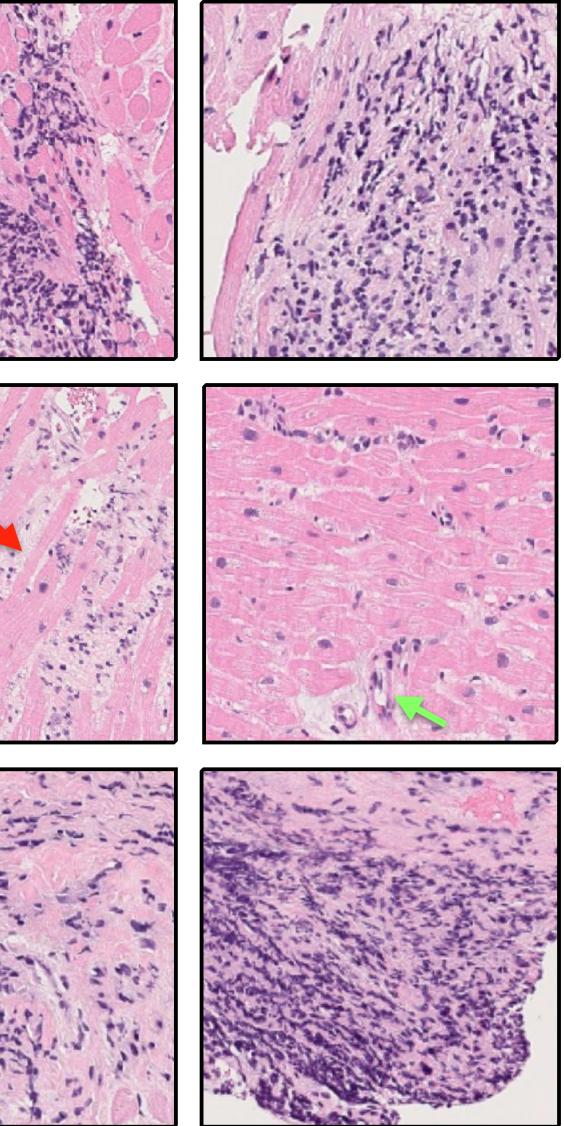


Abnormal tissue









Acute Cellular

- Lymphocyte infiltrates in muscle tissue
- Homogenous structure Comprised of T-cells

Antibody Mediated

- Increased extracellular space + **edema Capillary** endothelial **changes** Increase cell damage More macrophages and necrosis

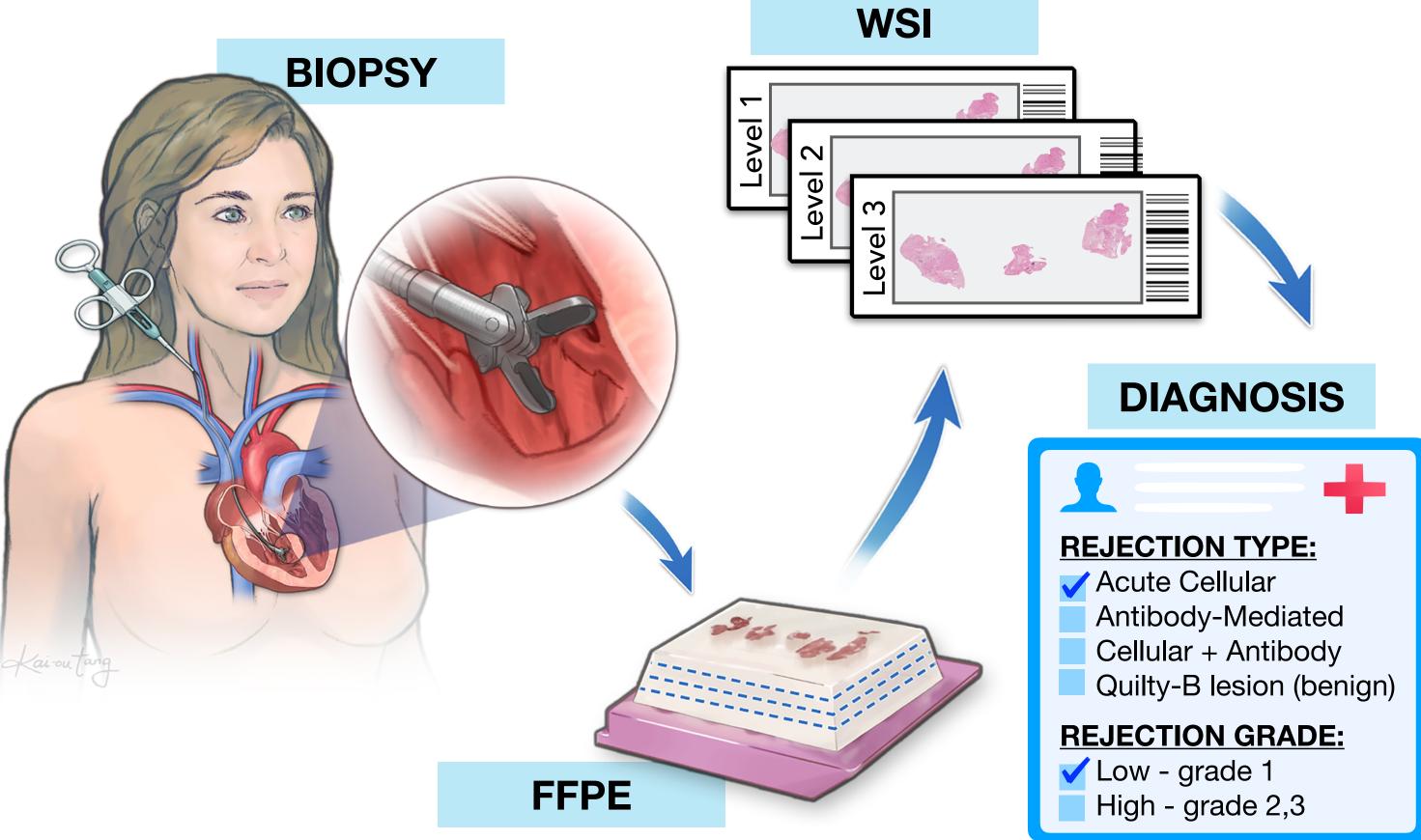
Quilty B Lesions

- Benign lesions
- Mixed B and T-cells, macrophages and plasma cells
- commonly mistaken for cellular rejections



MOTIVATION

- Early stages of rejections are **asymptomatic** → surveillance **Endomyocardial biopsy** (EMB) Gold-standard: manual assessment H&E-stained biopsies:
 - detection and subtyping of rejections (acute cellular, antibody-mediate, benign mimickers) and grading (I-III)
- Rejection type & grade determines the immunosuppressive treatment regime



[1] Concordance among pathologists in the second cardiac allograft rejection gene expression observational study (CARGO II) In: Transplantation 94.11 (2012), pp. 1172–1177

CHALLENGES:

Substantial inter-rater variability [1]:

- <71 % agree if recipient is rejecting the heart
- <28 % agree on the grade of advance rejections
- 19 % unable to reach majority agreement

Misinterpretation:

- under/over treatment with immunosuppressives
- unnecessary follow-up biopsies

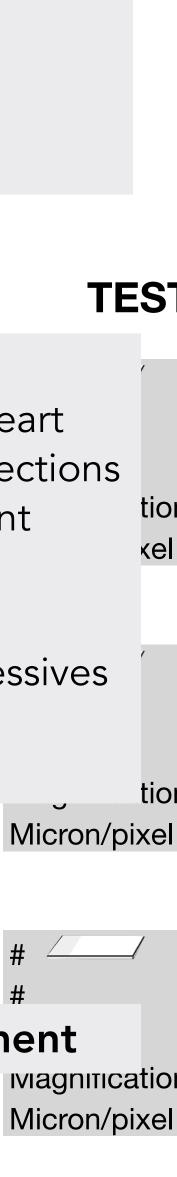
worse outcomes

Scanner	HAMAMATSU	
Magnification	40x	N
Micron/pixel	0.2206	

AIM:

Objective and automated EMBs assessment

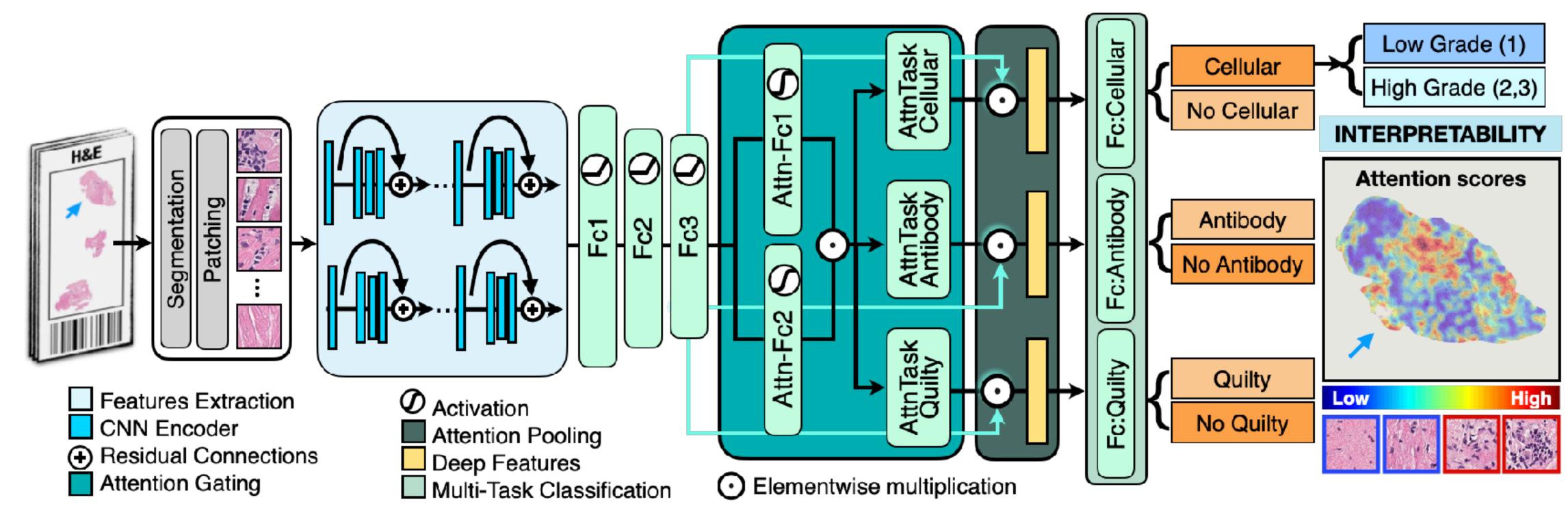
DDEDICTIONIC





Cardiac Rejection Assessment Neural Estimator

- **Input:** H&E-stained EMBs whole-slide-images (WSIs)
- Multi-task, multi-label model: simultaneously identifies presence and type of the rejection (cellular, antibody, and/or quilty lesions). Separate classifier estimate rejection grade
- Multiple-instance learning: use patient diagnosis as only label
- (avoid pixel-level annotations, supports large-scale deployment)
- Attention scores, reflecting relevance of each biopsy region, enable visual interpretation of the model's predictions





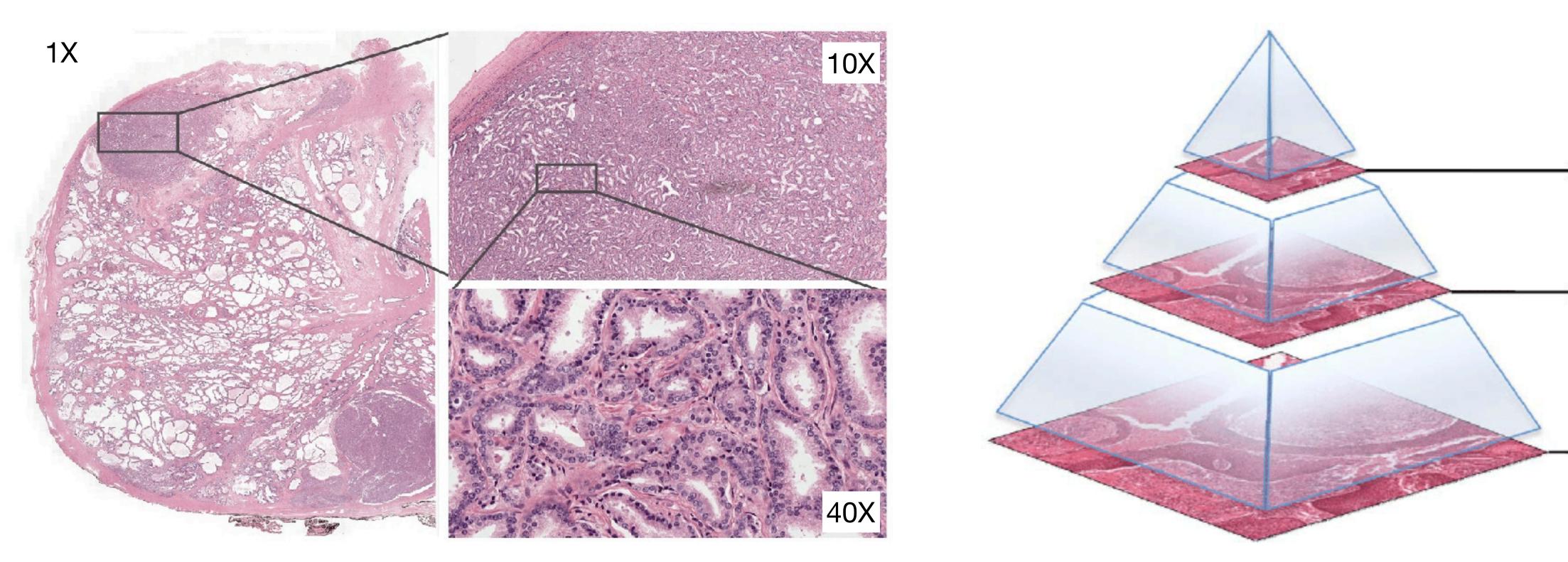
Lipkova et al. Nature Medicine (2022)





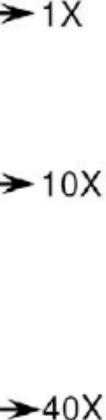
Digital Pathology: Whole Slide Images (WSIs)

- 1 WSI ~ 1 billion pixels !!!
- 100 WSI has more pixels than <u>whole</u> ImageNet
- Difficult to train AI directly on WSI

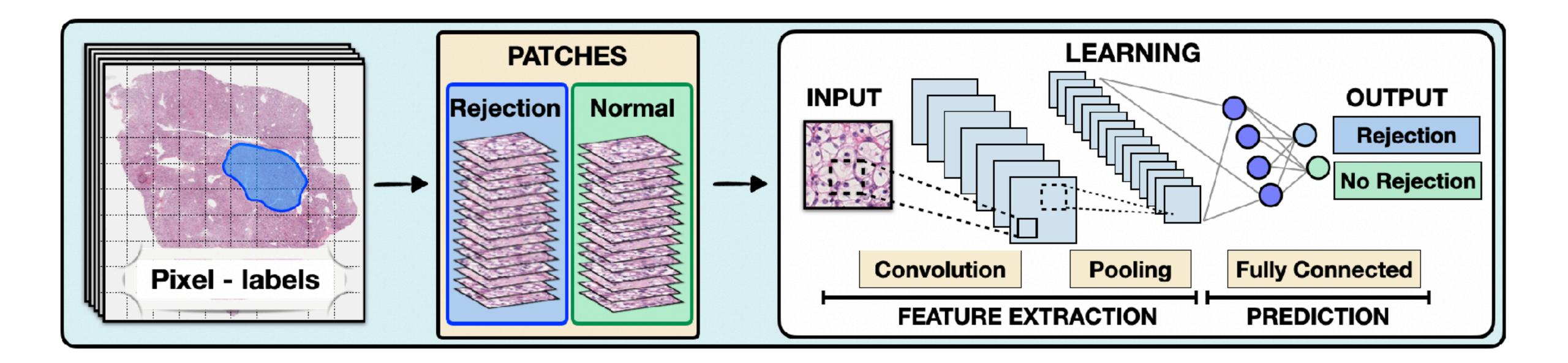


- High resolution scan of an entire tissue section (0.25 - 0.5 microns per pixel)





Typical Deep Learning for Pathology

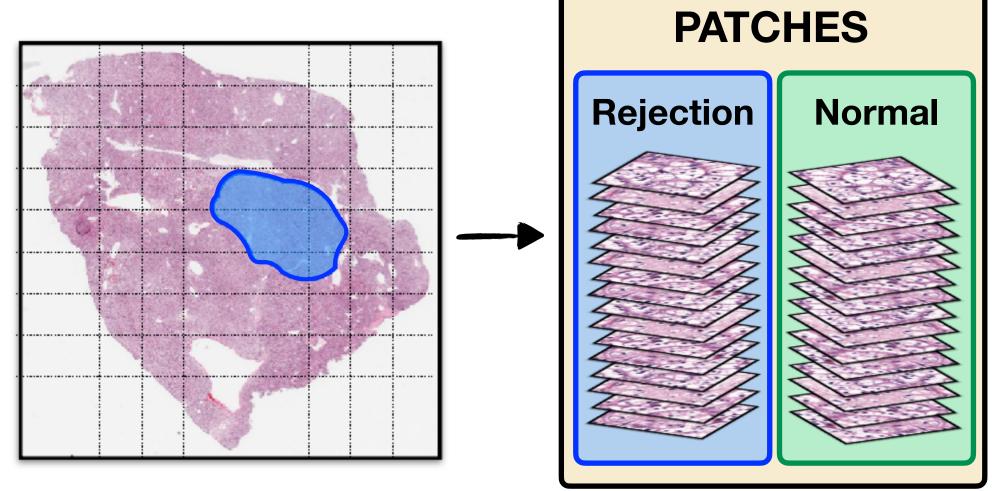


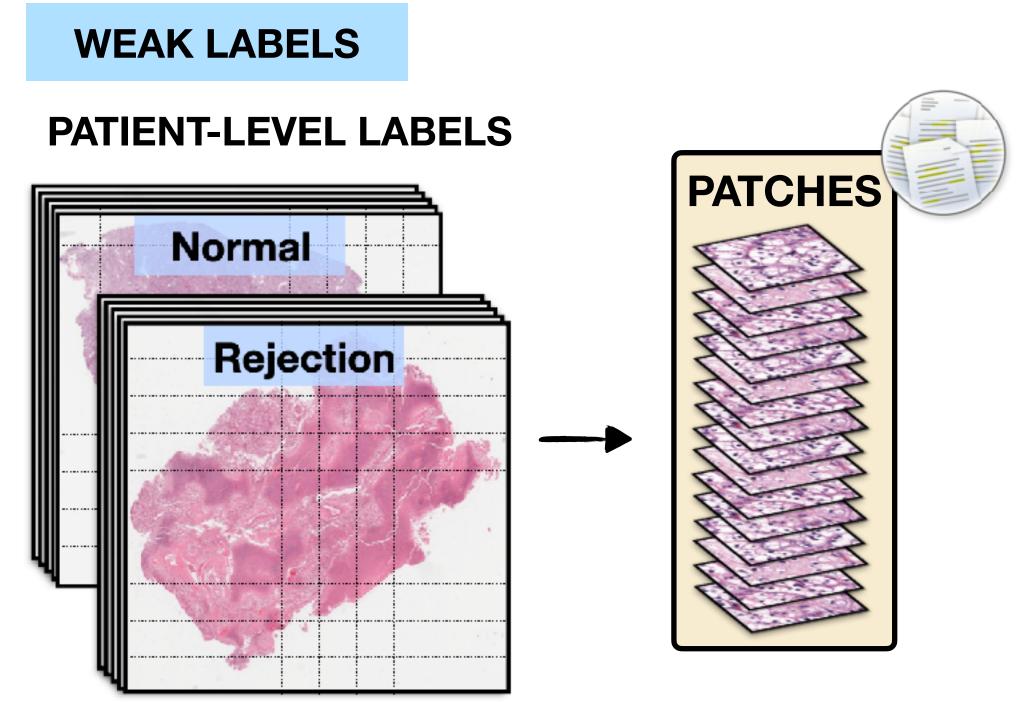
Laborious and time consuming to annotate gigapixels large histology images
Disease borders not always well defined → inter-rater variability → bias
Predictive regions for some tasks (e.g. treatment response) might be unknown
Possible data imbalance: small proportion of image contain the disease (needle-in-haystack problem)
Image annotation is not part of standard clinical practice

Strong vs Weak Supervision

STRONG LABELS

PATCH-LEVEL LABELS





Model alone must discover which tissue regions and which features are predictive for rejections.

Analogy with Natural Images

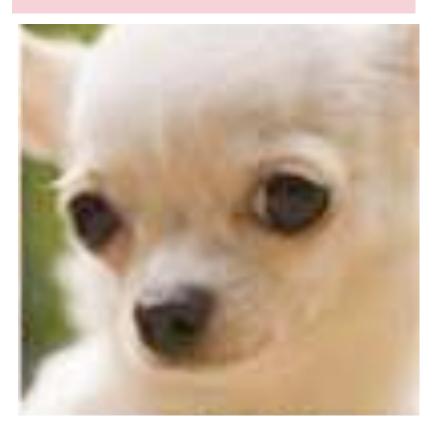
STRONG LABELS

Label for each input

Muffin



Chihuahua



WEAK LABELS

Label for bag of inputs

Contains Chihuahua

No Chihuahua



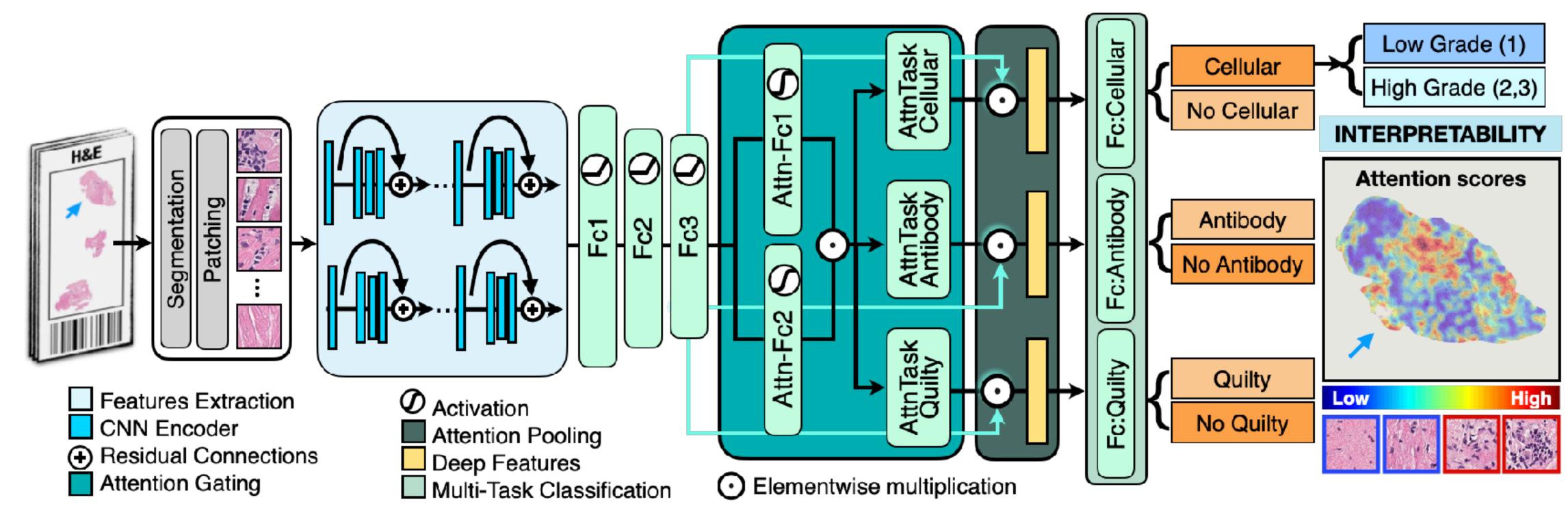
The model alone has to discover which image items and features correspond to chihuahua





Cardiac Rejection Assessment Neural Estimator

- **Input:** H&E-stained EMBs whole-slide-images (WSIs)
- Multi-task, multi-label model: simultaneously identifies presence and type of the rejection (cellular, antibody, and/or quilty lesions). Separate classifier estimate rejection grade
- Multiple-instance learning: use patient diagnosis as only label
- (avoid pixel-level annotations, supports large-scale deployment)
- Attention scores, reflecting relevance of each biopsy region, enable visual interpretation of the model's predictions



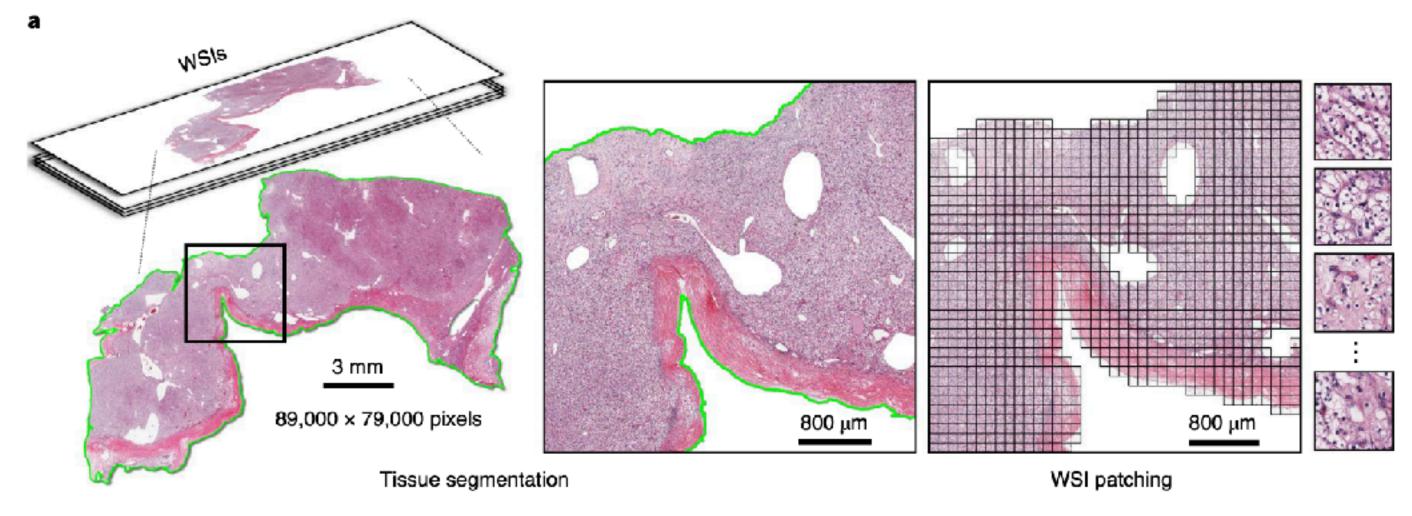


Lipkova et al. Nature Medicine (2022)

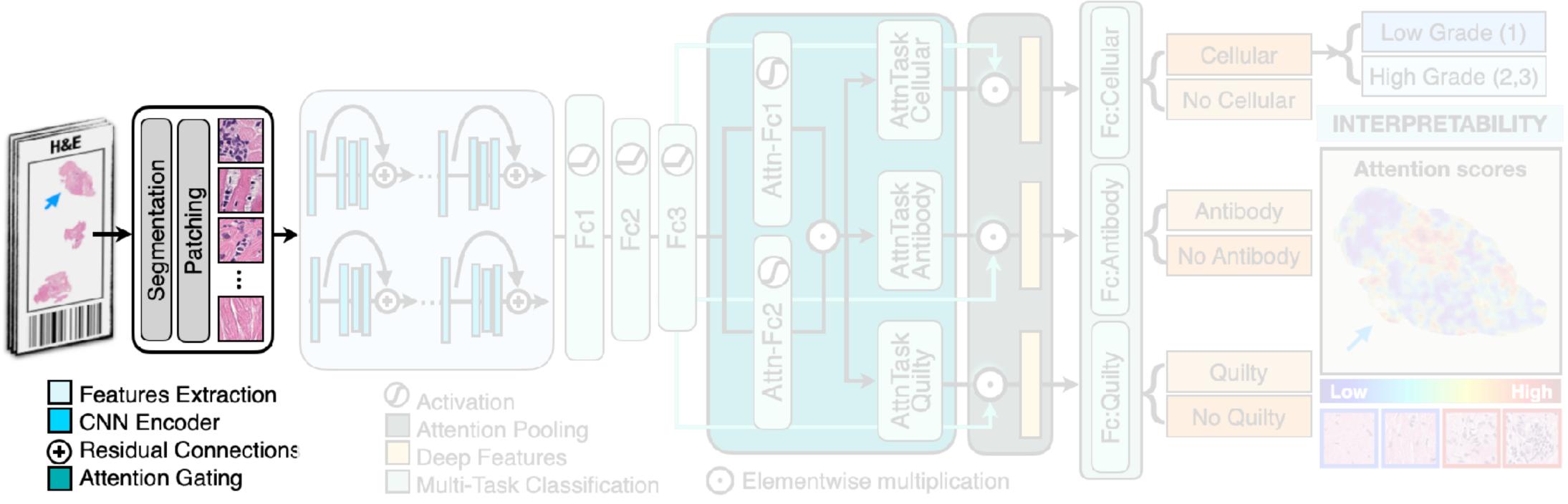




PREPROCESSING

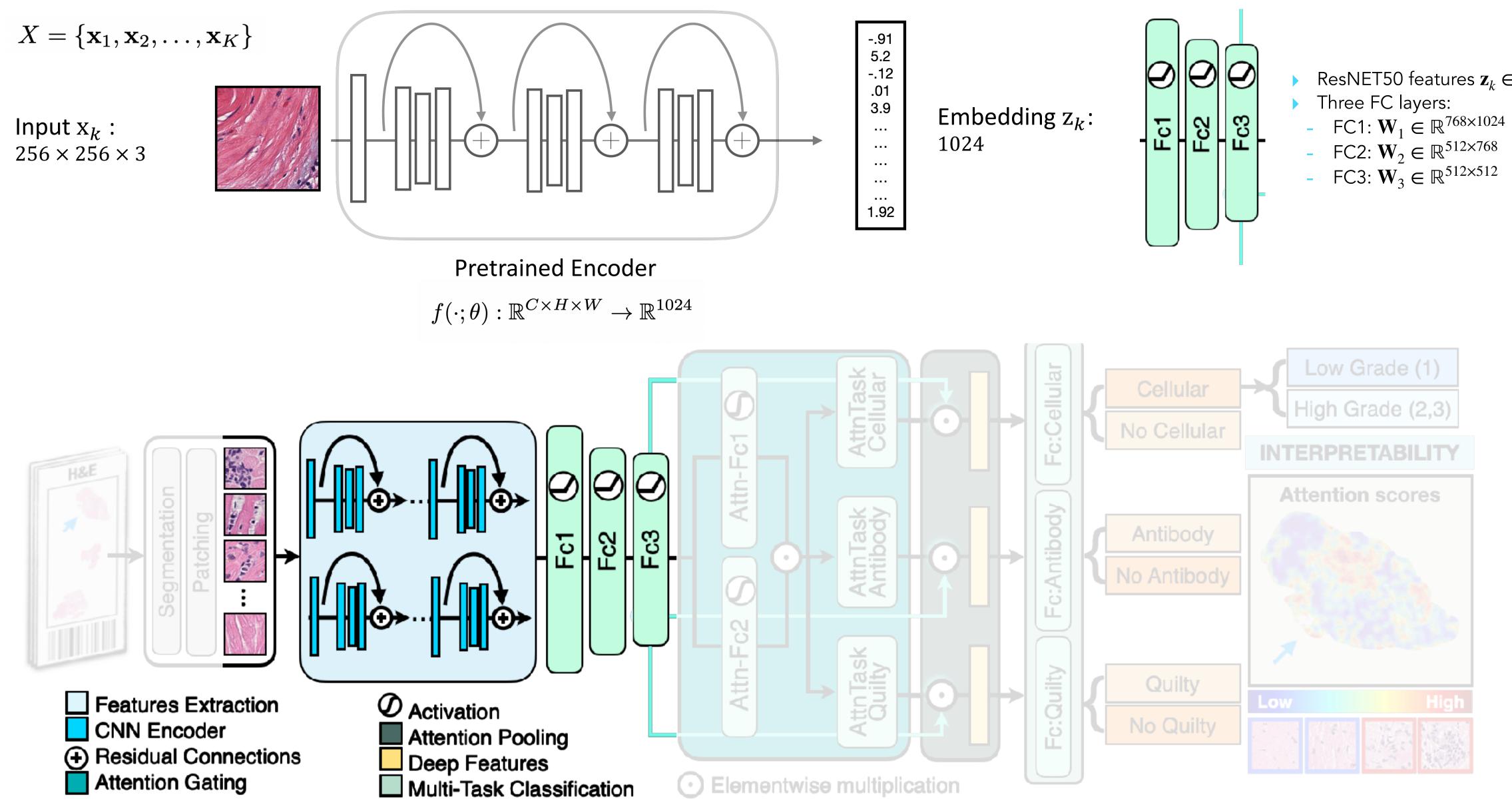


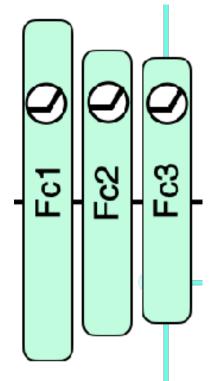
\approx 1 Billion Pixels!



EMBEDDINGS

Patch-level representation of patch k from {1,...,K}

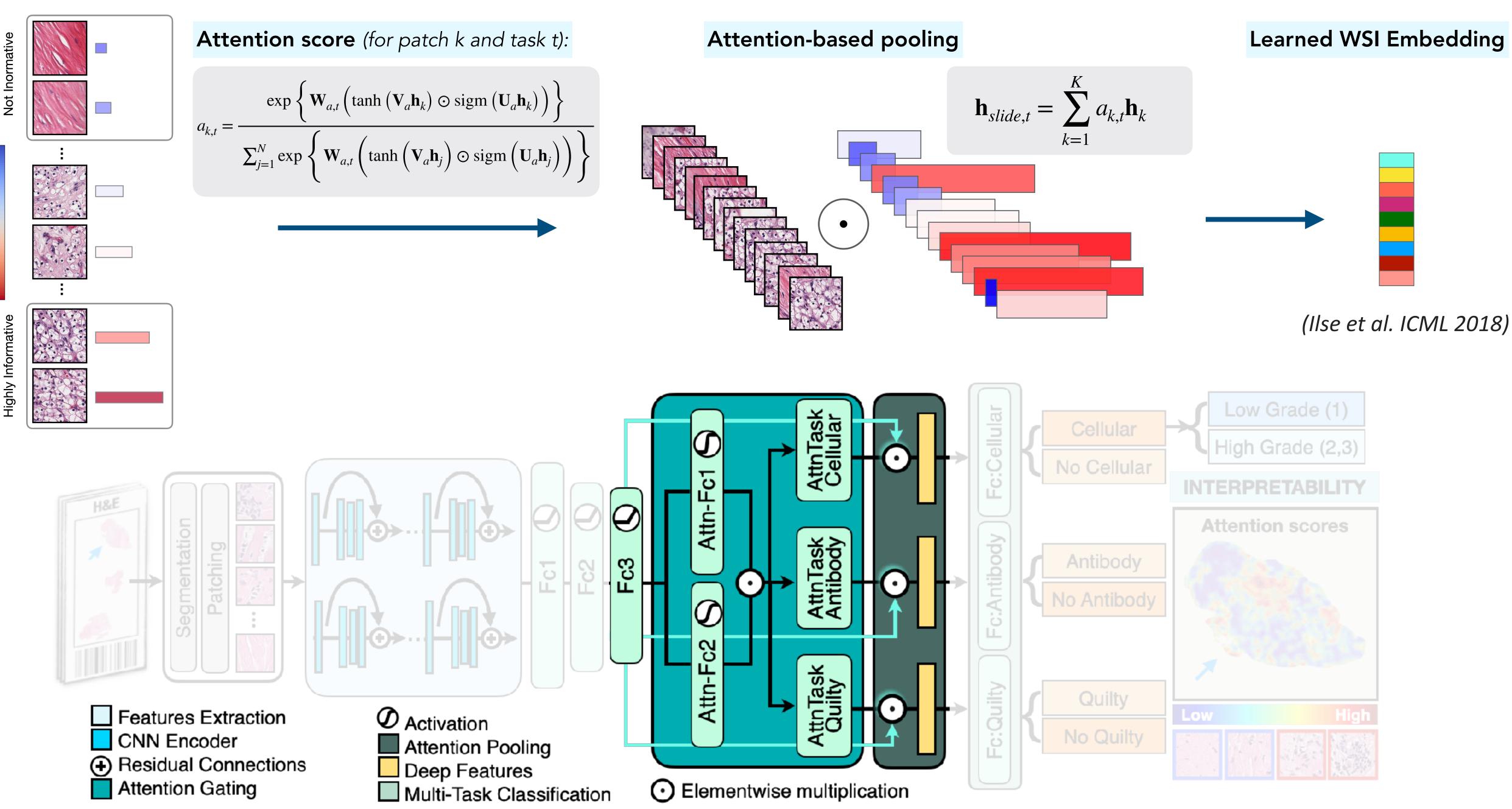




- ResNET50 features $\mathbf{z}_k \in \mathbb{R}^{1024}$

- FC2: $\mathbf{W}_2 \in \mathbb{R}^{512 \times 768}$

ATTENTION LEARNING



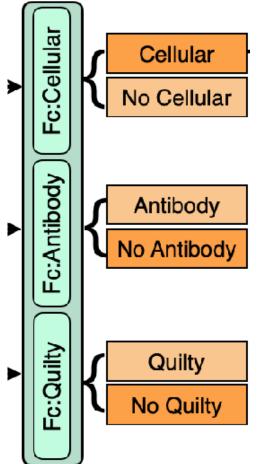
Learned WSI Embedding





MULTI-TASK CLASSIFIER

Learned WSI Embedding



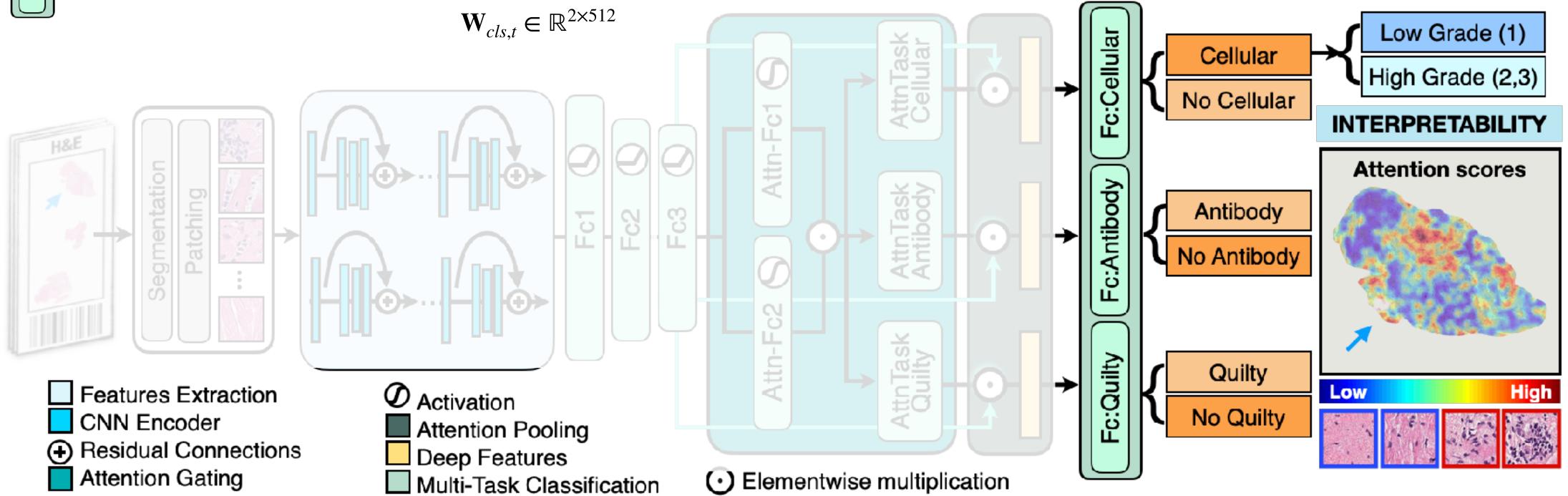
Slide-level representations for task *t*:

$$\mathbf{h}_{slide,t} = \sum_{k=1}^{K} a_{k,t} \mathbf{h}_{k}$$

Slide-level predictions for task t:

 $\mathbf{p}_{t} = \text{Softmax}(\mathbf{W}_{cls,t} \,\mathbf{h}_{slide,t} + \mathbf{b}_{cls,t})$

$$\mathbf{W}_{cls,t} \in \mathbb{R}^{2 \times 512}$$



REJECTION GRADE

Same MIL model, just single-task

INTERPRETABILITY

WSI attention heatmaps

• Elementwise multiplication

Study Design



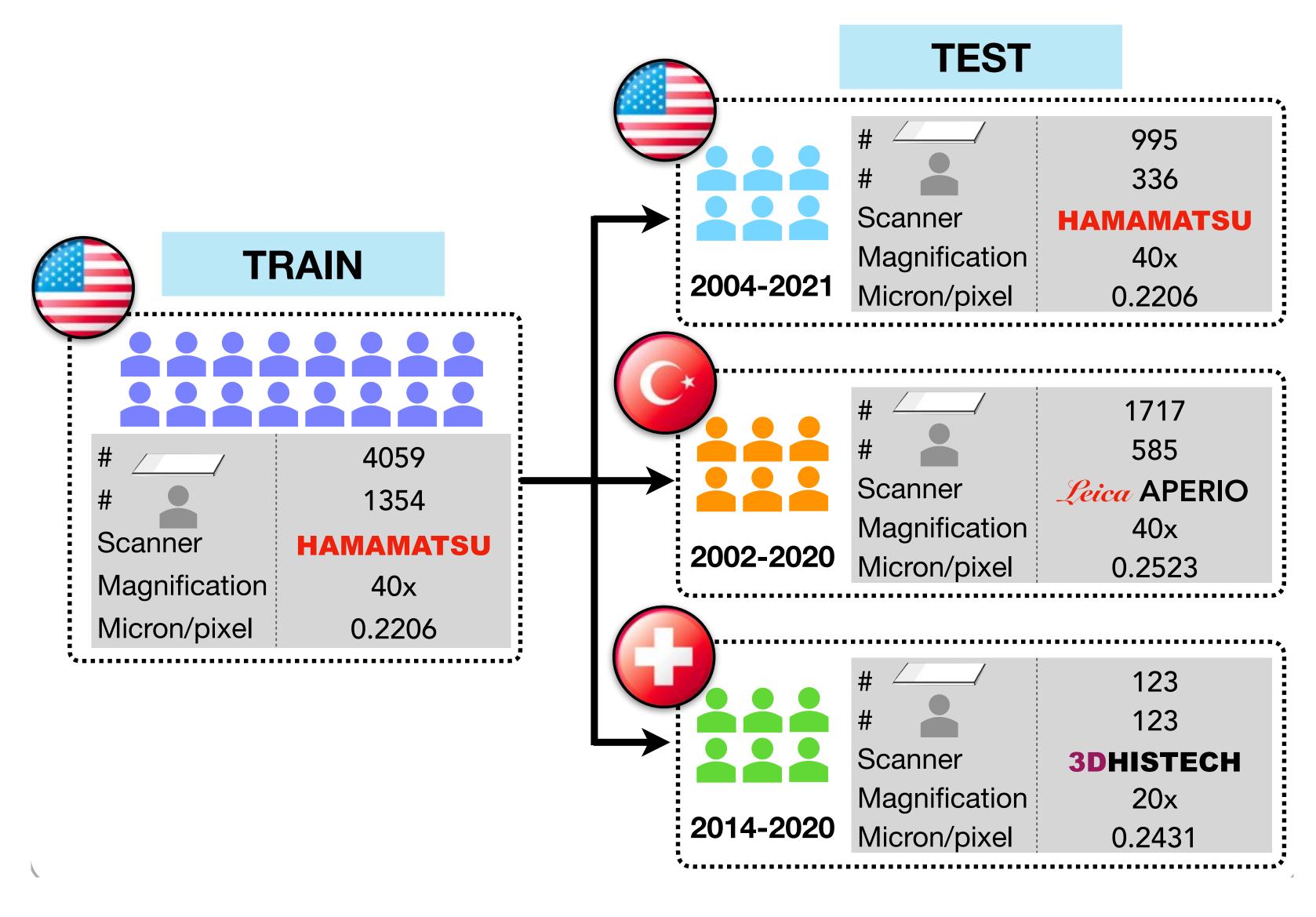
- micron/pixel,
- etc

The model is trained on subset of data collected in USA

70/10/20% split (balance diagnosis)

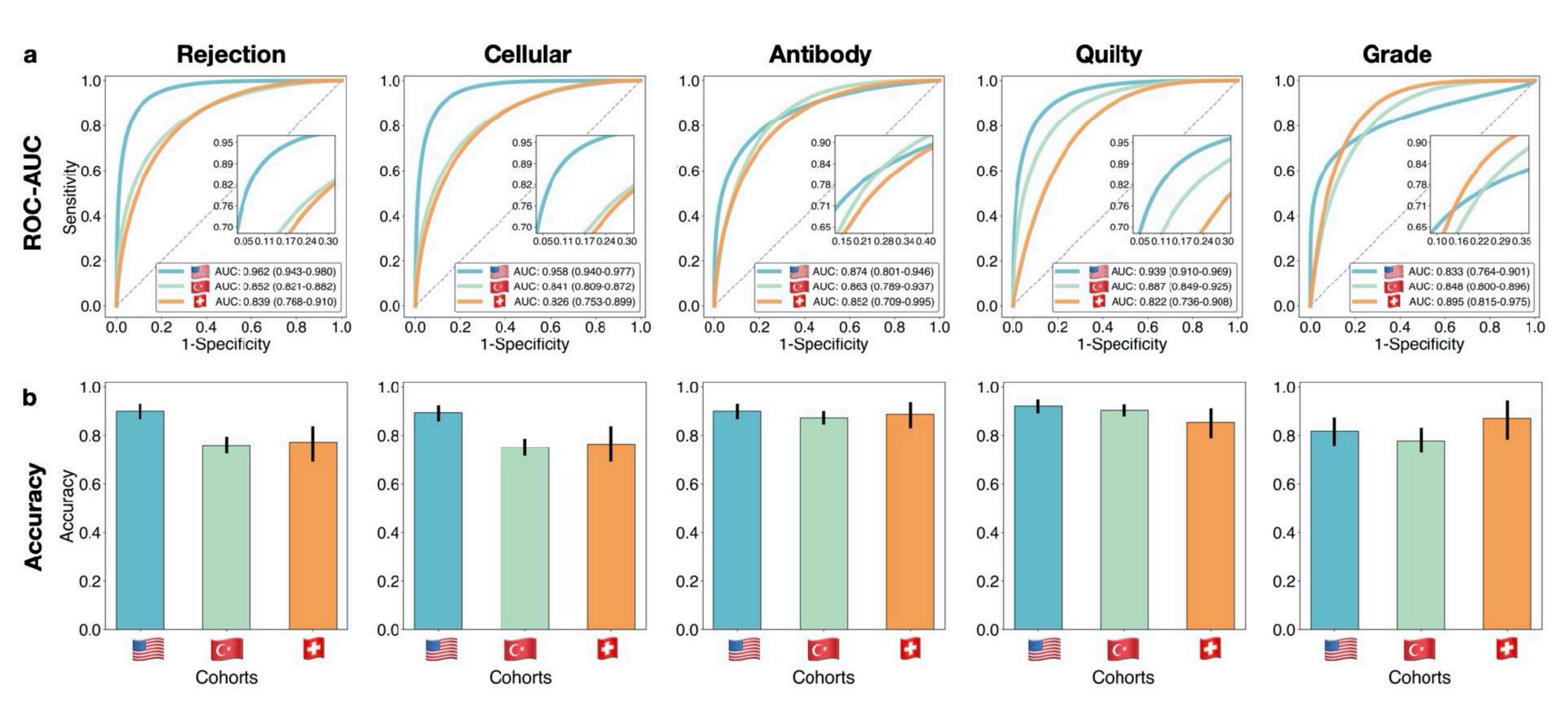
to external cohorts

domain-specific adaptations



PREDICTIONS Lipkova et al. Nature Medicine (2022)

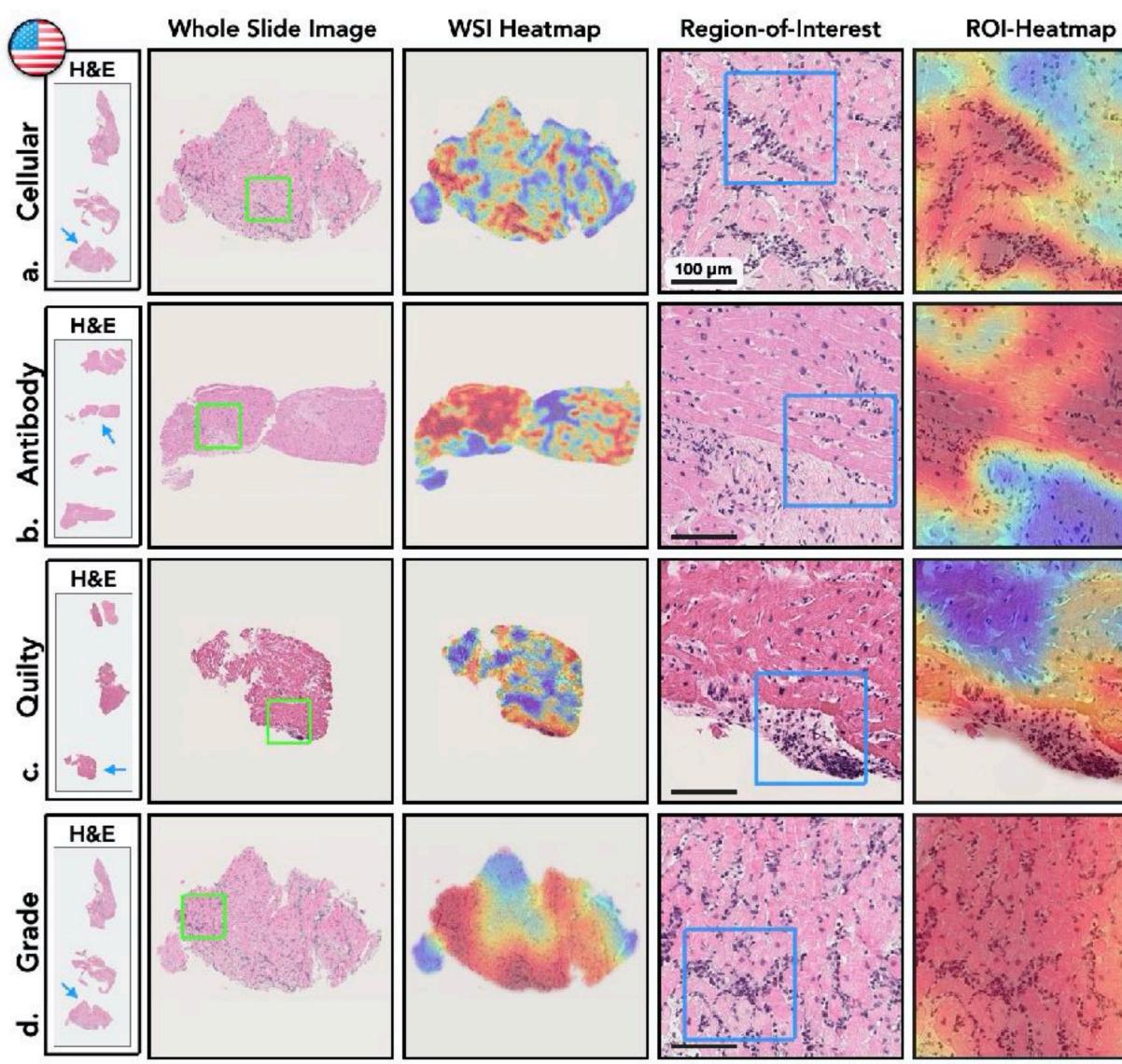
Evaluation & Results



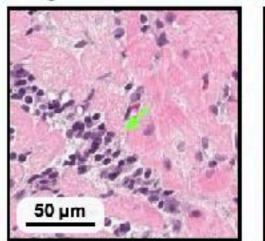
Lipkova et al. Nature Medicine (2022)

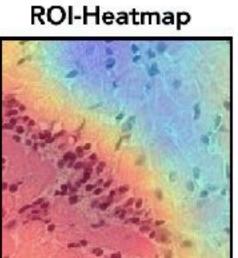


Interpretability

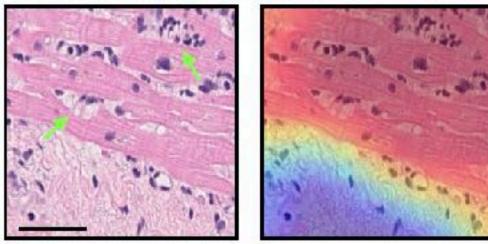


Region-of-Interest

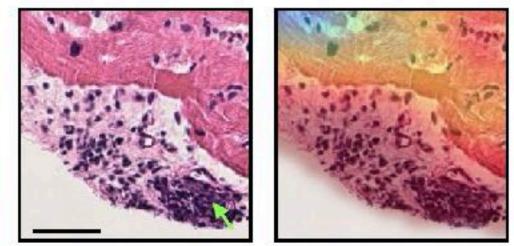




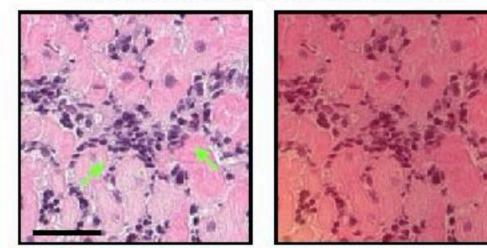
Interstitial lymphocytic infiltrate with myocyte damage



ial edema with mixed inflammatory infiltrate



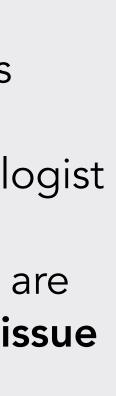
Dense, benign, endocardial lymphocytic infiltrate



Multifocal myocyte injury with diffuse mixed inflammatory

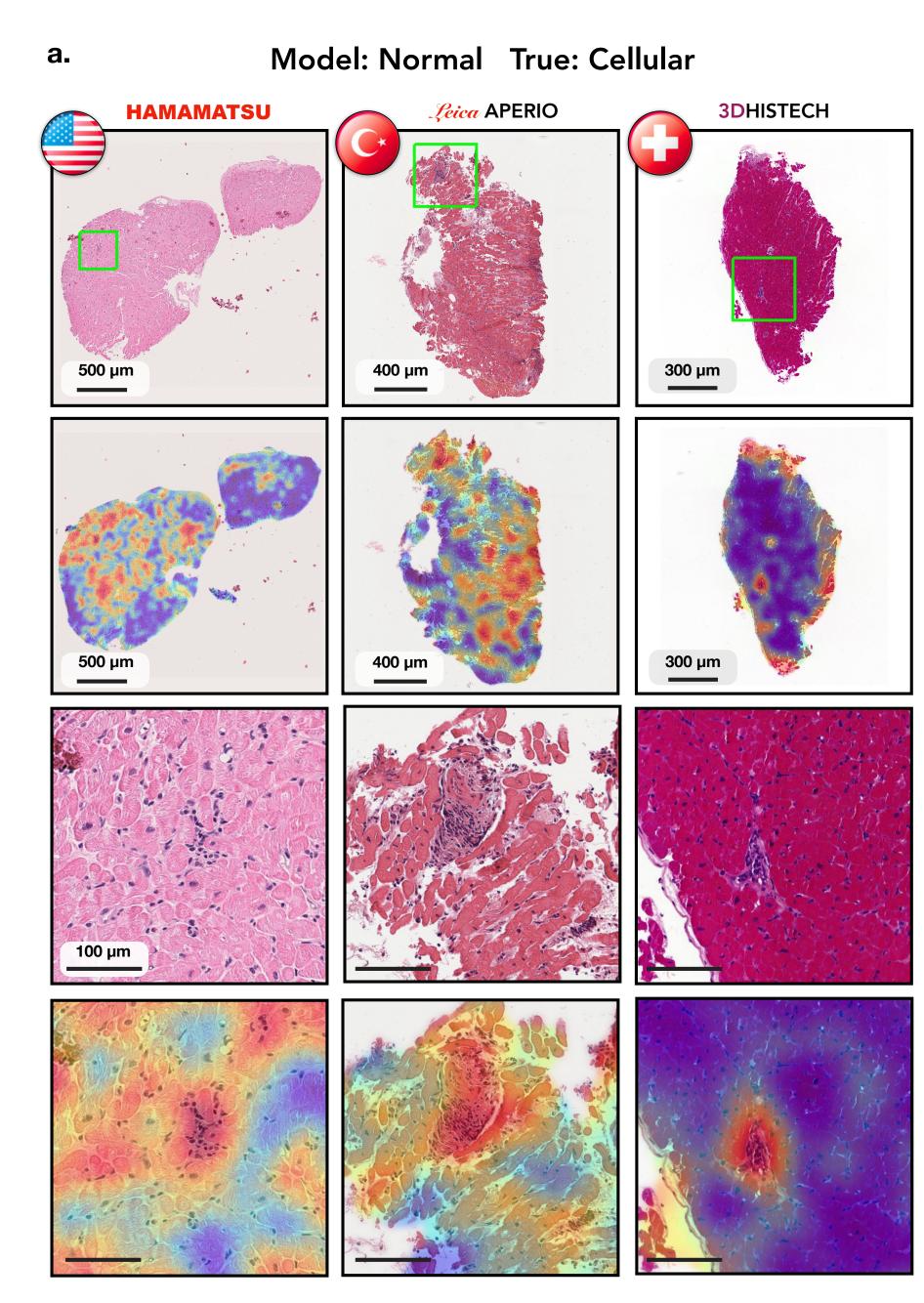
- High-attention (red) regions correspond to rejection morphology used by pathologist for diagnosis
- Low-attention (blue) scores are assigned mostly to **benign tissue**

Lipkova et al. Nature Medicine (2022)

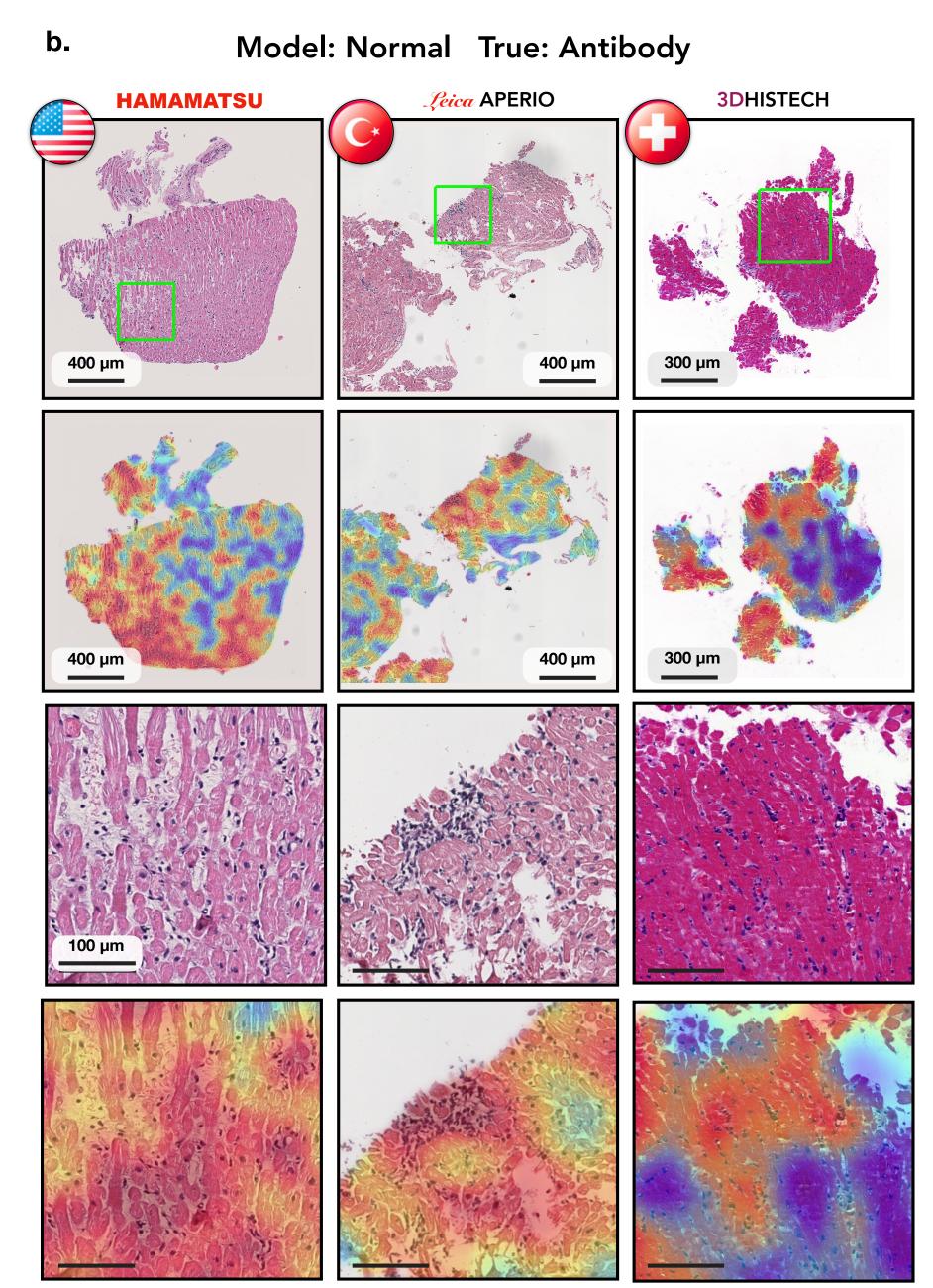




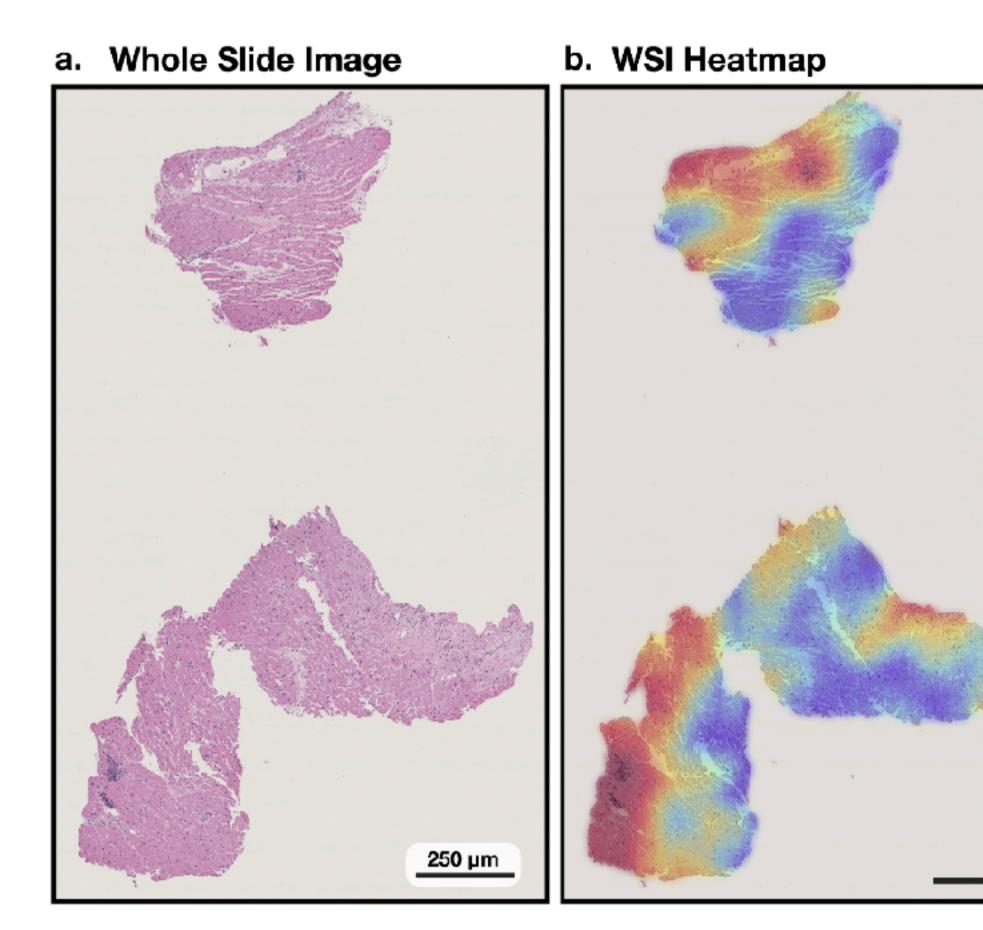




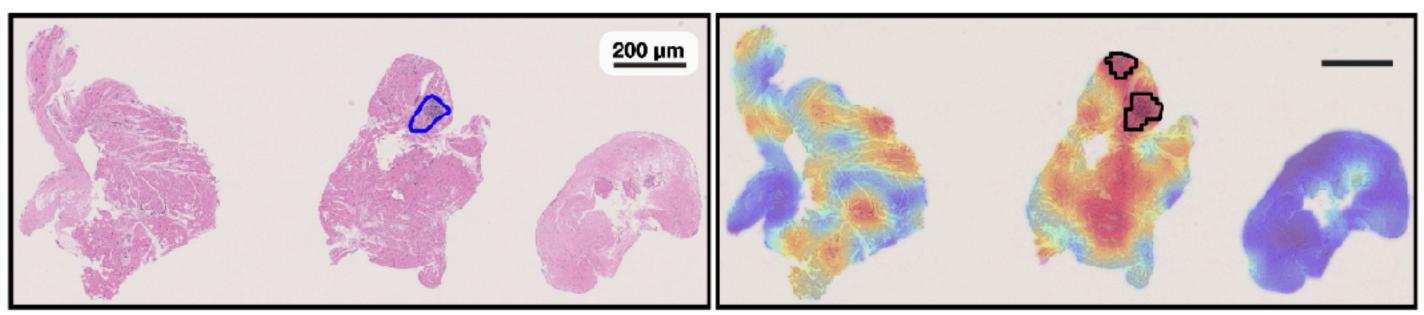
Assessment of Failure Cases



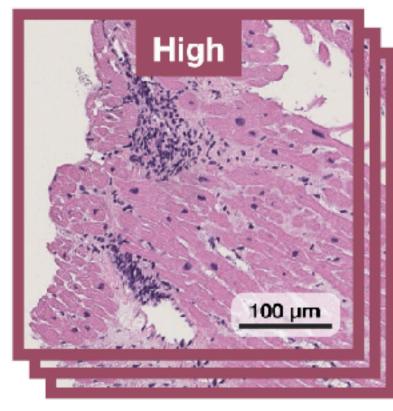
Quantitative Assessment of Interpretability

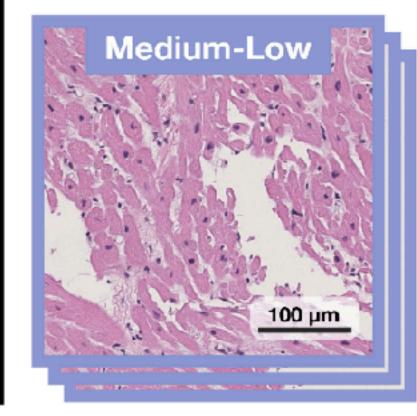


f. Pathologist annotation



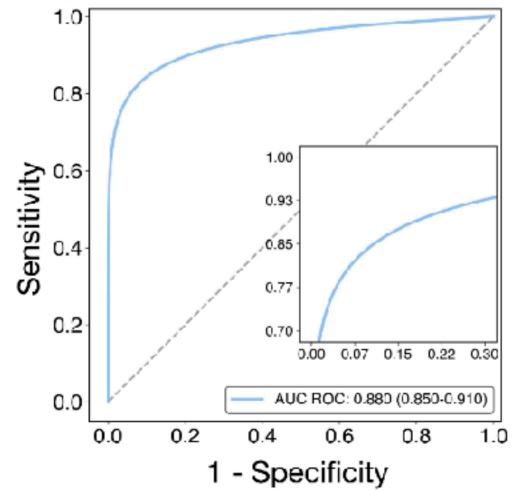
c. Patches





g. High-attention regions

d. Diagnostic Relevance

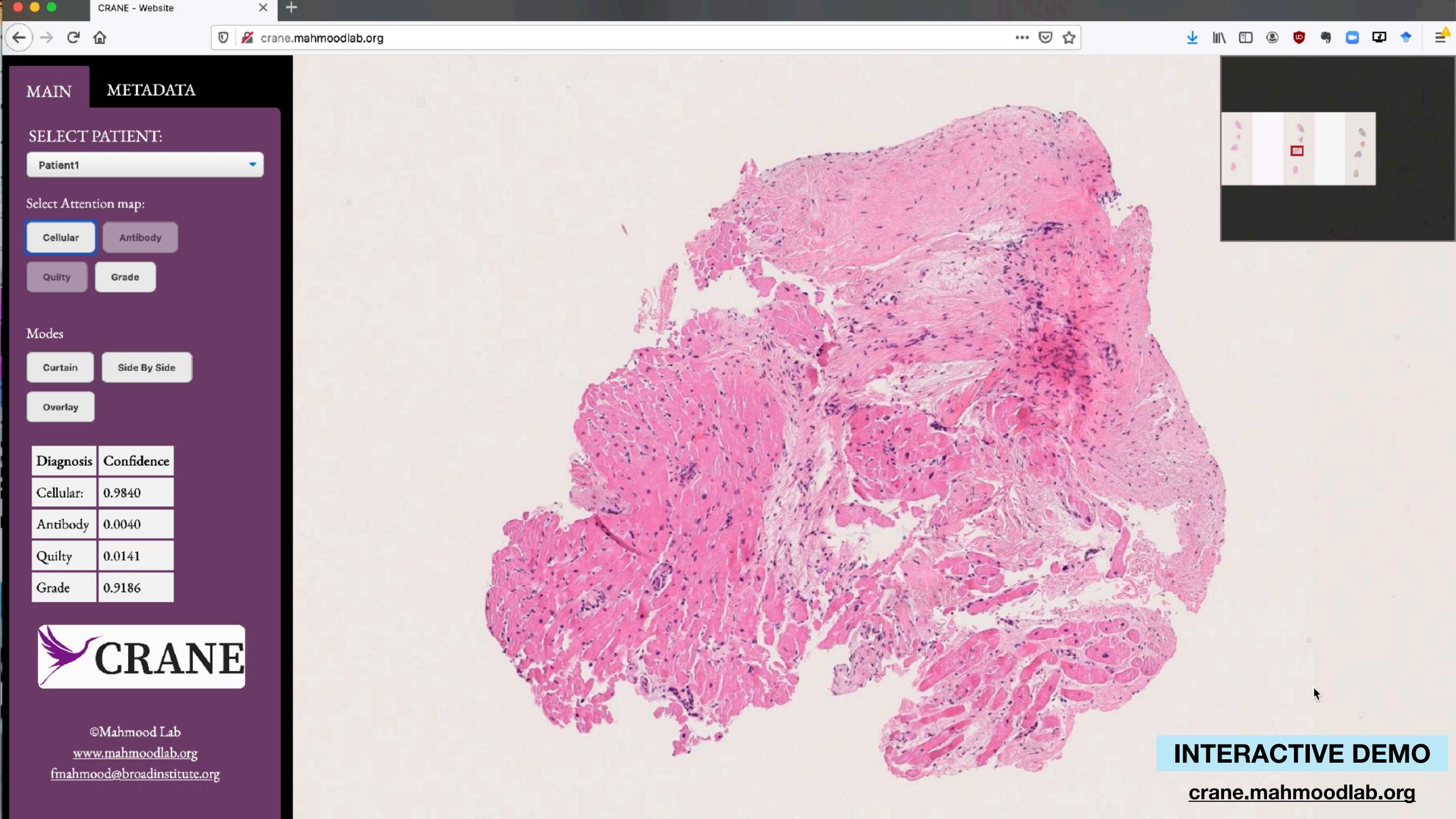


e. Patch-Level Scores

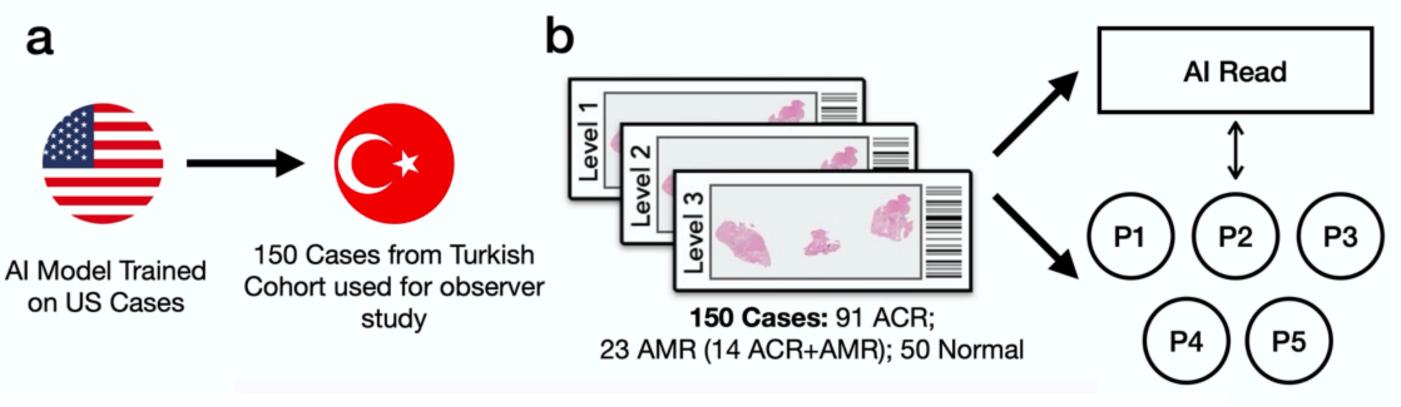
Tasks:	Accuracy	F1	κ
All	0.873	0.855	0.744
Cellular	0.925	0.914	0.848
Antibody	0.902	0.911	0.802
Quilty	0.809	0.729	0.596

h. Slide-Level Scores

Tasks:	Detection rate
All	0.922
Cellular	0.942
Antibody	0.901
Quilty	0.924



Comparison with human readers



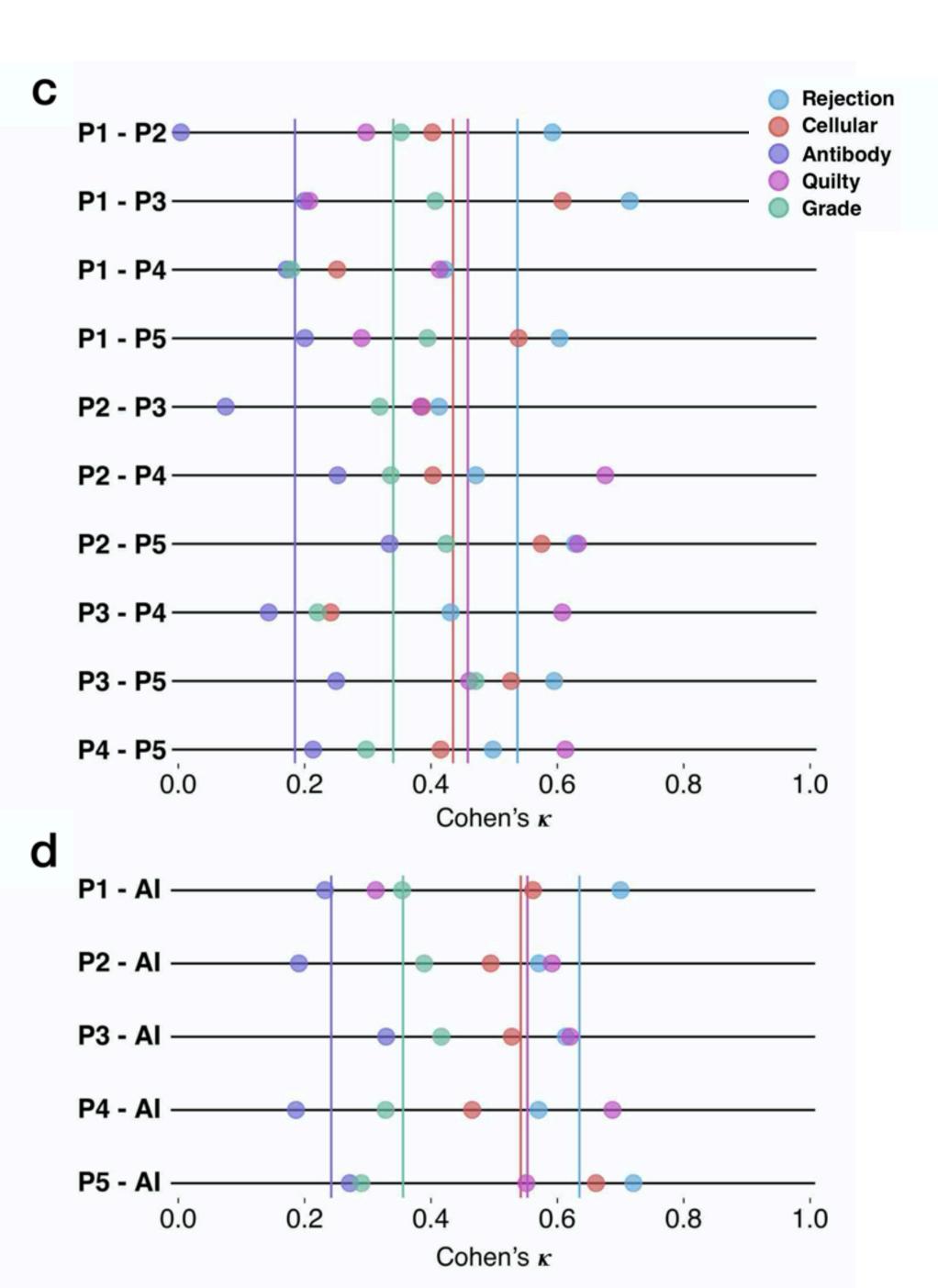
Cohens' κ (-1 to 1): inter-observer agreement: Agreement between expert is comparable to previous studies

For all tasks Al-predictions are not inferior to human experts:

avg. agreement on rejection between pathologists $\kappa = 0.537$ (moderate agreement)

> avg. agreement between **pathologists and model** $\kappa = 0.639$ (substantial agreement)

(avg. 10.5 years of experience)



Clinical Potential

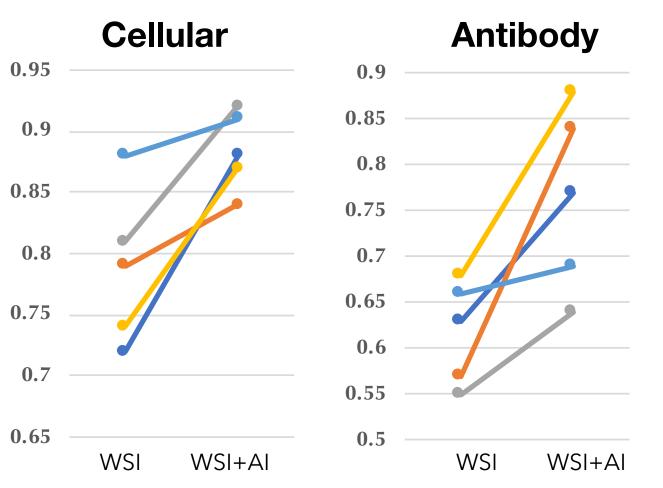
Ground-truth labels:

consensus of readers from the first study

Al-assistance:

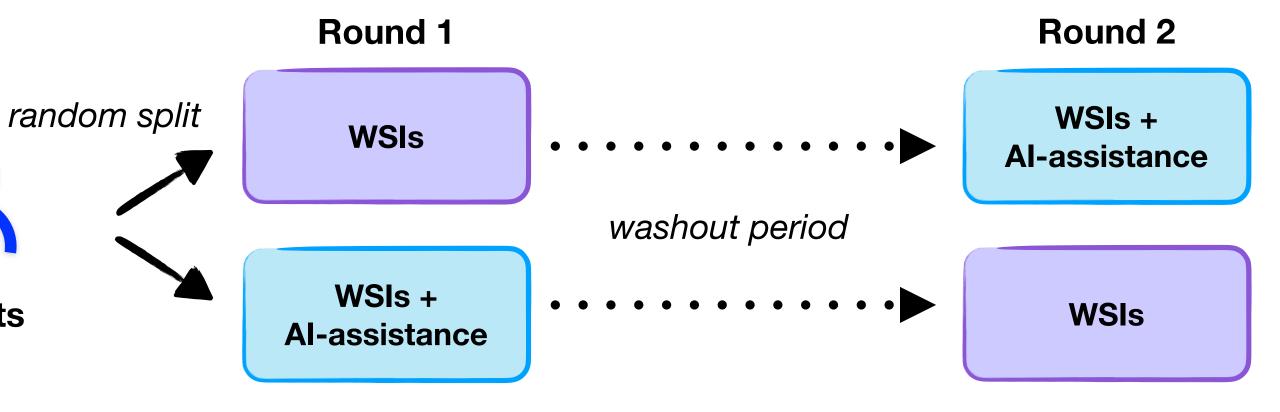
attention heatmaps as semitransparent layer at the top of H&E slide

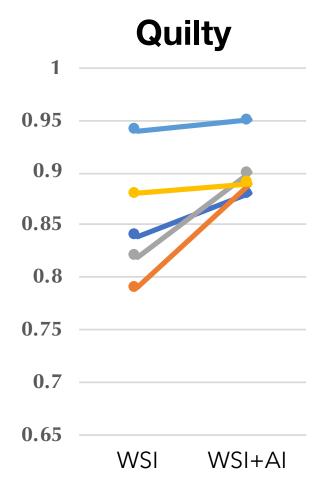
Pathologists

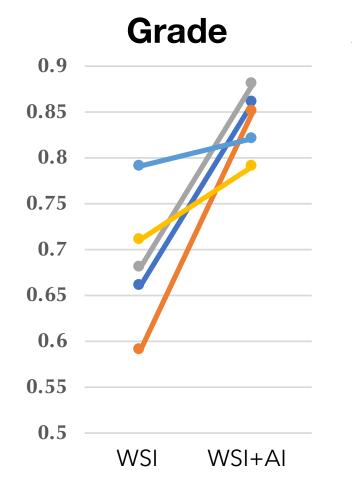


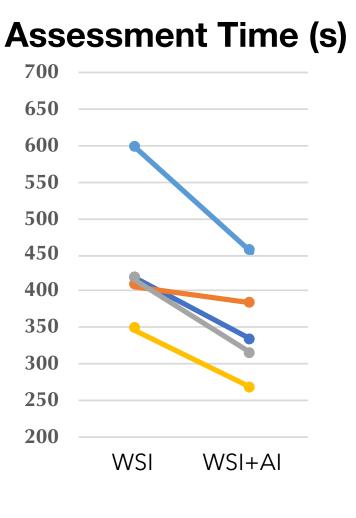
For all readers:

- **Increase accuracy**
- (i.e. reduce inter-rater variability)
- **Decrease assessment time**



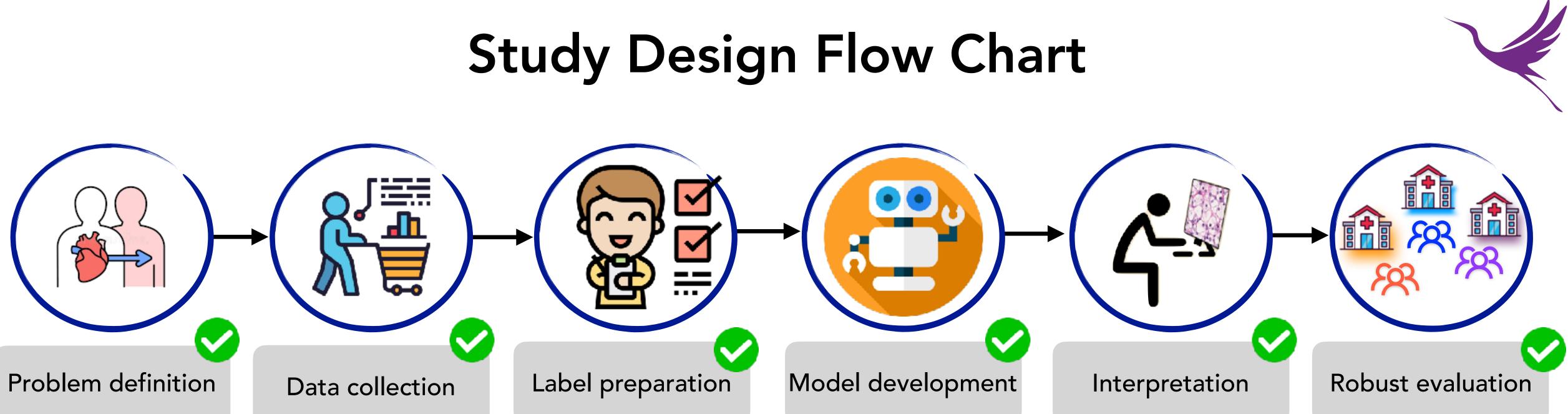




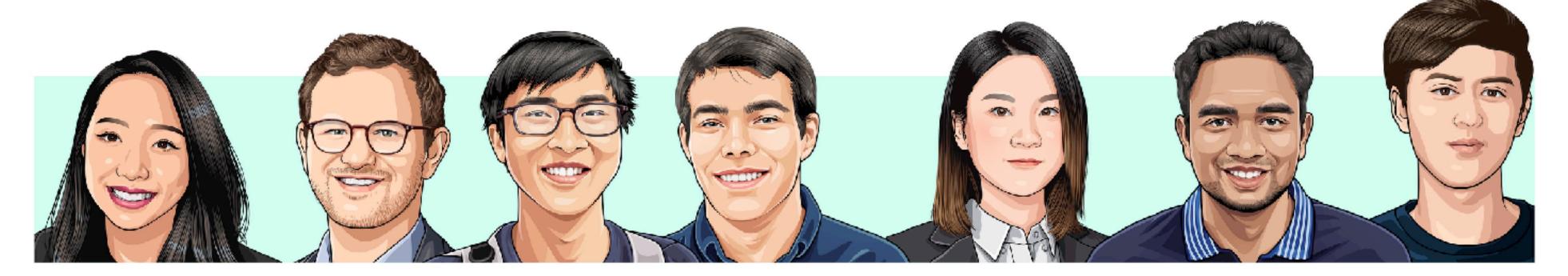


Lipkova et al. Nature Medicine (2022)













The Mahmood Lab







BRIGHAM AND WOMEN'S HOSPITAL

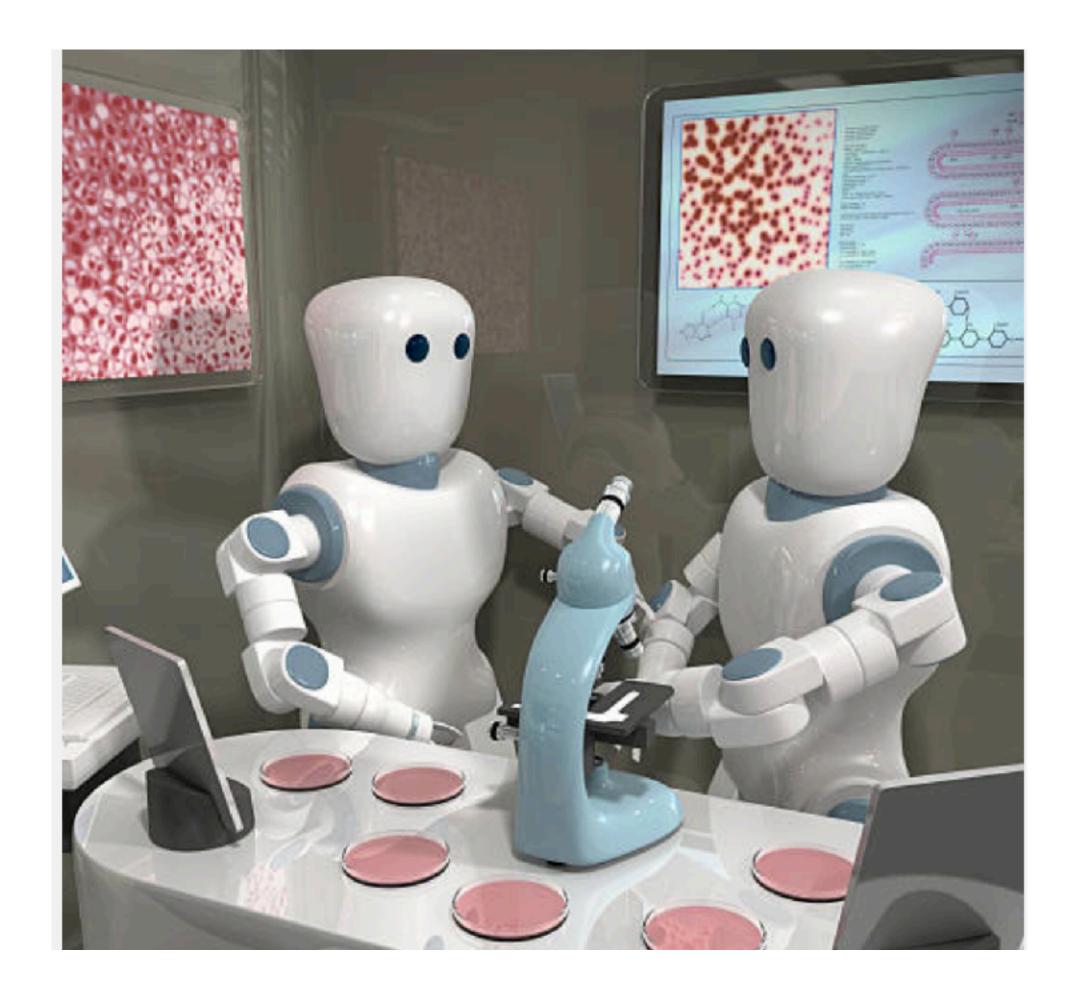






WERE HIRING!







UC Irvine Health School of Medicine