



## Machine Learning for Cancer Diagnostics, Prognostics and Prediction

### Fayyaz Minhas

Associate Professor, Department of Computer Science

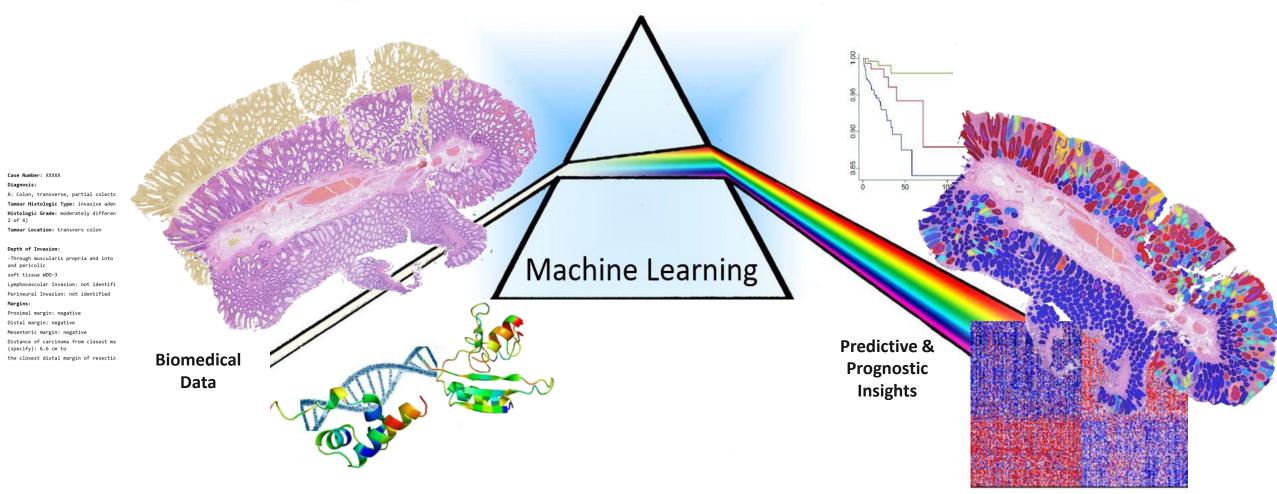
Deputy Director, Tissue Image Analytics Centre

Warwick Cancer Research Centre

University of Warwick



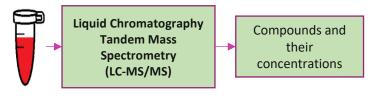




AI and ML to help discover and understand biology and pathology



• Al for Biology



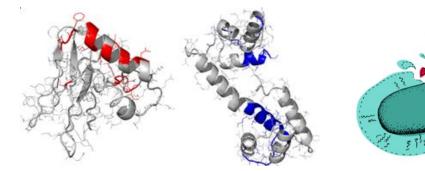
RAMClust: Spectral-matching-based annotation for metabolomics data

Nucleic Acids Research, 2020 1 doi: 10.1093/nar/gkaa219



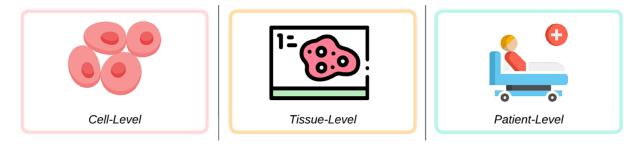
### Machine learning predicts new anti-CRISPR proteins

Simon Eitzinger<sup>1,†</sup>, Amina Asif<sup>2,3,†</sup>, Kyle E. Watters<sup>1,†</sup>, Anthony T. lavarone<sup>4</sup>, Gavin J. Knott<sup>1</sup>, Jennifer A. Doudna <sup>©1,5,6,7,8,9,\*</sup> and Fayyaz ul Amir Afsar Minhas<sup>2,10,\*</sup>



Prediction of Protein Interfaces, Drug-Protein Interactions and Protein Function (Anti-CRISPR proteins, Anti-Microbial & Hemolytic peptides, Amyloid & Prion activity, Phage antibacterial proteins)

### • Al for Pathology & Diagnostics



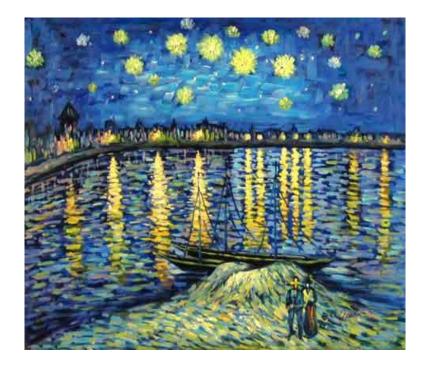


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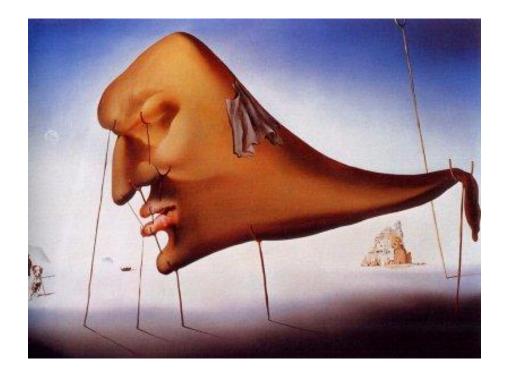
## Paintings by two different painters



## Who's painting is this?



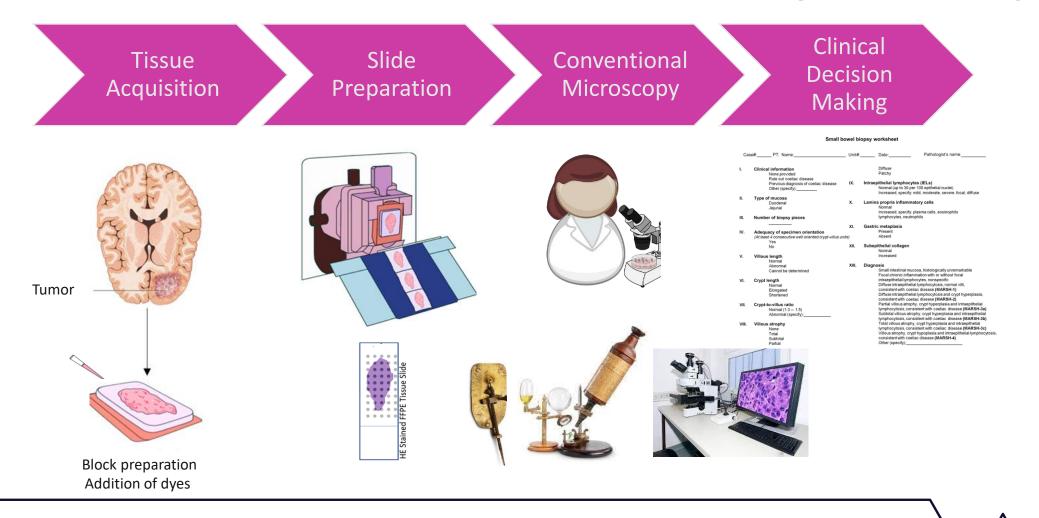
## And this?



*learning from data for generalization to unseen cases* **inductive inference** 

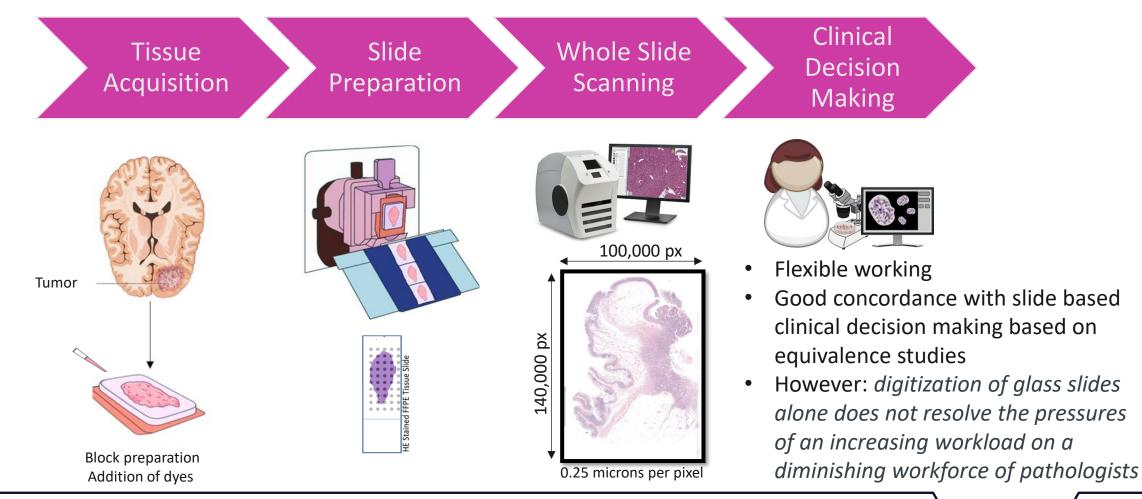
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## Workflow: Conventional Histopathology



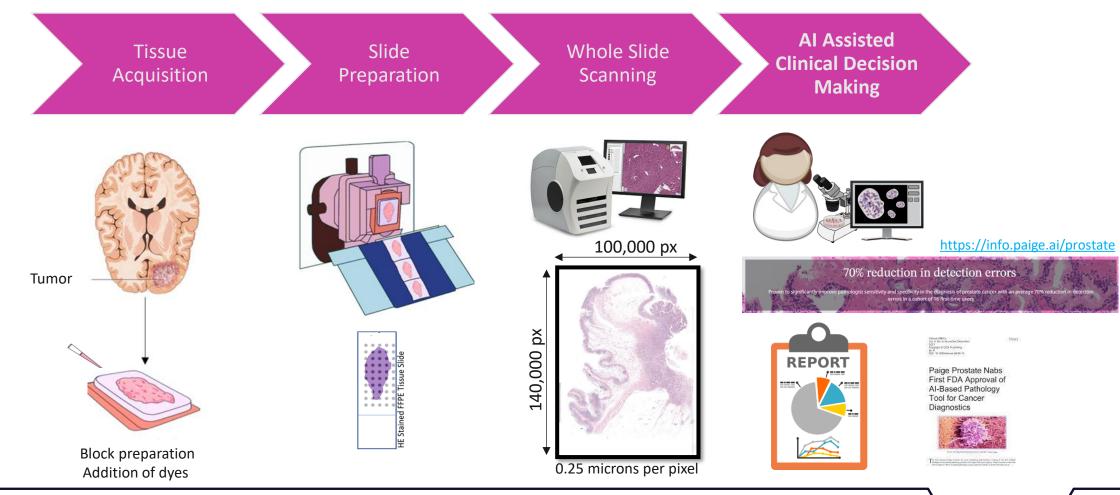
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# Workflow: Digital Pathology

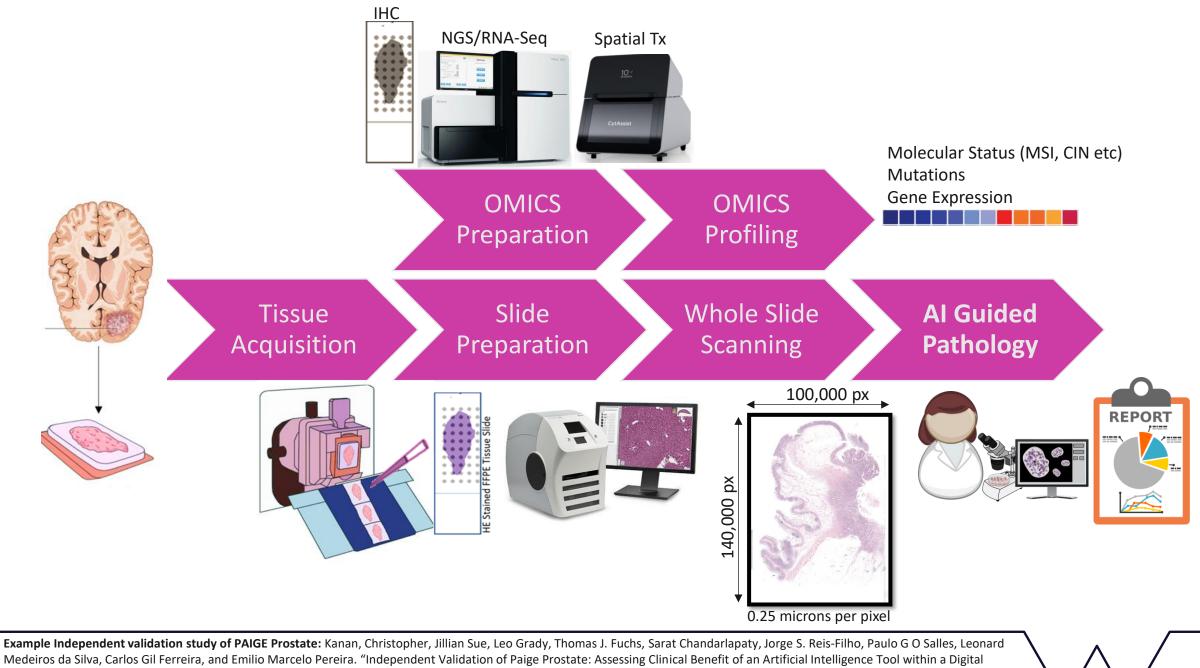


**Example equivalence studies:** Snead, David R. J., Yee-Wah Tsang, Aisha Meskiri, Peter K. Kimani, Richard Crossman, Nasir M. Rajpoot, Elaine Blessing, et al. "Validation of Digital Pathology Imaging for Primary Histopathological Diagnosis." *Histopathology* 68, no. 7 (June 2016): 1063–72. <u>https://doi.org/10.1111/his.12879</u>. Hanna, Matthew G., Victor E. Reuter, Meera R. Hameed, Lee K. Tan, Sarah Chiang, Carlie Sigel, Travis Hollmann, et al. "Whole Slide Imaging Equivalency and Efficiency Study: Experience at a Large Academic Center." Modern Pathology 32, no. 7 (July 2019): 916–28. <u>https://doi.org/10.1038/s41379-019-0205-0</u>.

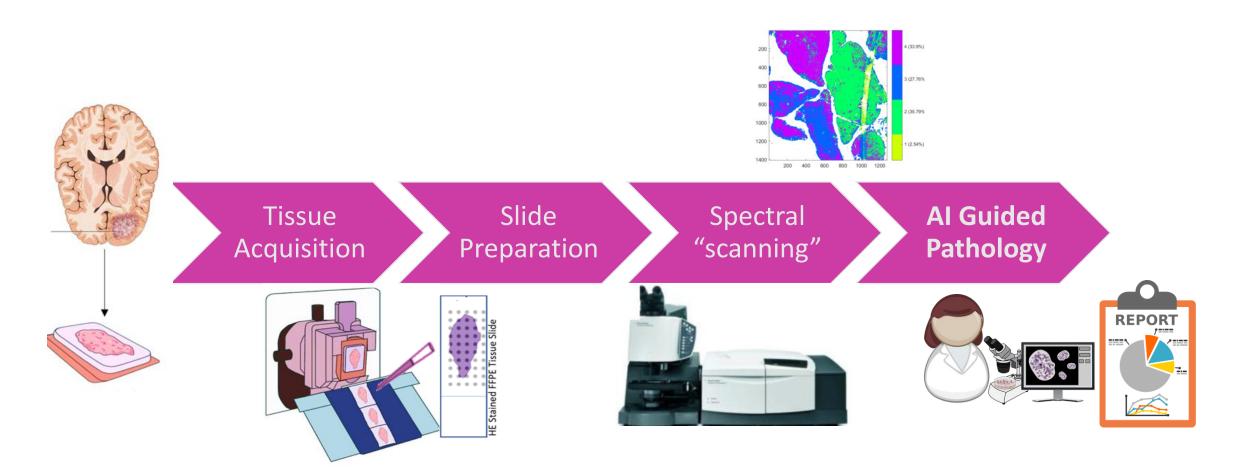
# Workflow: Computational Pathology



**Example Independent validation study of PAIGE Prostate:** Kanan, Christopher, Jillian Sue, Leo Grady, Thomas J. Fuchs, Sarat Chandarlapaty, Jorge S. Reis-Filho, Paulo G O Salles, Leonard Medeiros da Silva, Carlos Gil Ferreira, and Emilio Marcelo Pereira. "Independent Validation of Paige Prostate: Assessing Clinical Benefit of an Artificial Intelligence Tool within a Digital Diagnostic Pathology Laboratory Workflow." Journal of Clinical Oncology 38, no. 15\_suppl (May 20, 2020): e14076–e14076. https://doi.org/10.1200/JCO.2020.38.15\_suppl.e14076.

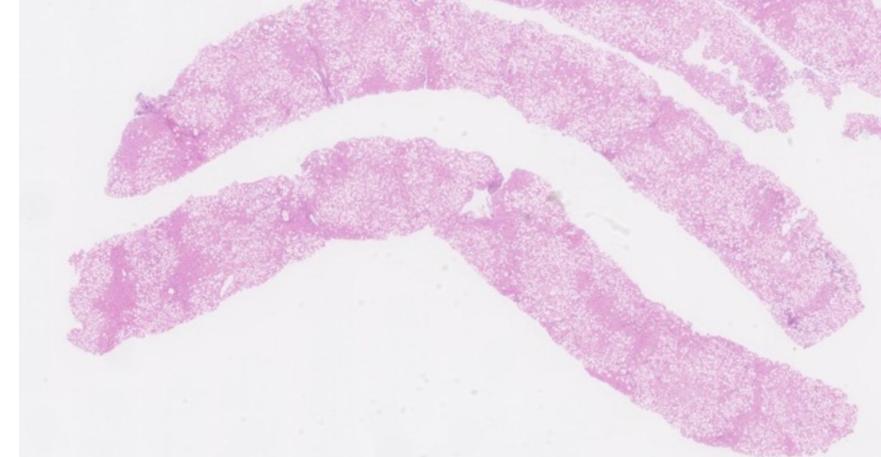


Medeiros da Silva, Carlos Gil Ferreira, and Emilio Marcelo Pereira. "Independent Validation of Paige Prostate: Assessing Clinical Benefit of an Artificial Intelligence Tool within a Digital Diagnostic Pathology Laboratory Workflow." Journal of Clinical Oncology 38, no. 15\_suppl (May 20, 2020): e14076–e14076. https://doi.org/10.1200/JCO.2020.38.15\_suppl.e14076.

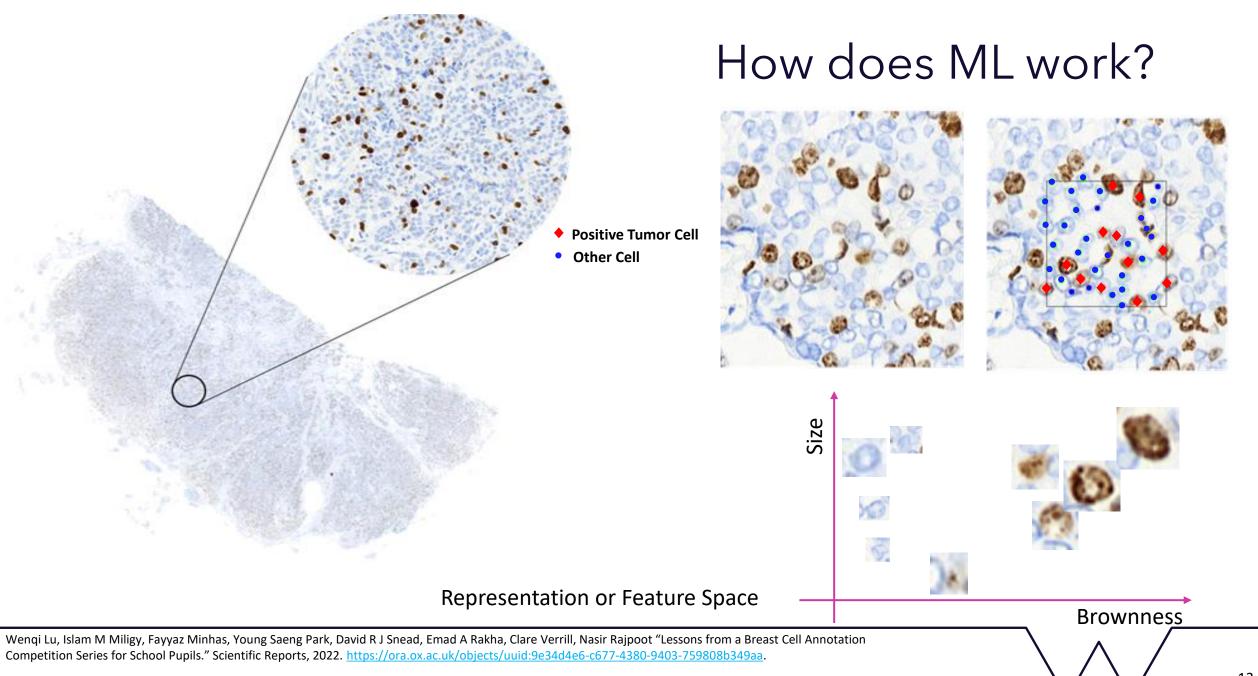


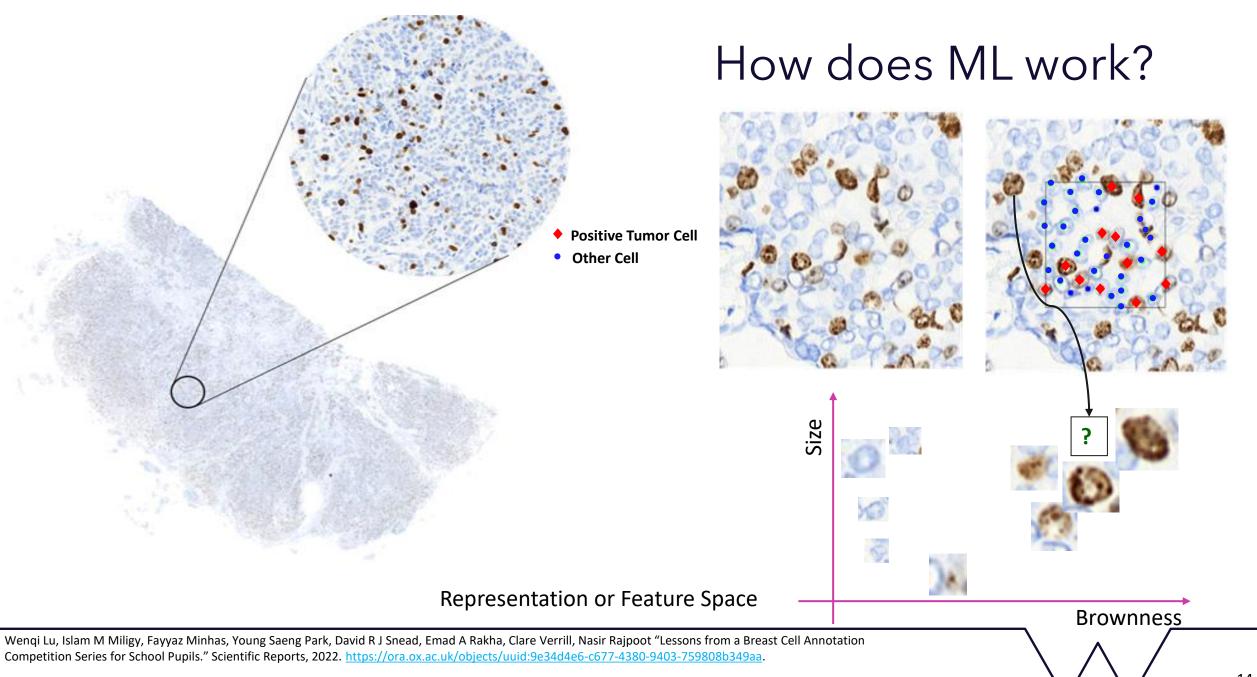
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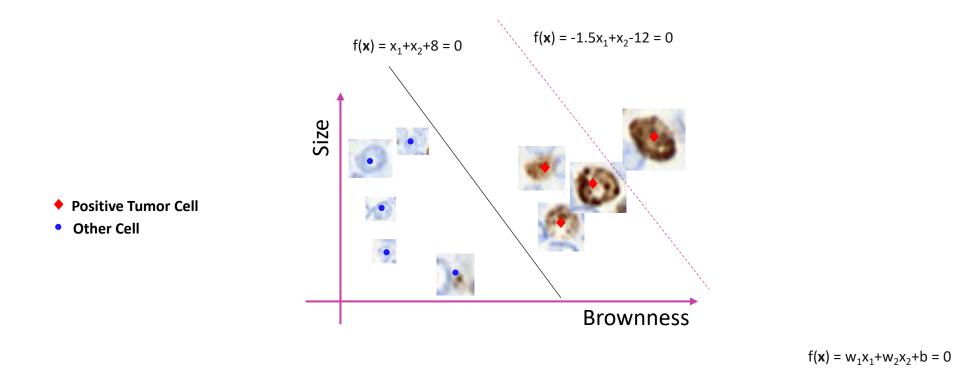
# Why is applying AI to pathology a good idea?



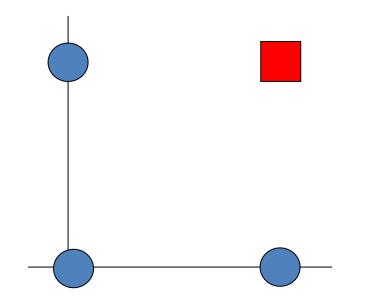
What percentage of cells in this sample are fat cells?







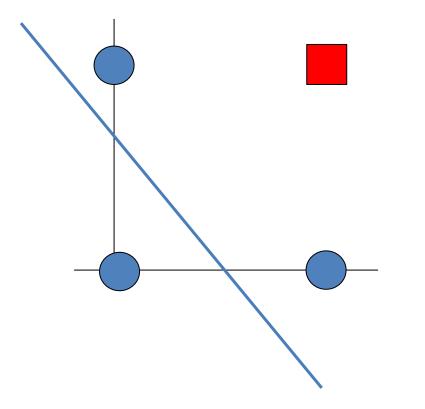
## Exercise



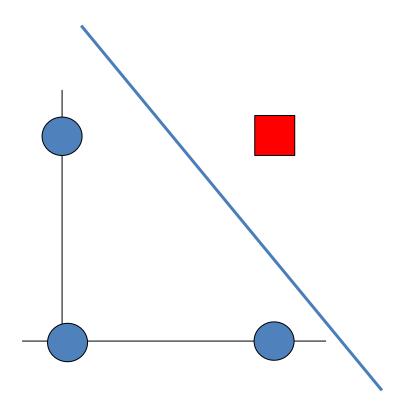
Deep Learning: From Philosophy to AGI

16 University of Warwick

## Exercise



## Exercise

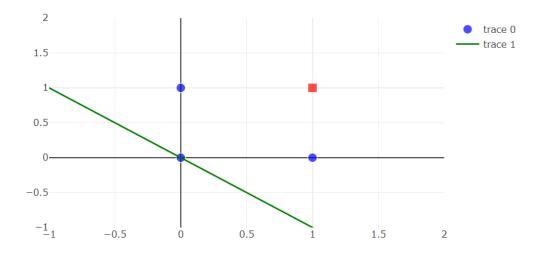


Deep Learning: From Philosophy to AGI

## Doing it interactively



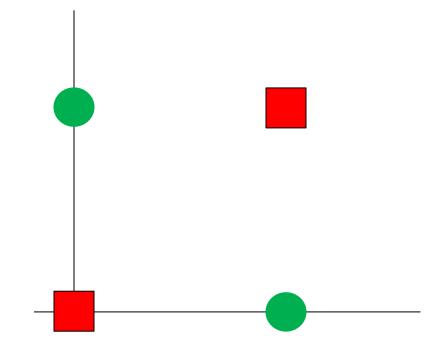
AND Classification Problem



https://foxtrotmike.github.io/CS909/AND-NEURON.html

**Deep Learning: From Philosophy to AGI** 

## But what if we have a linearly inseparable problem?



**Deep Learning: From Philosophy to AGI** 

Machine learning and deep learning involve discovering meaningful representations of input data and then using these representations to partition data for various tasks

an almost irresistible claim for damages for negligence. KENNETH S MULLARD

Bletchingdon, Oxon

#### BRITISH MEDICAL JOURNAL VOLUME 282 14 FEBRUARY 1981

#### Staffing crisis in pathology

SIR,-We are writing to express concern that the methods of teaching pathology to undergraduate medical students are both qualitatively and in outlook inappropriate for current demands. Pathology is the study of disease and as such is the basis of all medical practice. It is the one subject which encourages the strict discipline of the scientific method together with the empirical art of medicine to explain the clinically observed patterns of disease behaviour. There is an increasingly scientific approach to medicine which needs to be reflected in and emphasised by expansion of undergraduate teaching of pathology. The size of undergraduate classes receiving instruction in pathology is generally increasing and there is a serious lack of first-hand experience of practical pathology. At a debate during the recent Pathological Society meeting (9 January) many of the senior academics admitted that large classes were exerting an increasingly detrimental effect on the teaching of pathology. As a direct consequence pathology teaching is in a decline; and the subject is becoming, apparently, more remote from the clinical practice of medicine.

We are concerned that the input to pathology training posts should be not only quantitatively but also qualitatively optimal to meet future demands. We strongly support Professors J R Anderson and J R Tighe (15 November, p 1370), who have pointed out that, although the immediate problem in pathology is a scarcity of suitably trained individuals for the large number of consultant posts shortly to become vacant through retirement, the problem in the future will become compounded by the shortage of SHO and registrar posts to feed the senior registrar grade. However, further factors should be considered and include that undefined proportion of those entering the junior grades who will not complete their higher specialist training in pathology. We also feel that it is extremely important that sufficient posts should be available within pathology departments to provide opportunities for those primarily working in other specialties to benefit from a first-hand experience of pathology. Therefore we further emphasise the urgent need for an increase in posts at the SHO and registrar grade. It is important, however, to avoid focusing on this single measure as being the answer to the future security of pathology. It is evident to us, from personal communications with the directors of pathology in many of the undergraduate teaching departments, that there is widespread concern to provide undergraduates with a proper exposure to pathology. We believe that medical students should have the opportunity to receive practically based tutorials in all branches of pathology. Pathology should be taught as being directly relevant to the clinical situation. In particular, we believe that the general principles of pathology should be introduced from the earliest stage of the undergraduate medical training. In addition, the teaching of pathology should be combined with the teaching of anatomy, histology, physiology, and biochemistry. Symington has stressed the

# Why is applying ML to pathology a good idea? <u>Business Need</u>

35%

30%

25%

waiting 15%

Percent v

5%

2009 2010

days

>62

- Pathologist Recruitment
- Aging workforce
- Subjectivity
- Time & Effort in Quantification

"... workforce census from the Royal College of Pathologists showed that a quarter of all histopathologists are over 55, most of whom are expected to retire by 2023. Furthermore, an all-time low number of trainee doctors are choosing to specialize in pathology with only 3 percent of NHS histopathology departments having enough staff to meet clinical demand."



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2015 2015 2016 2016 2016



### HISTOPATHOLOGY WORKFORCE SURVEY

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Cancer waiting

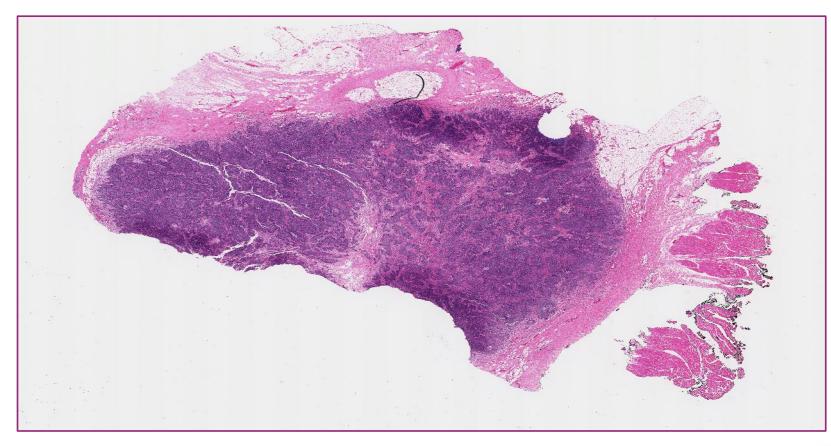
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# Why is applying ML to pathology a good idea?



Given training data, Can we predict:

> Normal or Abnormal? Classification Grade? Regression **Tumour Regions?** Segmentation Quantifying cells Quantification **Receptor Status?** Classification **Mutation Status?** Classification Survival? Drug Response? Similar images? Retrieval **Biomarkers?** Pattern Discovery

### Spectrum of Mundanity



IHC scoring Finding tumor Mitotic counting

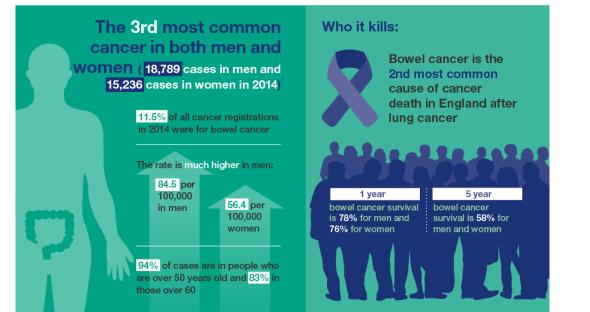
Outcome based subtyping Response to therapy Mutation status MSI positive HPV positive

## A case study

# The problem

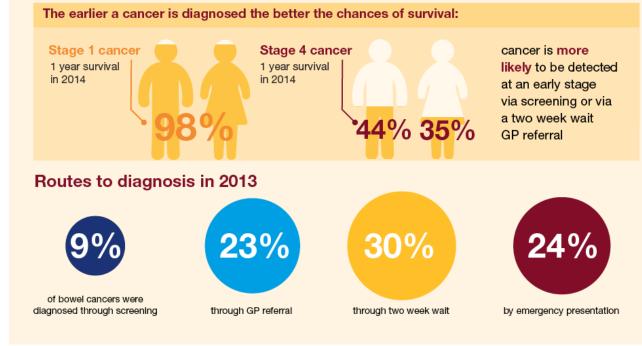
Colorectal Cancer

#### Who it affects



### Early detection of bowel cancer

#### Diagnosis

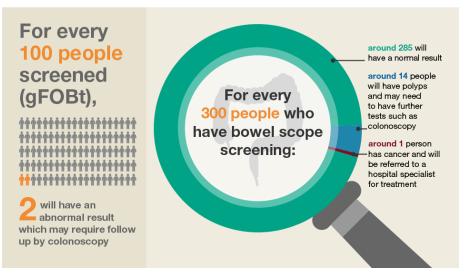


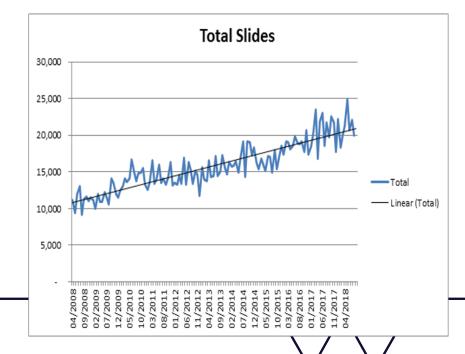
https://www.gov.uk/government/publications/health-matters-preventing-bowelcancer/health-matters-improving-the-prevention-and-detection-of-bowel-cancer

# **Bowel Cancer Screening**

- In England, screening is offered every 2 years for men and women aged 60 to 74 (or above on request)
- gFOBt test (replaced by Faecal Immunochemical Test (FIT)
- Followed by Colonoscopy
- Pathology Examination
  - Endoscopic biopsies from the large bowel account for 8% of the requests in NHS cellular pathology laboratories.
  - Of these samples, approximately 40% are reported as normal

What happens after screening?





# PathLAKE CoBI Project

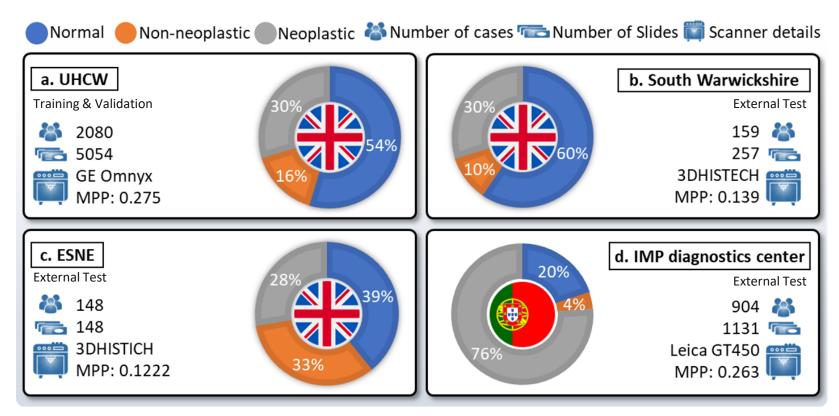
- WIBGI
  - We could "screen out" or "rule out" abnormalities using AI based computational pathology algorithms using whole slide images as input
- What do we need for this?
  - Training and validation data with labels
  - Develop algorithms



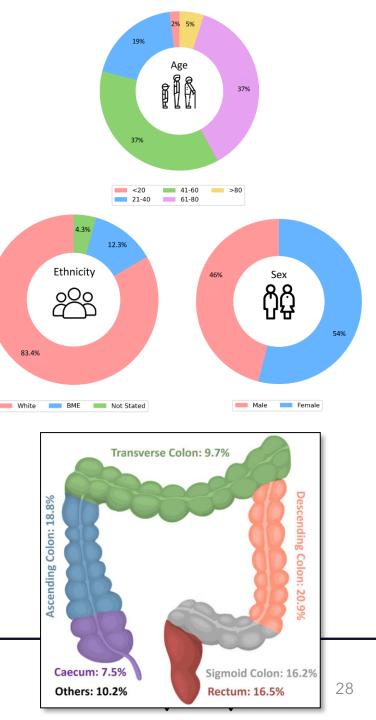
Special thanks to all the funders, researchers and collaborators for support and involvement in the work presented on these slides



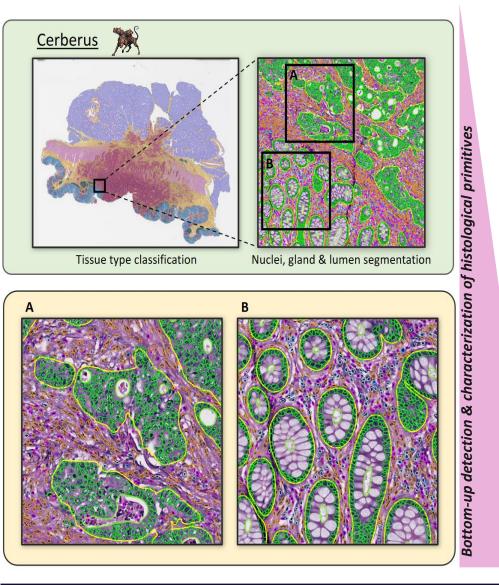
## Data



UHCW used as primary development set colorectal (colonoscopy) biopsies were audited over a 5-year period from 2012 to 2017, reviewed by 13 pathologists with diagnostic consensus



### Two Flavours of CPath



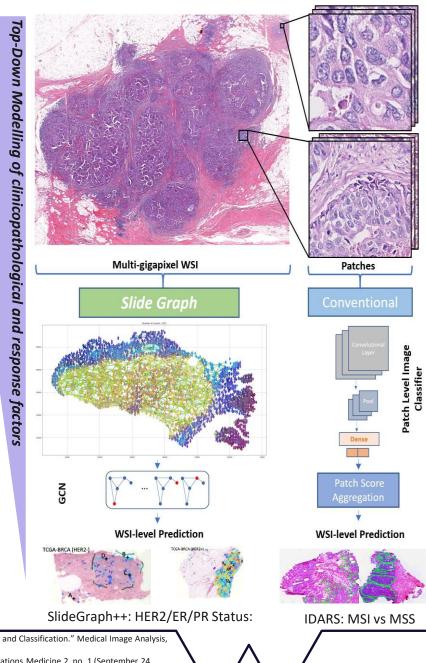
Pros and Cons (for Top-Down Style) Lesser annotations needed Multi-modality integration Versatility of scientific questions

Large(r) amount of WSIs Poor interpretability & explainability Robustness Impact of confounders



- Feature Mining
- Segmenting histological structures
- Detecting nuclei and Classifying cells
- Getting annotations (nuclear, regional)





Graham, Simon, Quoc Dang Vu, Mostafa Jahanifar, Shan E. Ahmed Raza, Fayyaz Minhas, David Snead, and Nasir Rajpoot. "One Model Is All You Need: Multi-Task Learning Enables Simultaneous Histology Image Segmentation and Classification." Medical Image Analysis, November 11, 2022, 102685. https://doi.org/10.1016/j.media.2022.102685.

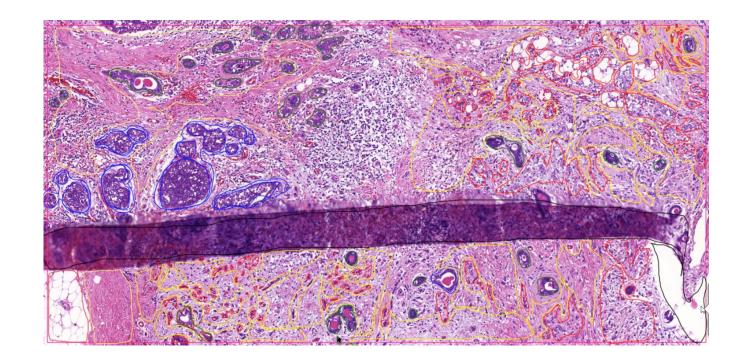
Pocock, Johnathan, Simon Graham, Quoc Dang Vu, Mostafa Jahanifar, Srijay Deshpande, Giorgos Hadjigeorghiou, Adam Shephard, et al. "TIAToolbox as an End-to-End Library for Advanced Tissue Image Analytics." Communications Medicine 2, no. 1 (September 24, 2022): 1–14. https://doi.org/10.1038/s43856-022-00186-5.

Lu, Wenqi, Michael Toss, Emad Rakha, Nasir Rajpoot, and Fayyaz Minhas. "SlideGraph+: Whole Slide Image Level Graphs to Predict HER2Status in Breast Cancer." ArXiv:2110.06042 [Cs], October 12, 2021. <u>http://arxiv.org/abs/2110.06042</u>. Bilal, Mohsin, Shan E. Ahmed Raza, Ayesha Azam, Simon Graham, Mohammad Ilyas, Ian A. Cree, David Snead, Fayyaz Minhas, and Nasir M. Rajpoot. "Development and Validation of a Weakly Supervised Deep Learning Framework to Predict the Status of Molecular 29

Pathways and Key Mutations in Colorectal Cancer from Routine Histology Images: A Retrospective Study." The Lancet Digital Health 3, no. 12 (December 1, 2021): e763–72. https://doi.org/10.1016/52589-7500(21)00180-1.

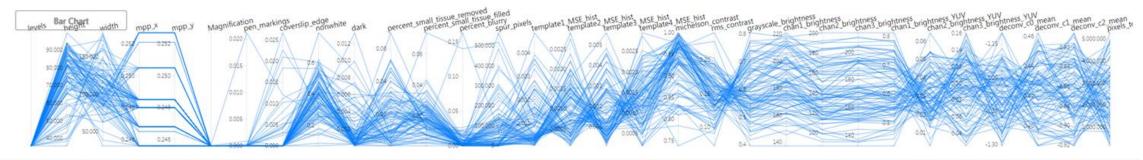
# Pre-processing and Image QC

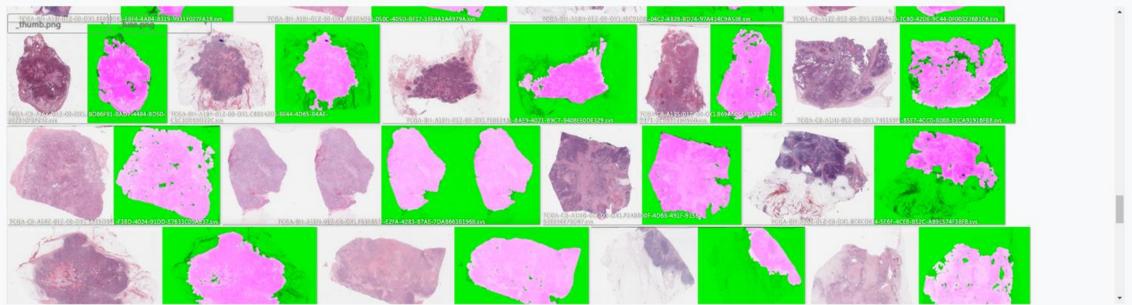
- Stain variation normalization
- Image quality control
  - Blur detection
  - Loss of cover slip
  - Tissue folding
  - Pen Marking



Reset

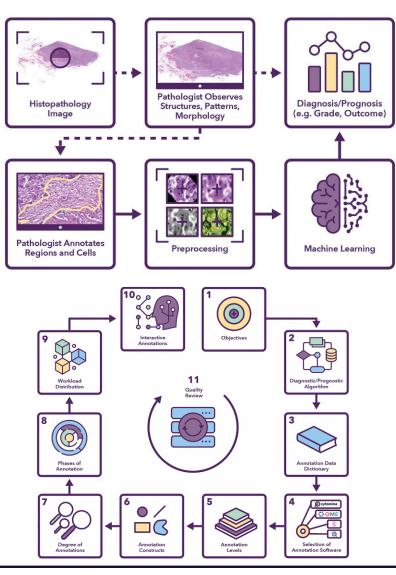
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TCGA-D8-A1JJ-01Z-00-DX1.a986b48f-b295-4d7a-b778-ce829cdf9c38.svs		4	89863	84727	0.248	0.248	40	0	0	0.40365625618646767	0.00038530934490148993	0.02	Delete
TCGA-D8-A1JJ-01Z-00-DX2.7D20F308-7DC6-4367-9459-3AC4C654E7F7.svs		4	74635	83663	0.2527	0.2527	40	0	0.02097230770845992	0.6022809295769842	0.0069632641758784015	0.02	Deselect
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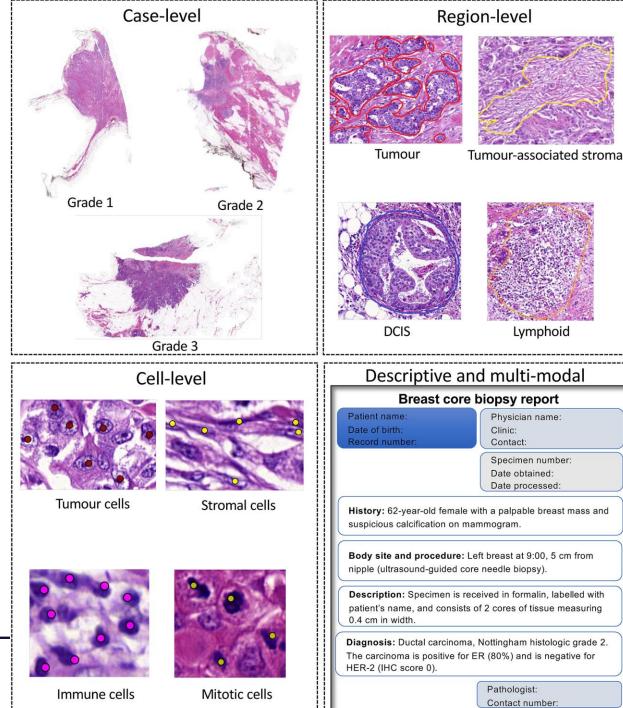


Janowczyk, Andrew, Ren Zuo, Hannah Gilmore, Michael Feldman, and Anant Madabhushi. "HistoQC: An Open-Source Quality Control Tool for Digital Pathology Slides." *JCO Clinical Cancer Informatics* 3 (April 16, 2019). <u>https://doi.org/10.1200/CCI.18.00157</u>.

### Semantic Annotation Guidelines

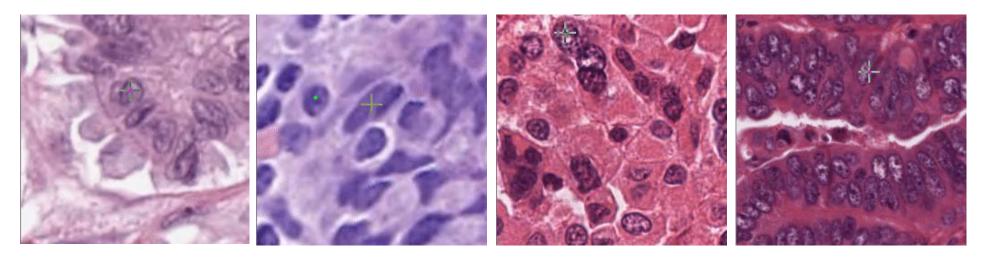


Wahab, Noorul, Islam M Miligy, Katherine Dodd, Harvir Sahota, Michael Toss, Wenqi Lu, Mostafa Jahanifar, et al. "Semantic Annotation for Computational Pathology: Multidisciplinary Experience and Best Practice Recommendations." *The Journal of Pathology: Clinical Research*. <u>https://doi.org/10.1002/cjp2.256</u>.



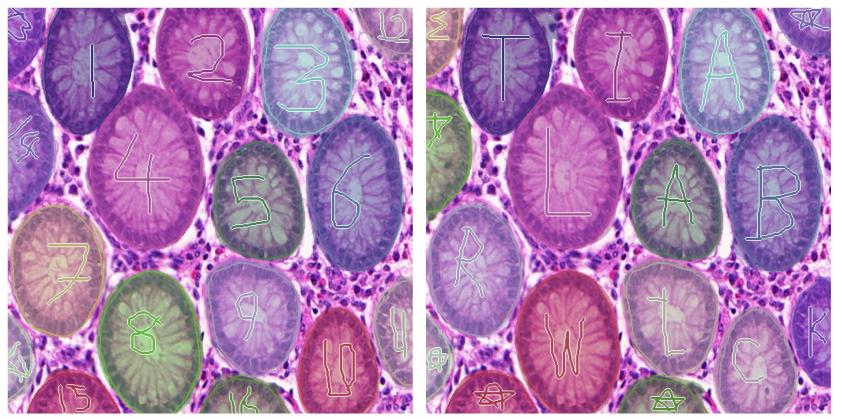
## **AI-Assisted Annotations**

# **NuClick** used for collecting the segmentation masks from point clicks



Koohbanani et al., Medical Image Analysis (Oct 2020) & Gamper et al., Arxiv (Apr 2020)

## **NuClick+**: Al-Assisted Annotation of Glands

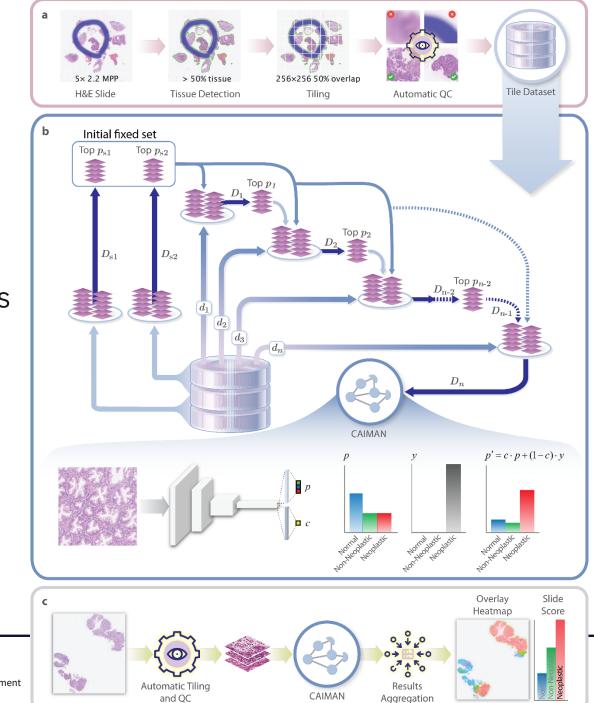


Koohbanani et al., Medical Image Analysis (Oct 2020)

## CAIMAN

Colorectal AI Model for Abnormality detectioN

- A weakly supervised top-down approach
  - Using whole slide image level labels for training
  - Based on our previous method for MSI detection in CRC

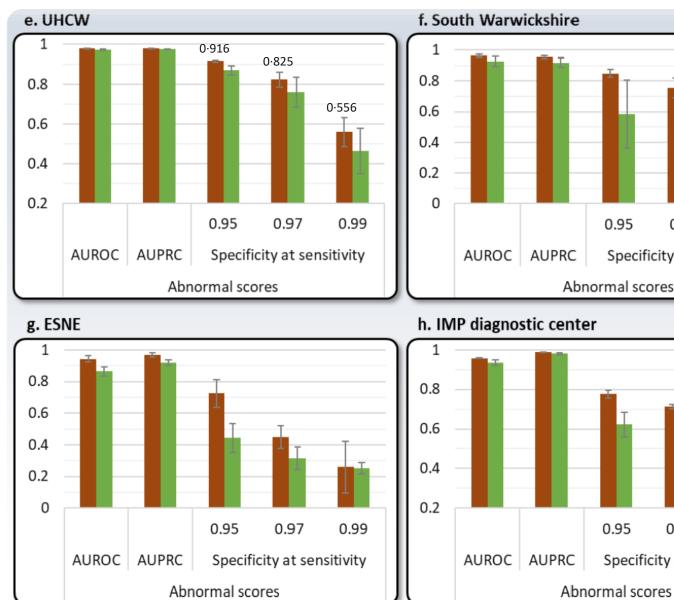


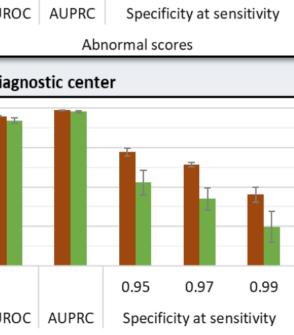
Bilal, Mohsin, Yee Wah Tsang, Mahmoud Ali, Simon Graham, Emily Hero, Noorul Wahab, Katherine Dodd, et al. "Development and Validation of Artificial Intelligence-Based Prescreening of Large-Bowel Biopsies Taken in the UK and Portugal: A Retrospective Cohort Study." The Lancet Digital Health 5, no. 11 (November 1, 2023): e786–97. https://doi.org/10.1016/S2589-7500(23)00148-6. Bilal, Mohsin, Shan F, Ahmed Raza, Avesha Azam, Simon Graham, Mohammad Ilvas, Jan A Cree, David Snead, Favvaz Minhas, and Nasir M Baipoot, "Development

Bilal, Mohsin, Shan E Ahmed Raza, Ayesha Azam, Simon Graham, Mohammad Ilyas, Ian A Cree, David Snead, Fayyaz Minhas, and Nasir M Rajpoot. "Development and Validation of a Weakly Supervised Deep Learning Framework to Predict the Status of Molecular Pathways and Key Mutations in Colorectal Cancer from Routine Histology Images: A Retrospective Study." *The Lancet Digital Health*, October 19, 2021. <u>https://doi.org/10.1016/S2589-7500(21)00180-1</u>.

## Results

- Cross-validation
- External Testing
- No statistically significant differences in prediction scores across anatomical sites





0.95

0.97

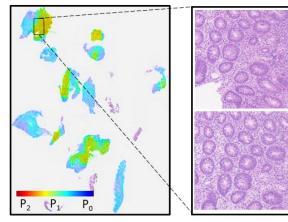
0.99

CAIMAN

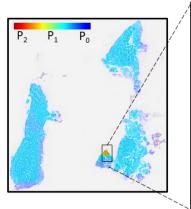
IDaRS

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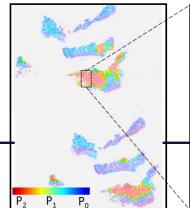
a. Non-neoplastic

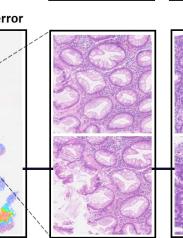


c. Neoplastic small fragment LGD

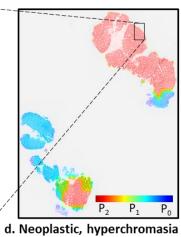


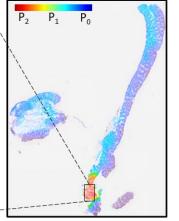
e. Diagnostic category error





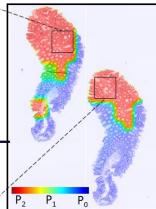


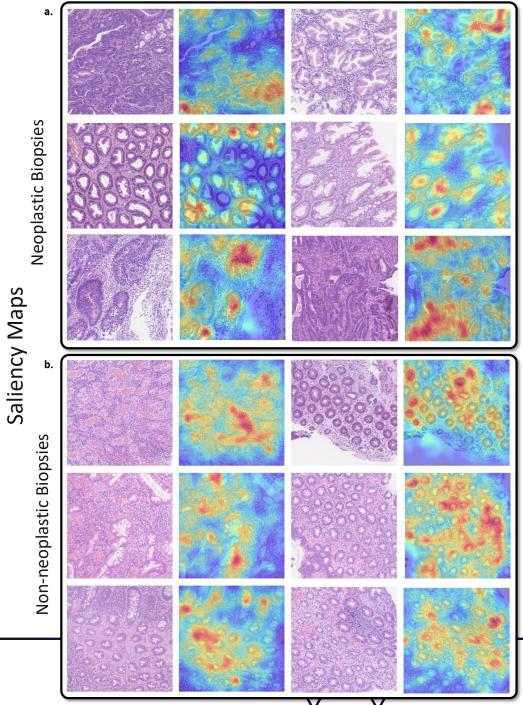


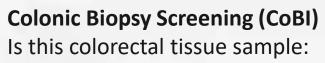


WSI Results

#### f. Diagnostic category error







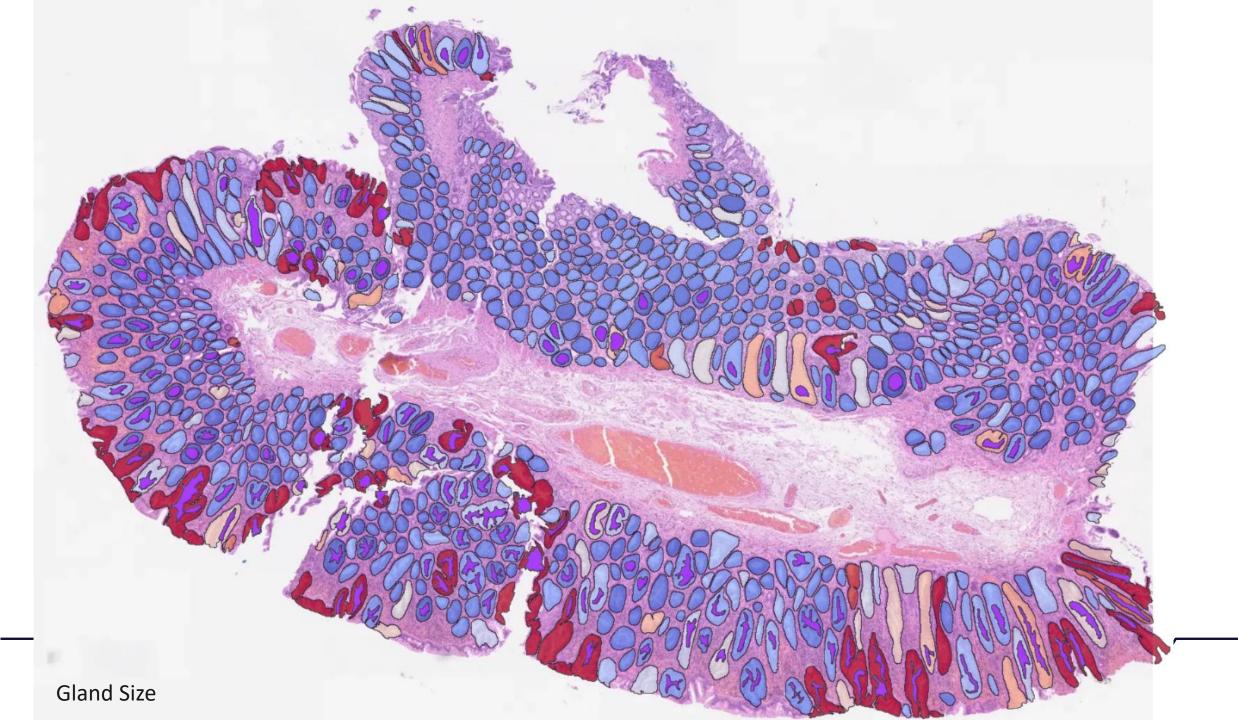


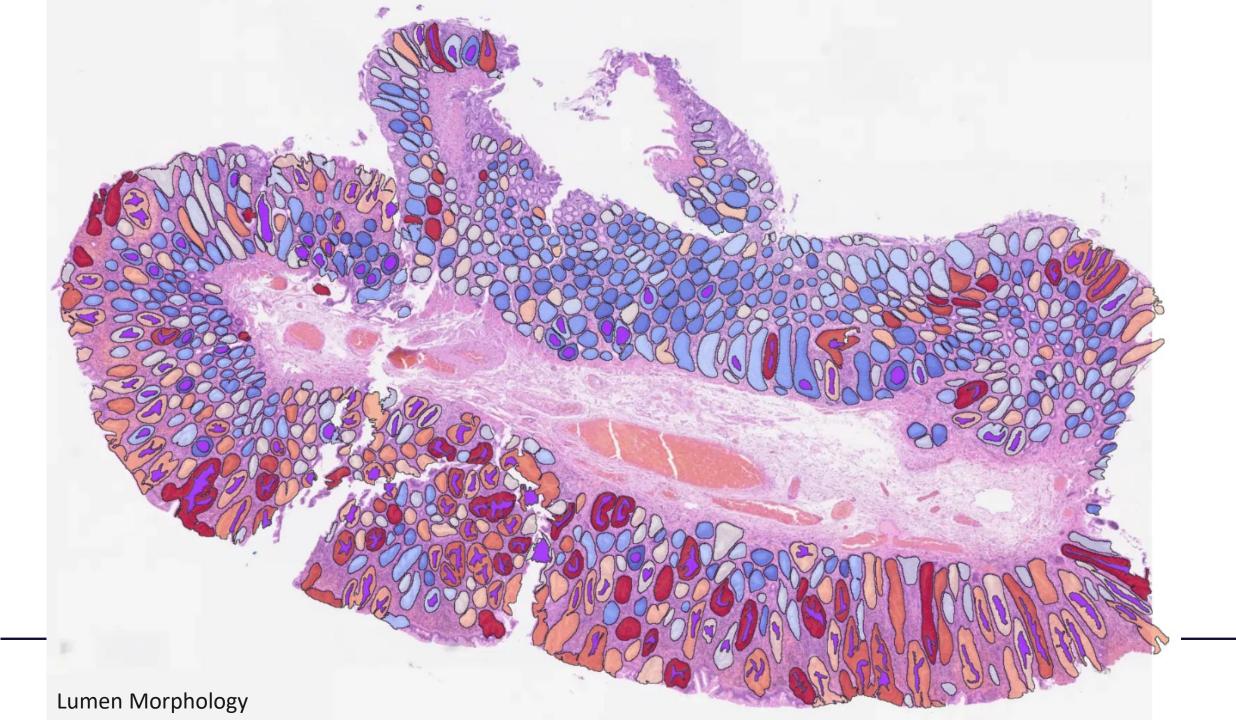
- Normal?
- Abnormal?

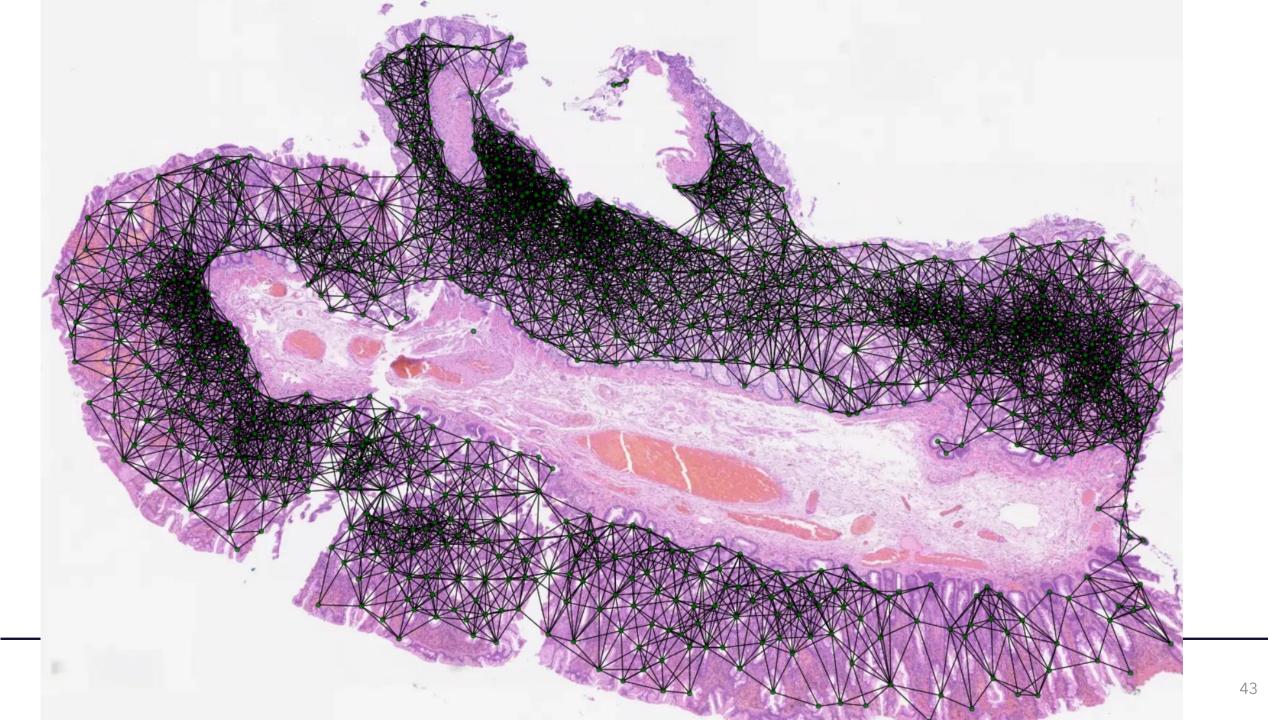
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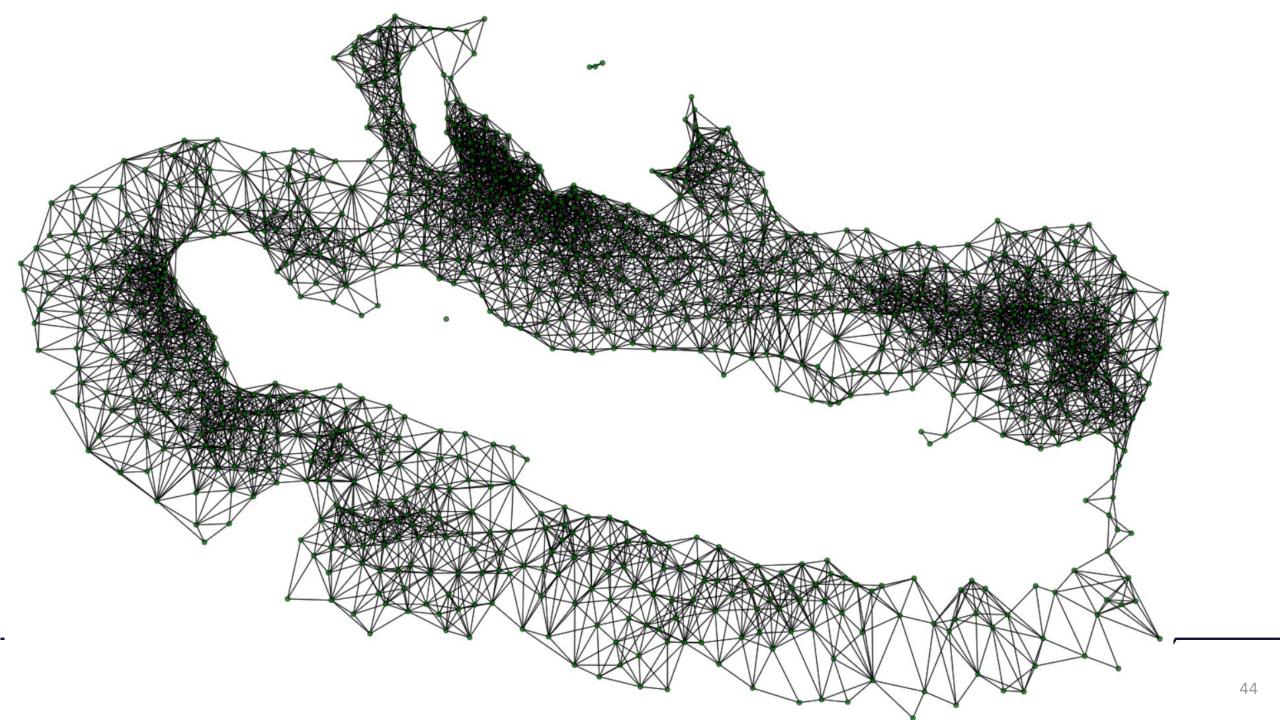
Plasma Cell

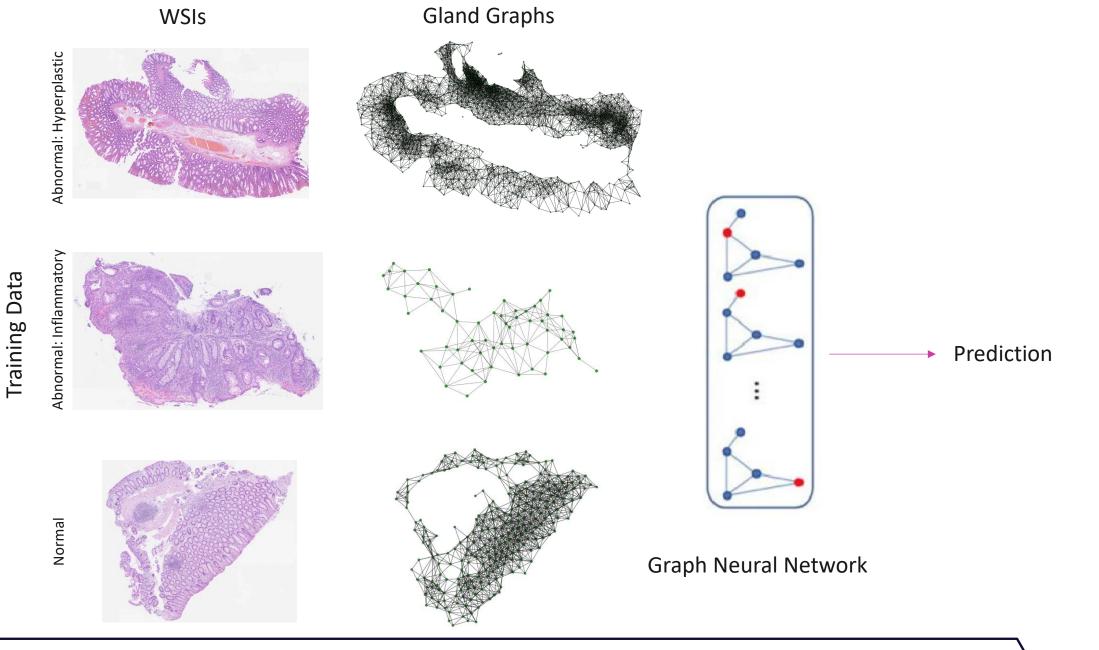
Lumen Eosinophil Neutrophil Gland Epithelial Cell

Lymphocyte Connective Cell 









*Whole Slide Images Are Graphs*, Fayyaz Minhas, 2020. <u>https://www.youtube.com/watch?v=Of1u0i7roS0</u>. https://iguana.dcs.warwick.ac.uk/

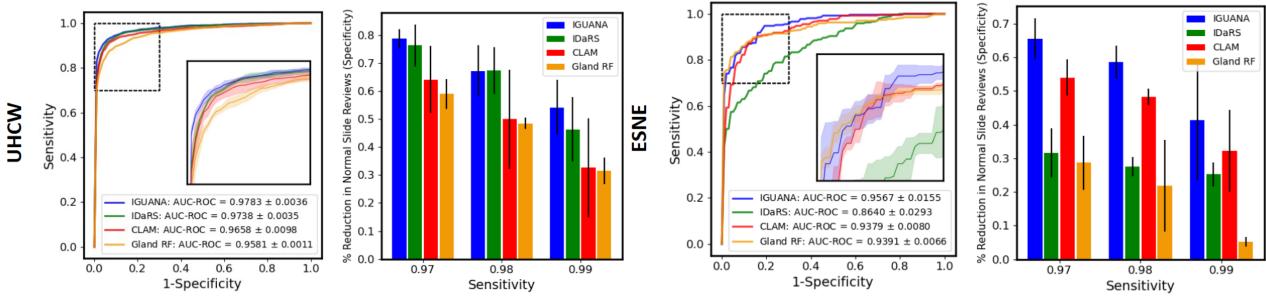
Automated Classification: Abnormal (Hyperplastic)

Leading to up to 33% reduction in review of normal slides at 99% sensitivity

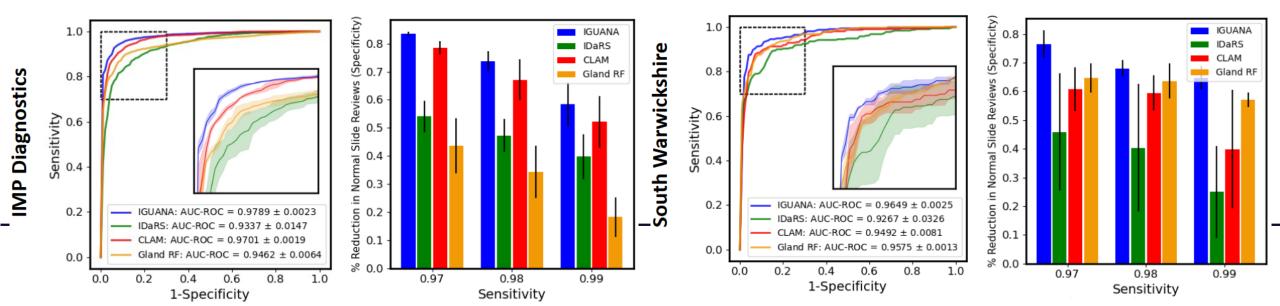


Graham, Simon, Fayyaz Minhas, ... Nasir Rajpoot "Screening of Normal Endoscopic Large Bowel Biopsies with Interpretable Graph Learning: A Retrospective Study." Gut, May 12, 2023. <u>https://doi.org/10.1136/gutjnl-2023-329512</u>.

#### **Internal Cross Validation**

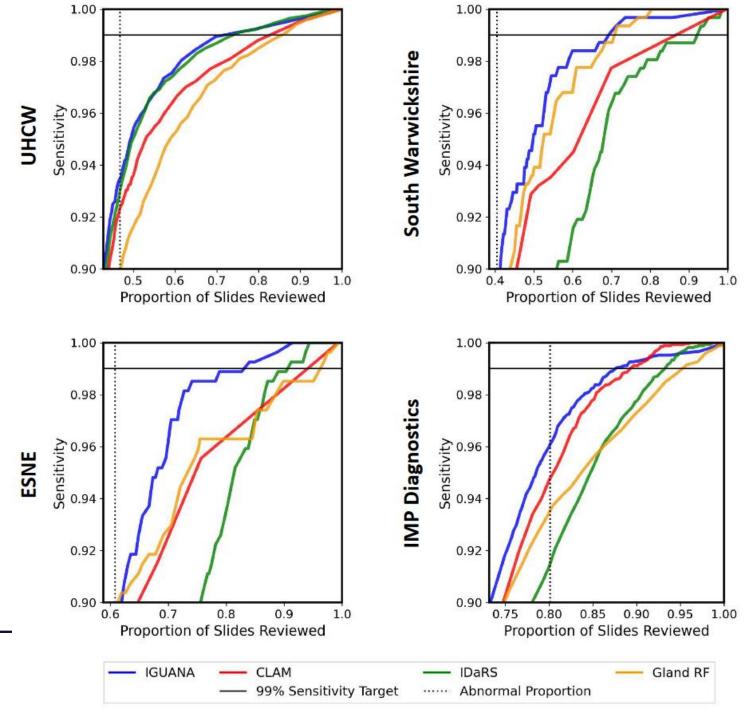


#### **Independent Validation**



### So what?

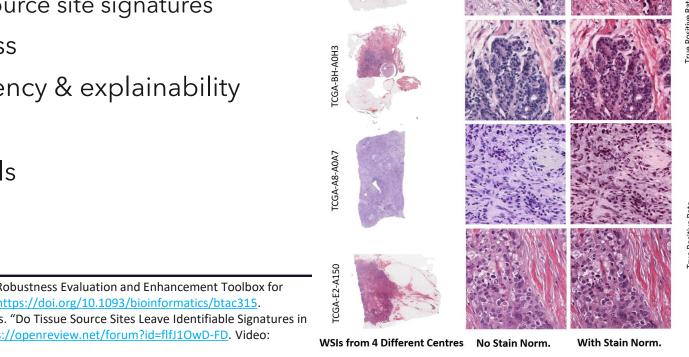
• How many slides need to be reviewed (to get a sensitivity of 99%)?

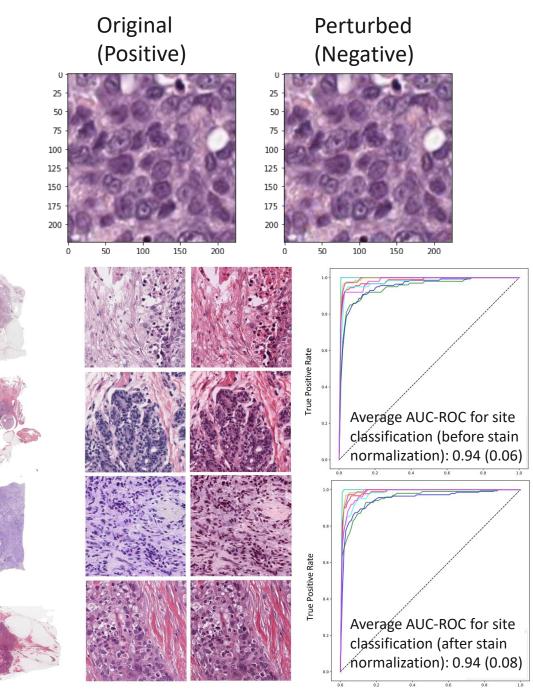


## Next Steps: CoBlx

**NIHR-COBIx:** Multi-site validation of automated AI tool for screening of large bowel endoscopic biopsy slides

- Larger validation
  - Data Quality
  - Ethnicity
  - Batch effects & Source site signatures
- Analysing robustness
- Improving transparency & explainability
- Health Economics
- Regulatory approvals
- Productization

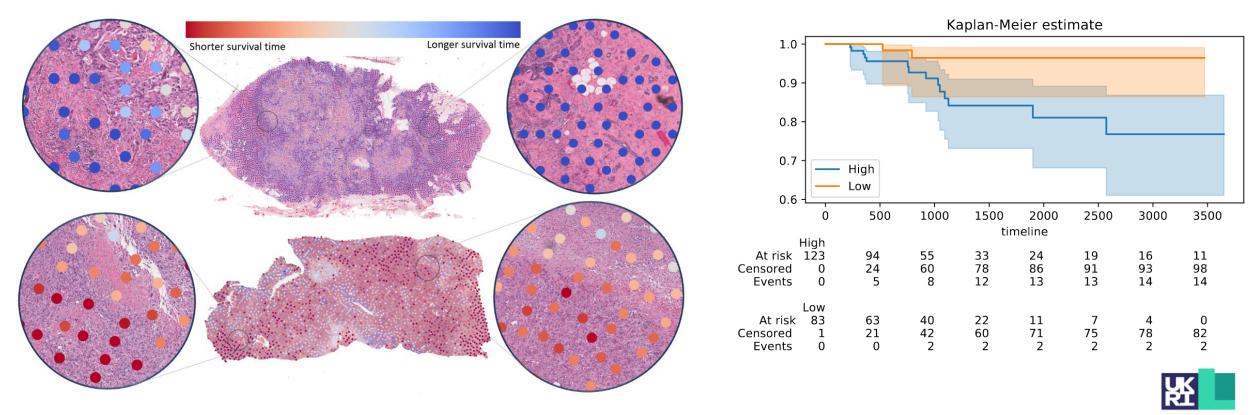




False Positive Rate

Foote, Alex, Amina Asif, Nasir Rajpoot, and Fayyaz Minhas. "REET: Robustness Evaluation and Enhancement Toolbox for Computational Pathology." Bioinformatics, May 9, 2022, btac315. <u>https://doi.org/10.1093/bioinformatics/btac315</u>. Dawood, Muhammad, Piotr Keller, and Fayyaz ul Amir Afsar Minhas. "Do Tissue Source Sites Leave Identifiable Signatures in Whole Slide Images beyond Staining?," ICLR-W TML4H, 2023. <u>https://openreview.net/forum?id=flfJ1OwD-FD</u>. Video: <u>https://www.youtube.com/watch?v=lg9g7xmZYPA</u>

## WSI graphs to predict patient survival

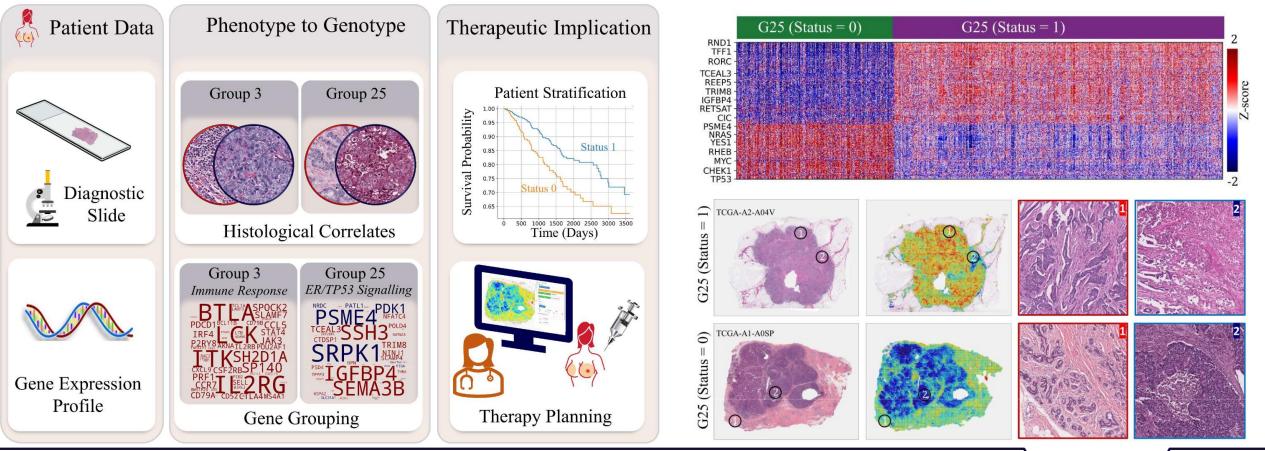


Engineering and Physical Sciences Research Council

Mackenzie, Callum Christopher, Muhammad Dawood, Simon Graham, Mark Eastwood, and Fayyaz ul Amir Afsar Minhas. "Neural Graph Modelling of Whole Slide Images for Survival Ranking." In *Proceedings of the First Learning on Graphs Conference*, 48:1-48:10. PMLR, 2022. https://proceedings.mlr.press/v198/mackenzie22a.html.

### Predicting gene expression from WSIs

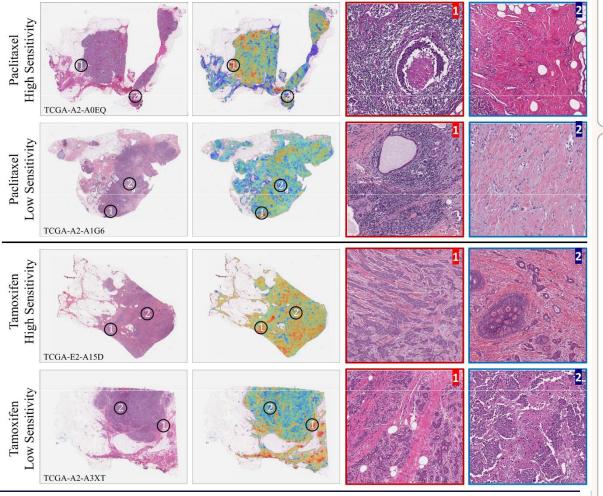




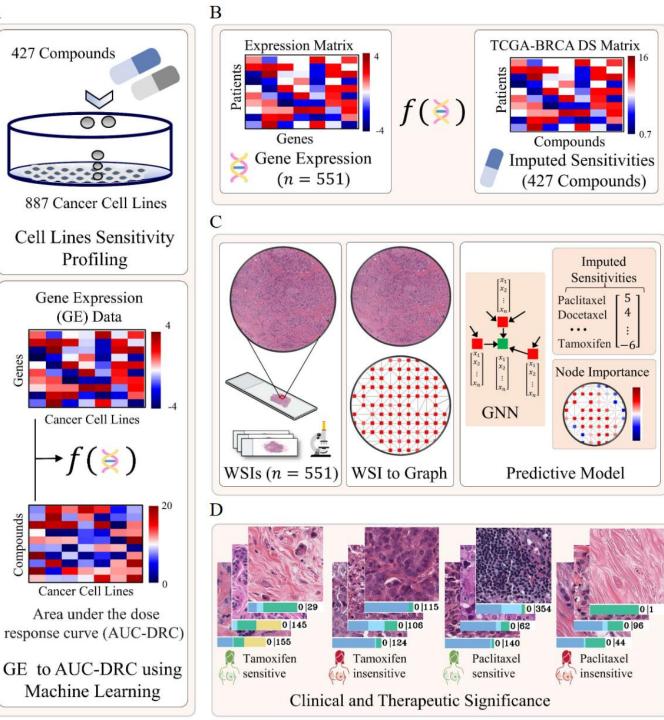
Dawood, Muhammad, Mark Eastwood, Mostafa Jahanifar, Lawrence Young, Asa Ben-Hur, Kim Branson, Louise Jones, Nasir Rajpoot, and Fayyaz ul Amir Afsar Minhas. "Cross-Linking Breast Tumor Transcriptomic States and Tissue Histology." *Cell Reports Medicine* 4, no. 12 (December 19, 2023). https://doi.org/10.1016/j.xcrm.2023.101313.

## WSI graphs to predict drug sensitivities

A



Dawood, Muhammad, Quoc Dang Vu, Lawrence S. Young, Kim Branson, Louise Jones, Nasir Rajpoot, and Fayyaz ul Amir Afsar Minhas. "Cancer Drug Sensitivity Prediction from Routine Histology Images." *Npj Precision Oncology* 8, no. 1 (January 6, 2024): 1–13. <u>https://doi.org/10.1038/s41698-023-00491-9</u>.



## Using Large Language Models

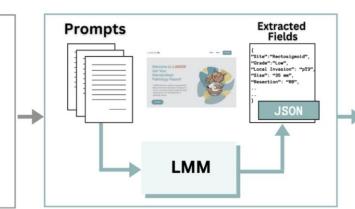
• For Standardization of Reports: LABIEB

#### **Pathology Report**



This is a moderately to poorly differentiated adenocarcinoma of the rectosigmoidal junction of grade G2 to 3, with tumor infiltration of the parietal layers into the subserous fatty connective tissue, with a regional lymph node metastasis (1/14) and tumor-free overview slices from all other resection material portions described.

Tumour stage of pT3, pN1 (1/14), MX, L1, R0.

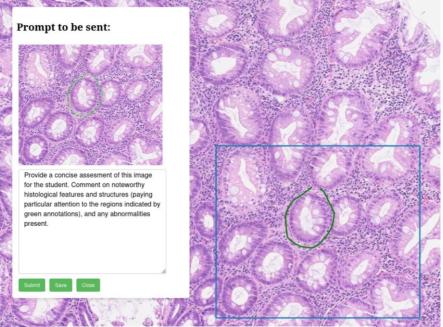


#### https://labieb.dcs.warwick.ac.uk/

#### Standardised Pathology Report

resection	or colorectal carcinoma specimens
Site of Tumour:	Number of lymph nodes:
Rectosigmoid	14
Tumour type:	Number of involved lymph nodes:
Adenocarcinoma	iymph nodes.
	1
Differentiation by worst	Lymph node status:
area:	pN1a
Low	
Local invasion:	Pathologically confirmed
pT3	distant metastatic disease:
	pMX
TNM (8 <sup>th</sup> edition)	Resection status:
pT3, pN1a, pMX	R0

#### • Image Interpretation: TIAViz



#### GPT-vision response:

This is an H&E (hematoxylin and eosin) stained histological image that appears to show a section of the gastrointestinal tract, most likely the colon or small intestine due to the presence of goblet cells, which are mucussecreting cells. The epithelial layer is columnar with regular nuclei, which is typical for this part of the digestive tract. The green annotation circles a gland in the epithelium. What's noteworth with sergion:

 In the annotated area, there is an intestinal gland (crypt of Lieberkfihm), which looks to be lined with regularly spaced columnare pithelial cells with basal, round to oval nuclei. The presence of goblet cells interspersed between the absorptive cells is typical for normal intestinal mucosa.

2. Within the crypt, there is no evidence of dysplasia, which is characterized by variations in cell size (aniso-cytosis), shape (pleomorphism), nuclear enlargement, hyperchromasia, and increased mitotic activity. The cells retain uniformity and there's no apparent crowding or stratification of nuclei, which is reassuring.

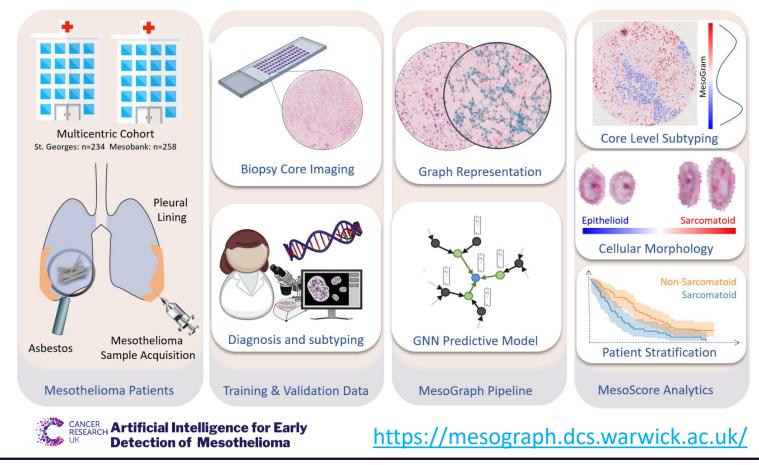
3. The surrounding stroma contains lamina propria, which shows a moderate number of inflammatory cells which can be normal, but an increased number could indicate mild inflammation. However, there doesn't appear to be significant infiltrate to suggest active inflammation or an inflammatory bowel disease.

4. The crypt architecture is maintained and there doesn't appear to be any evidence of cryptitis or crypt abscesses, which would suggest more specific inflammatory conditions such as ulcerative colitis.

Abnormalities: No overt abnormalities can be determined from this single region of the image, as the annotated crypt seems to have retained its normal histological architecture. Pathologists must, however, assess multiple fields and levels to establish a definite diagnosis and determine the absence of pathology accurately. For educational review always correlate microscopic findings with mecroscopic observations and clinical data

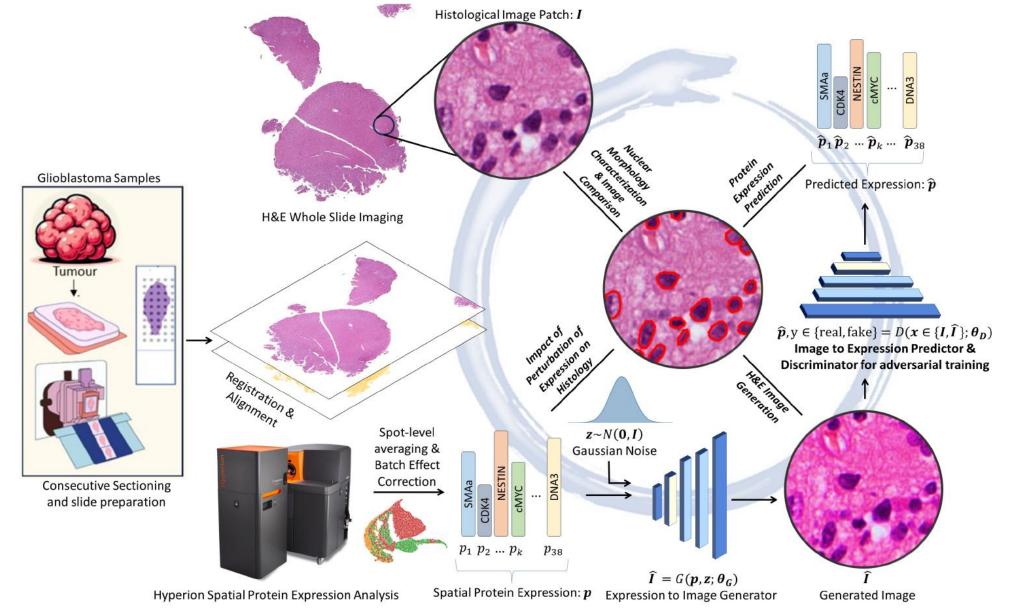
Alzaid, Ethar, Gabriele Pergola, Harriet Evans, David Snead, and Fayyaz Minhas. "Large Multimodal Model Based Standardisation of Pathology Reports with Confidence and Their Prognostic Significance." arXiv, May 3, 2024. <u>https://doi.org/10.48550/arXiv.2405.02040</u>. Eastwood, Mark, John Pocock, Mostafa Jahanifar, Adam Shephard, Skiros Habib, Ethar Alzaid, Abdullah Alsalemi, et al. "TIAViz: A Browser-Based Visualization Tool for Computational Pathology Models." arXiv, February 15, 2024. <u>http://arxiv.org/abs/2402.09990</u>.

## MesoGraph: subtyping mesothelioma



Epithelioid **Biphasic** Sarcomatoid

**Eastwood, Mark**, Heba Sailem, Silviu Tudor Marc, Xiaohong Gao, Judith Offman, Emmanouil Karteris, Angeles Montero Fernandez, et al. "MesoGraph: Automatic Profiling of Mesothelioma Subtypes from Histological Images." *Cell Reports Medicine* 0, no. 0 (October 9, 2023). <u>https://doi.org/10.1016/j.xcrm.2023.101226</u>. Epithelio

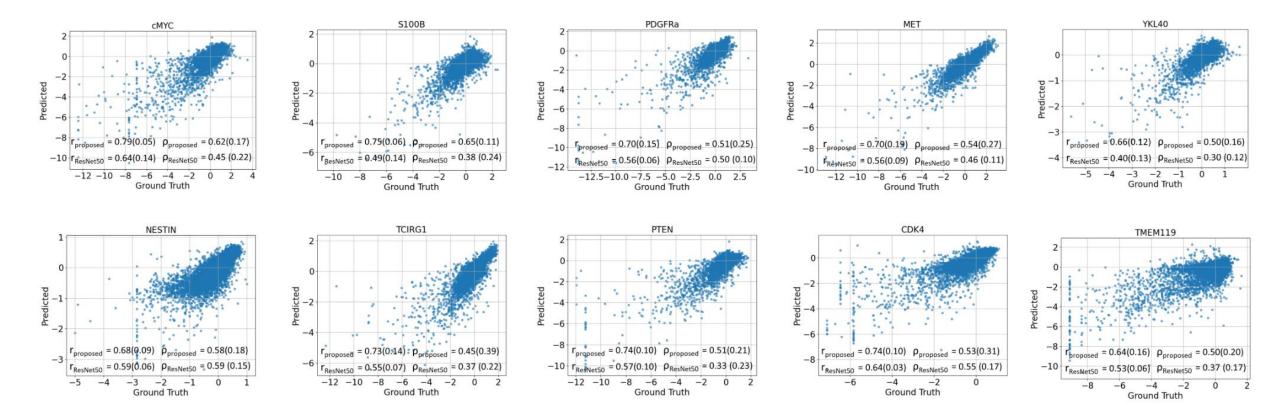


Can we predict spatial expression from imaging?

Can we generate images from expression profiles?

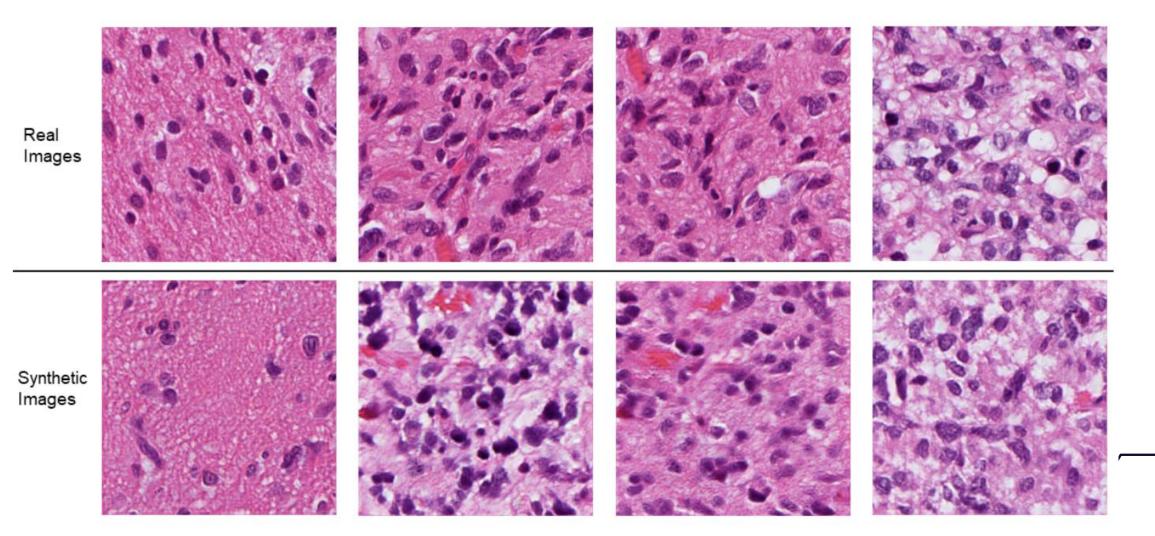
Deshpande, Srijay, Sokratia Georgaka, Michael Haley, Robert Sellers, James Minshull, Jayakrupakar Nallala, Martin Fergie, et al., Fayyaz Minhas "Ouroboros : Cross-Linking Protein Expression Perturbations and Cancer Histology Imaging with Generative-Predictive Modeling." Proceedings of the 23rd European Conference on Computational Biology, 2024. <u>https://doi.org/10.1093/bioinformatics/btae399</u>. [In press]

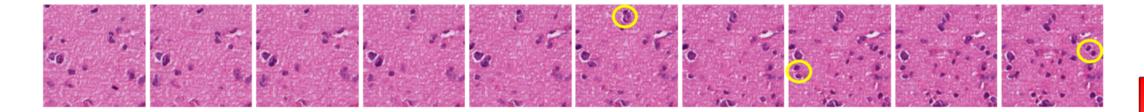
# SpecTx: Predicting protein expression from Imaging

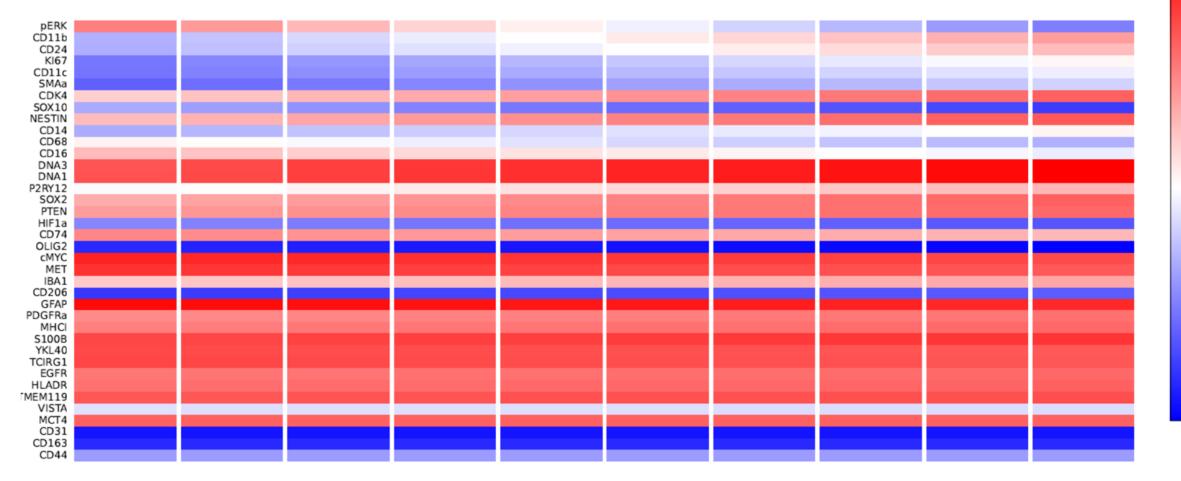


56

## SpecTx: Generating images from protein expression







Deshpande, Srijay, Sokratia Georgaka, Michael Haley, Robert Sellers, James Minshull, Jayakrupakar Nallala, Martin Fergie, et al., Fayyaz Minhas "Ouroboros : Cross-Linking Protein Expression Perturbations and Cancer Histology Imaging with Generative-Predictive Modeling." Proceedings of the 23rd European Conference on Computational Biology, 2024. <u>https://doi.org/10.1093/bioinformatics/btae399</u>. [In press] - 0

 $^{-1}$ 

-2

- - 3

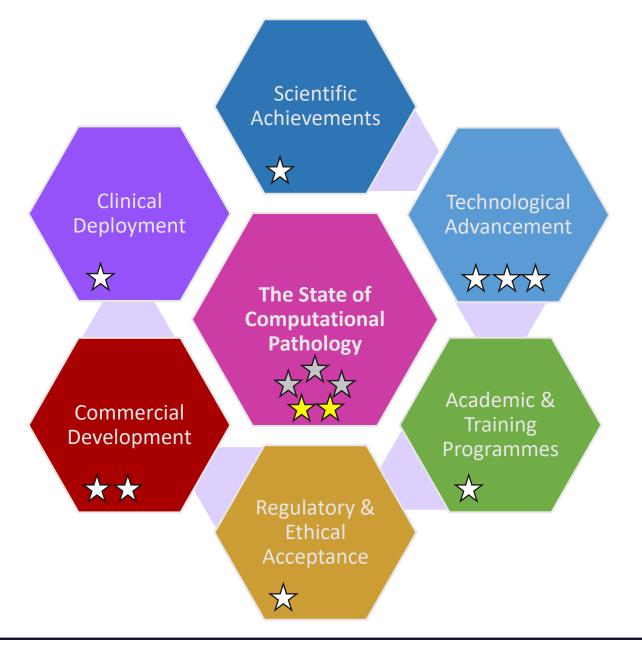
## The path for Cpath ahead

- 5 Pathologist Nobel Prize Winners [1]
  - Johannes Fibiger received the Nobel Prize in Physiology or Medicine in 1926 for the experiments in which he produced gastric carcinoma in rats by feeding them *Spiroptera*-infected cockroaches.
  - **George Whipple** was awarded the 1934 Nobel Prize in Physiology or Medicine for his discovery that a diet rich in liver cured pernicious anemia.
  - **Renato Dulbecco** was awarded the 1975 Nobel Prize for discovering by molecular techniques that the genetic material of viruses was incorporated into the genetic material of the transformed cells and represented the first phase in carcinogenesis.
  - **Baruj Benacerraf** received the 1980 Nobel Prize in Physiology or Medicine for his discovery of immune response genes.
  - John Warren received the Nobel Prize in Physiology or Medicine in 2005 in recognition of this discovery of the role of *H. Pylori* in ulcers.



Biology easily has 500 years of exciting problems to work on. It's at that level. (Knuth)







Towards Reinventing Pathology with AI