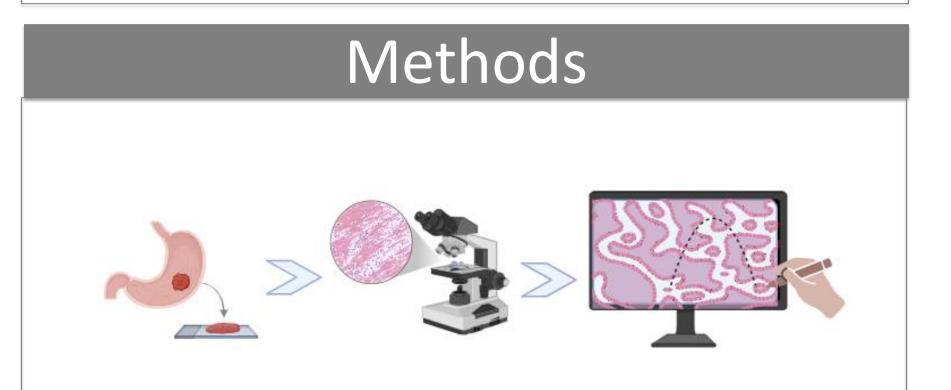


Florina Almarii^{1,2,3}, Corina-Elena Minciuna^{2,3}, Alexandra Ondu^{2,3}, Stefan Tudor², Mihai Tanase⁴, Dragos Stan⁵, Aurel Tonea², Alex Micu², Ovidiu Bitere^{2,3}, Teodora Ecaterina Manuc², Simona O. Dima^{2,3}, Vlad Herlea^{2,3}, Catalin Vasilescu^{2,3}

Introduction

carcinoma continues to Gastric pose а global health challenge, substantial despite advancements in treatment. Consequently, accurate pathological timely evaluation remains and paramount. Artificial Intelligence (AI), particularly in image analysis, holds significant potential to histopathological workflows revolutionize by augmenting accuracy and efficiency [1].

The aim of the study was to develop an application capable of delivering an assisted the efficiency the diagnosis, enhancing Of pathologist's workflow.



manually experienced pathologists Four the slides according to annotated Lauren's classification of gastric cancer, into five classes using QuPath v0.4.4.

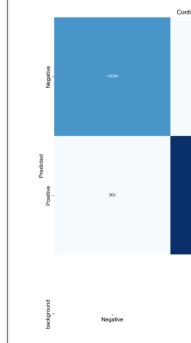
Image analysis was conducted following a threestep process:

I. Classify all patches in 2 classes, positive and negative, selecting only those that have confidence over 90%.

II. Classify all positive patches in 2 classes, intestinal and diffuse, selecting only those that have confidence over 90%.

III. Reconstruct the full image taking into consideration negative, diffuse and intestinal patches from the above

I. Metrics model 1



confusion matrix illustrates the Figure The performance of the classification model in distinguishing between the two categories of tiles: positive and negative.

II. Metrics model 2

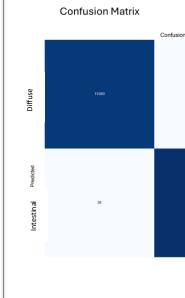


Figure 2: The confusion matrix illustrates the performance of the classification model in distinguishing between the two subtypes of gastric cancer, showing excellent performance with minimal misclassified tiles.

Figure 3: Illustration and predictions for model II

Revolutionizing Gastric Cancer Diagnosis: AI-Powered Image Analysis for Pathologists

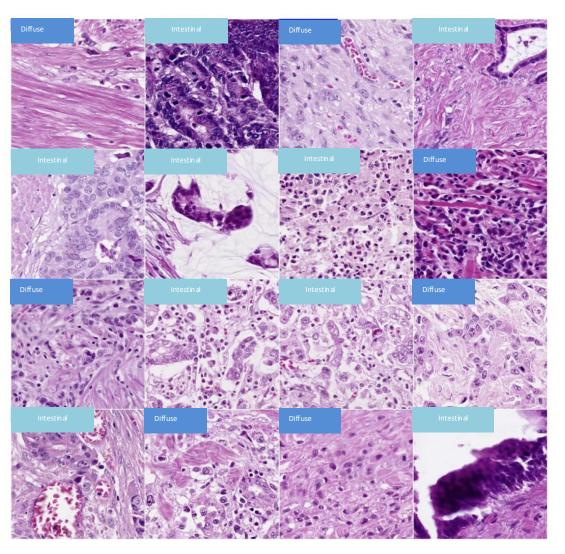
1. The University of Texas MD. Anderson Cancer Center, Houston, USA 2. Fundeni Clinical Institute, Bucharest, Romania 3. Carol Davila University of Medicine and Pharmacy, Bucharest, Romania 4. University of Bucharest, Romania 5. http://linkedin.com/in/dragos-stan-5a2909225

Results

nfusion Matrix								
253		- 20000	Classification prec			1-score	e support	
		- 15000	Negative Positive	0.98 0.99	0.98 0.99	0.98 0.99	15350 25166	
		- 10000	accuracy macro avg weighted avg	0.99 0.99	0.99 0.99	0.99 0.99 0.99	40516 40516 40516	
Positive	background	- 5000						

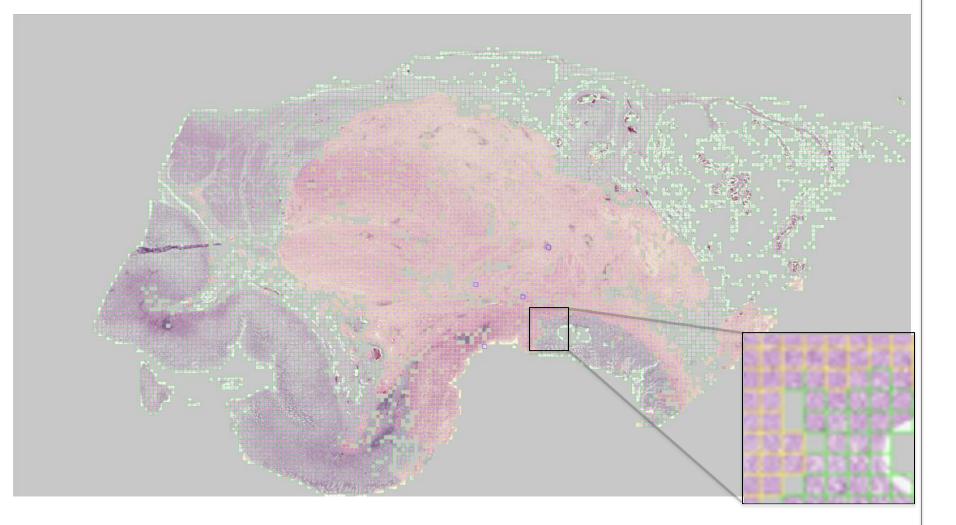
on Matrix 28	Classification Report: precision recall f1-score support								
	- 8000	Diffuse Intestinal	1.00 1.00	1.00 1.00	1.00 1.00	11618 12063			
	- 6000	mestinai	1.00	1.00	1.00	12000			
12035		accuracy			1.00	23681			
	- 4000	macro avg	1.00	1.00	1.00	23681			
		weighted avg	1.00	1.00	1.00	23681			
	- 2000								

of validation batches with identical labels

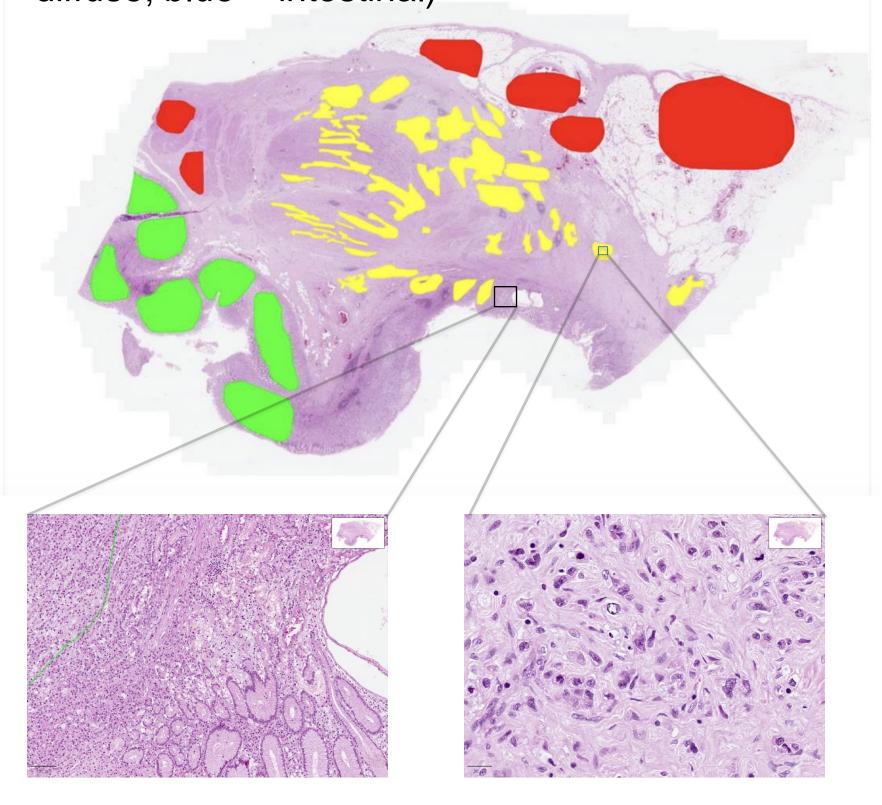


III. Image reconstruction and patch classification

Reconstructed (yellow = diffuse; green = negative)



= diffuse; blue = intestinal)



patches

Original annotation (red and green = negative; yellow

Figure 4: Image reconstruction including the unknown

Conclusions

This study explores the potential of artificial intelligence (AI) in revolutionizing histopathological workflows by automating tumor classification. By employing a pipeline inferencing approach using a supervised model to classify and map tumor regions in gastric carcinoma, the research emphasizes the applicability of AI in achieving precision and efficiency in pathology.

Future directions: Expanding on this model, our future research will explore automation of biomarker interpretation, predictive scoring for disease progression, and detection of visual indicators for treatment resistance.

References & Acknowledgements

This work was supported by a grant of the Ministry of Research, Innovation and Digitization, CNCS UEFISCDI, project number PN-III-P4-PCE-2021-1068, within PNCDI III. We extend our gratitude to Professor George Calin from the Department of Translational Molecular Pathology, MD Anderson, Houston, Texas, for his invaluable support.

Florina Almarii gratefully acknowledges financial support for research by the Fulbright Visiting Scholar Program, which is sponsored by the U.S. Department of State and the Romanian-American Fulbright Commission.

[1] Veldhuizen GP, Röcken C, Behrens HM, Cifci D, Muti HS, Yoshikawa T, Arai T, Oshima T, Tan P, Ebert MP, Pearson AT, Calderaro J, Grabsch HI, Kather JN. Deep learning-based subtyping of gastric cancer histology predicts clinical outcome: a multi-institutional retrospective study. Gastric Cancer. 2023 Sep;26(5):708-720. doi: 10.1007/s10120-023-01398-x. Epub 2023 Jun 3. PMID: 37269416; PMCID: PMC10361890.

