# COMPUTATIONAL PATHOLOGY: THE THIRD PATHOLOGY'S REVOLUTION CAN BE APPLIED TO CRC SCREENING?









(d) HG

Prof. Paola Cassoni

Pathology Unit, Dept. Medical Sciences

University of Turin, Italy







Adapted from Shafi, S., Parwani, A.V. *Diagn Pathol* (2023)



### ARTIFICIAL INTELLIGENCE

ENGINEERING OF MACHINES THAT MIMIC COGNITIVE FUNCTIONS



ABILITY TO PERFORM TASKS WITHOUT EXPLICIT INSTRUCTIONS AND RELYING ON PATTERNS



DEEP LEARNING

MACHINE LEARNING BASED ON ARTIFICIAL NEURAL NETWORKS

- Explicit programmig
- Predetermined features selection
- Multiple interactions pathologists/informaticians needed
- Time consuming

- Automated learning
- Freely available source codes of effective neural network architectures
- Superior results in most cases

## **COMPUTATIONAL PATHOLOGY "WORKFLOW"**

#### **Digital pathology**

The digitization of the traditional diagnostic process of analyzing cells and tissue with a microscope via whole-slide scanners and computer screens.

#### Whole-slide images

Digital images obtained by digitizing complete histopathological glass slides using a high-resolution scanner.

#### **Image segmentation**

The operation of decomposing the semantic content of an image into multiple segments, where each segment contains pixels belonging to the same semantic category (for example, the tumor region).

#### Model regularization

In machine learning, indicates the process of constraining a model's parameters to small values, discouraging complex models, therefore reducing the risk of overfitting the training data.



### Chan L. et al., *ICCV* (2019)

Weakly-annotated image patch



Loose Connective (C.L)

Seamented Patch

(Morphological)

(Functional) G.0

Seamented Patch

#### Patch-level annotations

- Epithelium (E.M.U) Exocrine Gland (G.O)
- Transport Vessel (T) Leukocytes (H.K) Lymphocytes (H.Y)
- Stratified Cuboidal
- Epithelium (E.T.U)
- Simple Cuboidal Smooth Muscle (M.M)
- White Adipose (A.W)
- Erythrocytes (H.E)
- Simple Squamous
- Epithelium (E.M.S)

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### **Deep learning overall workflow:**



# **Digital pathology for colorectal carcinoma**

- Distinction between tumor tissue and stroma (Kather JN et al. *Sci Rep* 2016)
- Outcome prediction (Bychkov D et al. Sci Rep 2018; Kather JN et al. PLoS Med 2019; Skrede O et al. Lancet 2020)
- Molecular profile prediction (Yamashita R et al., Lancet Oncol 2020; Sirinukunwattana K et al. Gut 2021; Bilal M et al. Lancet Digit Health 2021)



• Adenoma classification...

# **Adenoma classification**



### Original Investigation | Health Informatics Evaluation of a Deep Neural Network for Automated Classification of Colorectal Polyps on Histopathologic Slides

Jason W. Wei, BA; Arief A. Suriawinata, MD; Louis J. Vaickus, MD, PhD; Bing Ren, MD, PhD; Xiaoying Liu, MD; Mikhail Lisovsky, MD, PhD; Naofumi Tomita, MS; Behnaz Abdollahi, PhD; Adam S. Kim, MD; Dale C. Snover, MD; John A. Baron, MD; Elizabeth L. Barry, PhD; Saeed Hassanpour, PhD

JAMA Network Open. 2020;3(4):e203398. doi:10.1001/jamanetworkopen.2020.3398



# **Adenoma classification**

Table. Per-Class Comparison Between Local Pathologists and the Deep Neural Network Model in Classifying Colorectal Polyps on Internal and External Test Sets

	Internal test set (n = 157)							External test set (n = 238)					
	Local pathologists			Deep neural network			Local pathologists			Deep neural network			
Polyp type	Accuracy, %	Sensitivity, %	Specificity, %	Accuracy, %	Sensitivity, %	Specificity, %	Accuracy, %	Sensitivity, %	Specificity, %	Accuracy, %	Sensitivity, %	Specificity, %	
ТА	89.8	76.1	95.5	93.0	89.1	94.6	79.8	53.7	97.2	84.5	73.7	91.6	
TVA	94.3	88.2	95.8	95.5	97.1	95.1	81.5	100	77.7	89.5	97.6	87.8	
HP	89.8	76.9	94.1	92.4	82.1	95.8	91.6	80.8	96.8	85.3	60.3	97.5	
SSA	91.7	81.6	95.0	93.0	78.9	97.5	93.3	79.2	94.8	88.7	79.2	89.7	
Mean	91.4	80.7	95.1	93.5	86.8	95.7	86.6	78.4	91.6	87.0	77.7	91.6	

## Limitations:

- Lack of dysplasia grading
- Lack of normal tissue
- Lower performance during external testing



JAMA Network Open. 2020;3(4):e203398. doi:10.1001/jamanetworkopen.2020.3398

## High Performance Computing boosting Biomedical Applications



#### PARTNERS

22 partners from 9 European Countries

#### **Research Organisations**



### Aim

Provide **High Performance Computing (HPC)** power at the service of biomedical applications; and apply **Deep Learning** (DL) and **Computer Vision** (CV) techniques on large and complex biomedical datasets to support new and more

efficient ways of diagnosis, monitoring and treatment of diseases







This project has received funding from the European Union's Horizon 2020 research innovation programme under grant agreement No. 825111





# UniTOPatho





### Use Cases

14 pilot test-beds in 3 areas:

### Neurological diseases

- Migraine and Seizures prediction
- Major Depression
- Dementia
- Study of structural changes in lumbar spine
   pathology
- Population model for Alzheimer's Disease
- Epileptic seizures detection
- Objective fatigue assessment for multiple sclerosis patients

### Tumor detection and early cancer prediction

- Chest cancer detection
- Prostate tumor diagnosis
- Skin cancer melanoma detection

## Digital pathology and automated image annotation

- Classification of whole-slide histological images of colorectal biopsy samples
- CT brain perfusion maps synthesis
- Deep Image annotation
- Image Analysis and prediction for Urology

# **Colon cancer diagnosis**

## DeepHealth

Colon cancer is one of the most frequent causes of death. Screening programs can enable prompt diagnosis and treatment of this aggressive disease, but they also lead to higher caseloads and costs for the already strained European healthcare services. DeepHealth can help streamline pathological diagnosis of colon biopsies.



# **Dataset (WSI images)**

	HP	NORM	TA.HG	TA.LG	TVA.HG	TVA.LG	Total
Slides	62	30	34	232	44	55	457
$R_t$	158	112	145	777	264	245	1701
$A_t \left[ \mathrm{cm}^2 \right]$	9.91	18.38	7.94	71.74	60.45	41.86	210.29

- H&E slide acquired on the Hamamatsu Nanozoomer S210 scanner (200X)
- Manual annotation according to 6 classes:
  - NORM: normal tissue
  - HP: hyperplastic polyp
  - TA.LG: tubular adenoma, low-grade dysplasia
  - TA.HG: tubular adenoma, high-grade dyplasia
  - TVA.LG: tubulo-villous adenoma, low-grade dysplasia
  - TVA.HG: tubulo-villous adenoma, high-grade dysplasia



Perlo D. et al. MICAD 2021

# **Dysplasia grading**

	(a)	$\varphi = 600$	μm, gray	y-scale		(b) $\varphi = 600  \mu m$ , RGB							
			Predic	$\operatorname{cted}$			Predicted						
		HP	NORM	HG	LG			HP	NORM	HG	LG		
h	HP	0.85	0	0.05	0.1	h d	HP	0.75	0.05	0	0.2		
rut	NORM	0.12	0.75	0	0.12	rut	NORM	0	0.62	0	0.38		
: t	HG	0.02	0	0.63	0.35		HG	0	0.02	0.61	0.37		
5	LG	0.03	0.09	0.18	0.7	5	LG	0.03	0.06	0.15	0.76		

 Table 4. WSI inferences: confusion matrices.



Poor results in distinguishing
 TA versus
 TVA/VA



Perlo D et al. MICAD 2021

**(c)** LG

**(d)** HG

# **Multi-resolution analysis**

				Patch scale $\sigma$ [µm]					
Туре	100	800	1500	4000	7000	8000			
BA (6-class)	0.40	0.45	0.46	0.41	0.37	0.38			
NORM	0.70	0.66	0.72	0.76	0.78	0.71			
HP	0.81	0.92	0.85	0.70	0.60	0.69			
TA (HG+LG)	0.65	0.66	0.65	0.71	0.76	0.70			
TVA (HG+LG)	0.64	0.67	0.68	0.74	0.84	0.76			

**Table 2**: Preliminary experiments: overall BA for all of the sixclasses (first row) and BA for each polyp type, plus normal tissue.

 Adenoma type and dysplasia grade are best classified at different scales



### Limitations:

- Some entities missing (serrated adenomas, invasive adenocarcinomas,...)
- Larger dataset is warranted
- Lack of external validation

Barbano CA et al. IEEE ICIP 2021

## OPEN A promising deep learning-assistive algorithm for histopathological screening of colorectal cancer

Cowan Ho<sup>1,6</sup>, Zitong Zhao<sup>2,6</sup>, Xiu Fen Chen<sup>2,6</sup>, Jan Sauer<sup>3</sup>, Sahil Ajit Saraf<sup>3</sup>, Rajasa Jialdasani<sup>3</sup>, Kaveh Tadhipour<sup>3</sup>. Aneesh Sathe<sup>3</sup>. Li-Yan Khor<sup>2,4</sup>. Kiat-Hon Lim<sup>2,4</sup> & Wei-Qiand Leow<sup>2,4,5</sup>

SCIENTIFIC REPORTS, 2023 nature portfolio



**Figure 1.** A sample analysis on a whole slide image (WSI). Left: original image. Right: the segmentation model highlighted regions of the WSI as (1) likely benign or normal (green), (2) likely dysplastic (orange), and (3) likely malignant (red). The AI model also segmented blood vessels (pink) and inflammation (yellow), and these segmentations were taken into account for slide labeling.





**Figure 4** AUC curve from applying the AI model on the validation set of 150 WSIs. The system achieves an AUC of 91.7%.

### CONCLUSIONS

Upon evaluation of the output data from the validation set, its high AUC of 91.7 demonstrates a good concordance between the AI model and the expert pathologist. With a sensitivity of 97.4, it validates our AI model to function as a screening tool that minimizes false negatives extensively. This favour of sensitivity over specificity shows its usability in assistive workflow and has added practicality into the application of a clinical workflow. Since our AI model is designed solely as an assistive tool, the final diagnosis during reporting remains with the pathologist. This high sensitivity ensures with greater certainty that all lesions suspicious for



# To sum up:

CPATH is the third revolution in pathology and when applied CRC SCREENING may helps to **identify** and **distinguish benign and cancerous glands, as well as low and high dysplasia**→ implementing these technoloy in the diagnostic workflow would result in a <u>reduction</u> of turn around time, workload, misdiagnosis and lab costs however some challenges are still open.

### **FUTURE PERSPECTIVES ON CPATH**

- Implement novel tool to minimize errors
- Improve the pathology units worldwide (e.g. slide scanner, IT unit ecc...)
- Train pathologists to use WSI and CPATH









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