

People's Democratic Republic of Algeria
Ministry of Higher Education and Scientific Research
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Faculty of Exact Science
Computer Science Department



Revolutionizing Lung Cancer Diagnosis with LungPathAl: An Al-Powered Digital Pathology Platform

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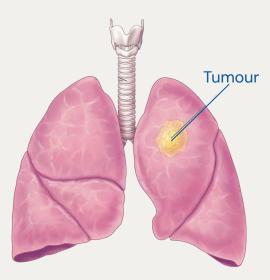
01

Introduction

Introduction

Lung cancer is a major health concern in Algeria. In 2022, it ranked among the most diagnosed cancers, with **5,040 new cases**.

Among men, it had the **highest cancer-related death rate**, with an incidence of **19.5** and mortality of **17.9 per 100,000** (GLOBOCAN, 2022).



Cancer Statistics in Algeria (2022)



Number of new cases

64 713

Number of deaths

35 778

Number of prevalent cases (5-year)

177 718

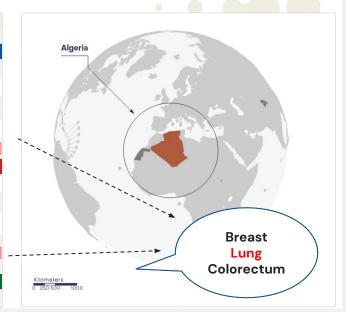
Statistics at a glance, 2022



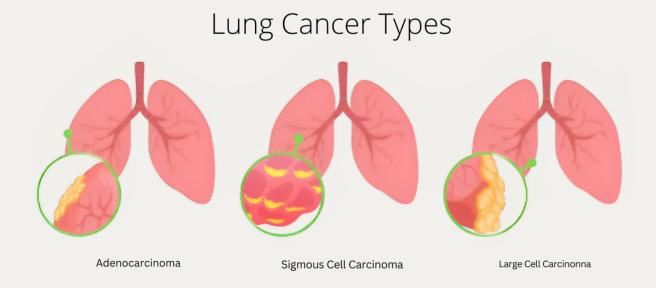
Population	22 912 687	22 437 454	45 350 141
Incidence*			
Number of new cancer cases	29 387	35 326	64 713
Age-standardized incidence rate	130.6	152.2	141.2
Risk of developing cancer before the age of 75 years (cum. risk %)	14.1	15.1	14.6
Top 3 leading cancers (ranked by cases)**	Colorectum Prostate	Breast Colorectum Thyroid	Breast Colorectum Lung
The Court of the C			

Lung Colorectum Bladder

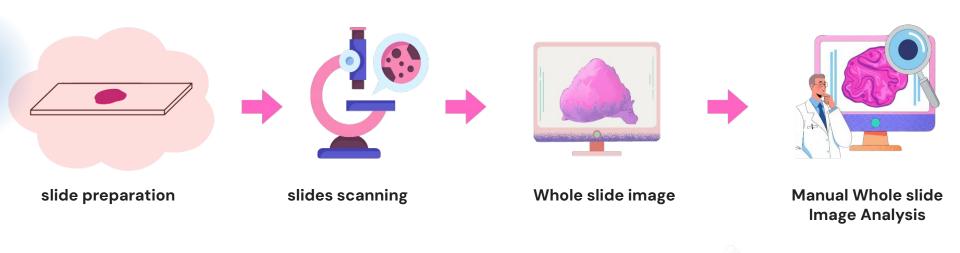
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Mortality*			
Number of cancer deaths	18 809	16 969	35 778
Age-standardized mortality rate	82.7	73.1	77.7
Risk of dying from cancer before the age of 75 years (cum. risk %)	8.7	7.8	8.2
Top 3 leading cancers (ranked by deaths)**	Lung Colorectum Bladder	Breast Colorectum Cervix uteri	Breast Lung Colorectum
Prevalence*			
5-year prevalent cases	72 408	105 310	177 718



Introduction | Lung Cancer Types



Introduction | TRADITIONAL SLIDE DIAGNOSIS



Limitations of Traditional Lung Cancer Diagnosis



Time-intensive Process

Manual examination of tissue samples creates significant diagnostic delays.



Diagnostic Variability

Substantial inter-observer differences among pathologists affect consistency.



Volume Challenges

Rising cancer incidence is overwhelming existing diagnostic capacity.



Limited Standardization

Inconsistent sample preparation and interpretation practices across institutions

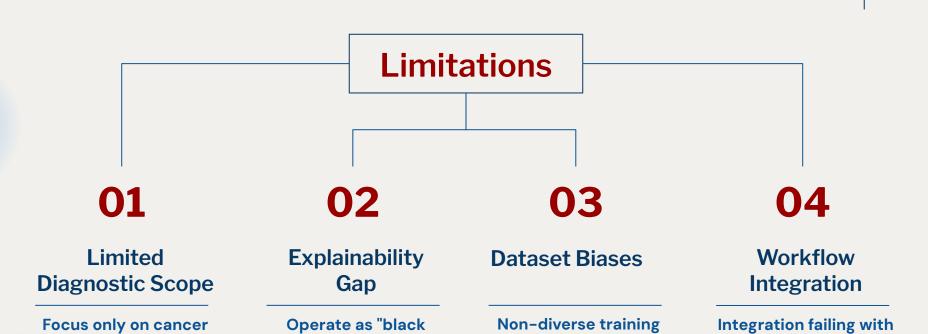
02

Problem Statement

Al Solutions in Literature: Limitations

subtypes and fail to

detect normal tissue



data leads to poor

performance across

varied demographic

groups

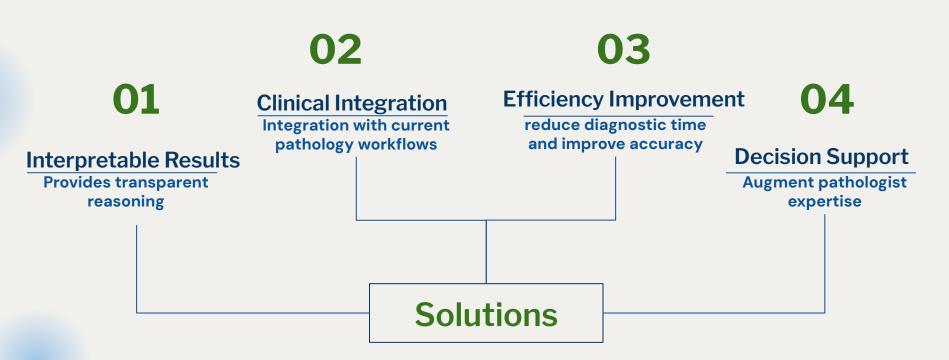
boxes," limiting clinical

trust and adoption

existing clinical

systems

Proposed Solution



03

Related Work

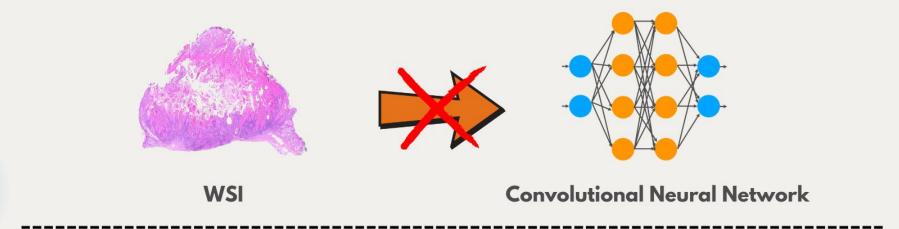
Overview of Leading MIL Methods

Approach	Year	Learning Paradigm	Training/Inference Speed	Clinical Interpretability Quality
CLAM	2021	Weakly Supervised + Clustering	Moderate/Fast	Good
TransMIL	2021	Correlated instance learning	Slow/Moderate	Normal
DSMIL	2021	Self-supervised + supervised	Fast/Fast	Normal
CAMIL	2025	Channel-aware feature learning	Moderate/Fast	Normal

Our work

Adopt CLAM architecture to classify lung cancer subtypes. Train and Evaluate the model using the public CPTAC dataset. Generate heatmaps highlighting regions of interest (ROIs) using attention scores. Integrate the model into a web application. Validate heatmaps quality by comparing model outputs with pathologist-provided annotations.

Why Not Traditional CNN?



04

Methodology

Operational steps

2- Preprocessing

- Tissue Segmentation
- Patching
- **Feature Extraction**

4- Explainability (XAI)

- **Attention Scores**
- Generate Heatmap over WSI

1- Data Acquisition

Whole Slide Images (WSIs) collected from public dataset (TCGA)

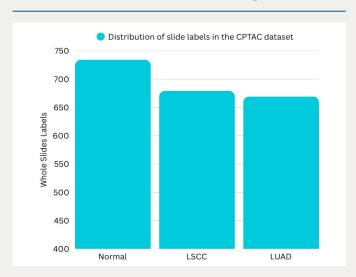
3- Deep Learning Model 5- Model Validation

Train a model that classifies each slide into LUAD, LSCC or Normal with high accuracy

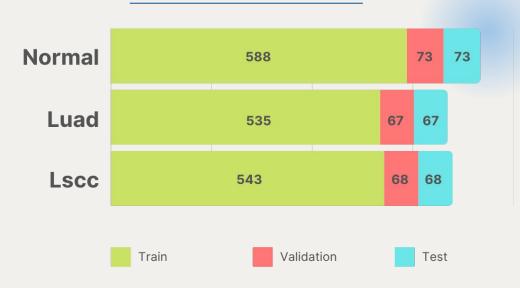
Validate the model with unseen data

1- Data Acquisition

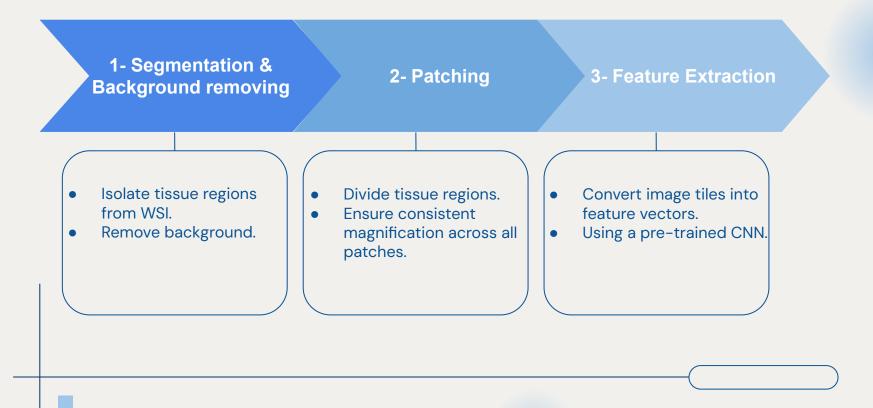
Data-set Description



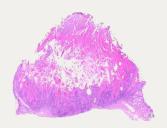
Data-set Split



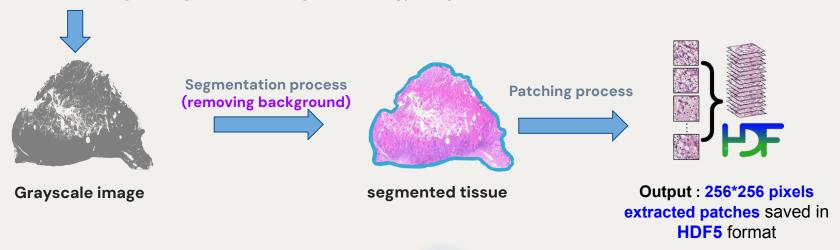
2- Preprocessing



Step 1: Removing background and Patching



Whole slide image: a high-resolution digital pathology image (in svs format).



Segmentation Params

Parameter	Value	Description		
Segmentation Level	-1	Resolution level used for segmentation -1 indicates the lowest available resolution level in the WSI		
Saturation Threshold	8	determine if a pixel belongs to tissue based on its color saturation		
Close	4	Size of the morphological closing operation to fill small holes and gaps in tissue regions		
Area Threshold	100	Minimum area threshold for keeping a tissue region. Regions smaller than this are removed		
Area holes	16	Minimum hole area to be considered for removal from tissue mask		
Max_n_holes	8	Maximum number of holes allowed per tissue region		

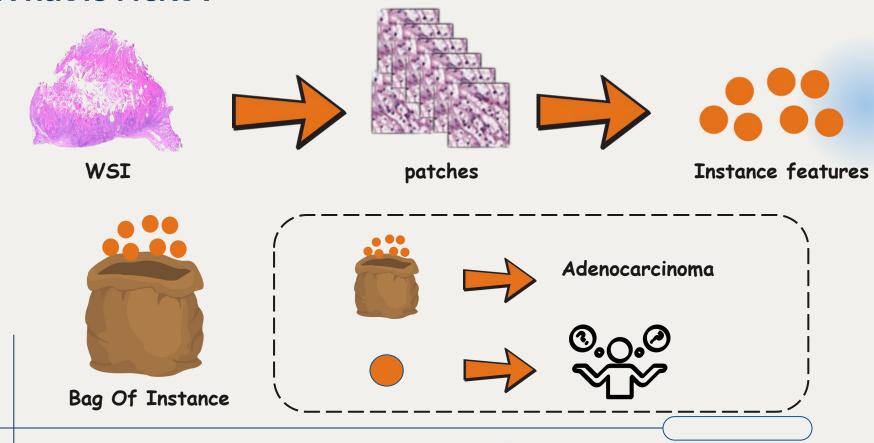
Patching Params

Parameter	Value	Description		
Patch Level	0	The resolution level of the WSI from which patches are extracted (0 is is for the best details)		
Patch Size	256	Size of each patch		
Step Size	256	This control how far to move when extracting the next patch it determines the overlap between adjacent patch		

Step 2: Feature Extractions



What is Next?



3- Deep Learning Model Single-Branch

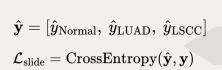


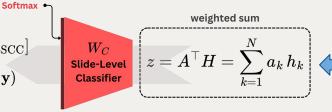


Components	Single Branch	Multi Branch	
Attention Branches	Single Attention Pathway (one attention score per patch)	Multiple Attention Pathways (class- specific attention score per patch)	
Slide Representation	A single aggregated slide-level representation	3 class-specific slide-level representations (Normal , Luad , Lscc)	
Slide-Level Classifier	A single classifier acts on the aggregated slide-level representation	3 parallel classifiers act on 3 class- specific representations	
Patch-Level Classifier (Clustering)	A single patch-level objective based on general attention.	N parallel patch-level objectives based on class-specific attention.	

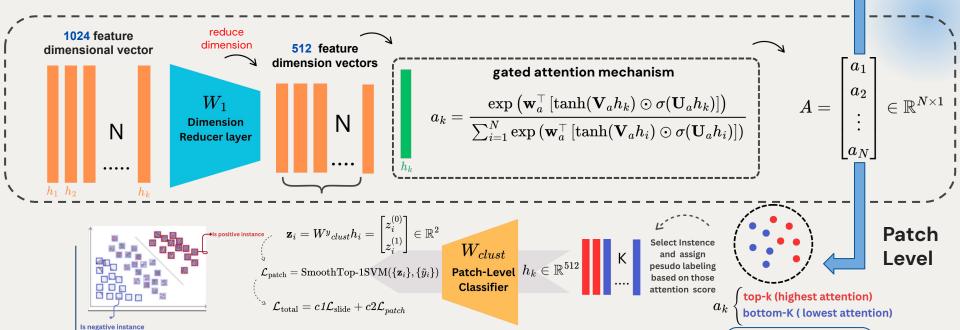
3- Deep Learning Model

Single Branch





Slide Level



3- Deep Learning Model

$$\hat{\mathbf{y}} = [\hat{y}_{ ext{Normal}},~\hat{y}_{ ext{LUAD}},~\hat{y}_{ ext{LSCC}}]$$

Softmax •

Multi Branch

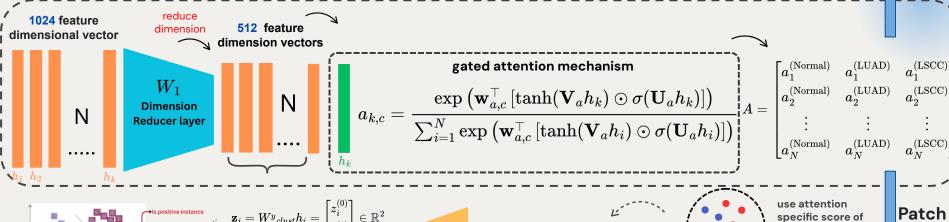
 $\mathcal{L}_{\mathrm{slide}} = \mathrm{CrossEntropy}(\hat{\mathbf{y}}, \mathbf{y})$

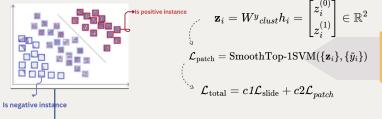
weighted sum

 W_C

$$Z = \mathbf{A}^ op \mathbf{H} = egin{bmatrix} \sum_{k=1}^N a_k^{ ext{(Normal)}} h_k \ \sum_{k=1}^N a_k^{ ext{(LUAD)}} h_k \ \sum_{k=1}^N a_k^{ ext{(LSCC)}} h_k \end{bmatrix} = egin{bmatrix} z^{ ext{(Normal)}} \ z^{ ext{(LUAD)}} \ z^{ ext{(LSCC)}} \end{bmatrix}$$

Slide Level







Select Instence and assign pesudo labeling based on those attention score



specific score of the true class Level

top-k (highest attention)

bottom-K (lowest attention)

Training HyperParameters

Parameter	Value		
Maximum epochs	200		
Learning rate	0.0002		
Optimizer	Adam Cross-entropy (ce) Smooth Top-1 SVM		
Bag loss	Cross-entropy (ce)		
Instance loss	Smooth Top-1 SVM (svm)		
Bag weight	0.7		
Dropout rate	0.25		
Early stopping	True		

04

Experimental Results

Results | Evaluation Metrics:

Single Branch

Accuracy = 89%



	auc	accuracy	precision	recall	f1_score	specificity
Normal	0.9962455606	0.971153846	0.9466666667	0.9726027397	0.9594594595	0.9703703704
Luad	0.9611516884	0.913461538	0.9152542373	0.8059701493	0.8571428571	0.9645390071
Lscc	0.9712184874	0.913461538	0.8378378378	0.9117647059	0.8732394366	0.9142857143

Multi Branch

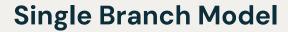
Accuracy = 88%



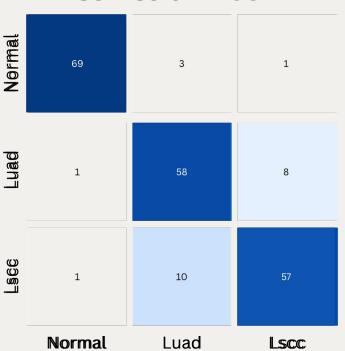
	auc	accuracy	precision	recall	f1_score	specificity
Normal	0.998782344	0.971153846	0.971830985	0.9452054795	0.9583333333	0.9851851852
Luad	0.96305705	0.894230769	0.816901408	0.8656716418	0.8405797101	0.9078014184
Lscc	0.970378151	0.903846153	0.86363636	0.8382352941	0.8507462687	0.9357142857

Results

Confusion Matrix



Multi Branch Model



Results | ROC Curve:

Class 0 (Normal): AUC = 1.00

Class 1 (LUAD): AUC = 0.96

Class 2 (LSCC): AUC = 0.97



Pathologist business card



Pathologist validation

Pathologist Validation Report

Al-Generated HeatMap Analysis for Lung Pathology (LungPathAl)

Validating Pathologist: Dr. Khelifi N Ep karasad

Institution/Affiliation: Ex assistante at EPH biskra / CHU at beni messous

Board Certifications: Anatomic Pathology

Date of Validation: 20/05/2025

-

Al System Under Review: LungPathAl Heat Map

Developers: Ouamane Takieddine, Cherifi Kacem, Guesbaya Islem

Supervisor: ABIR belaala

Cases Reviewed: 10 lung pathology specimens

- Lung Adenocarcinoma (LUAD)
- Lung Squamous Cell Carcinoma (LSCC)
- Normal Lung Tissue

Validation Method: Independent pathologist review comparing Al-generated heat map outputs against standard histopathological assessment.

Based on my review as a board-certified pathologist with expertise in lung pathology, I hereby validate that:

The LungPathAI heat map generation system produces clinically relevant outputs that accurately identify pathologically significant areas in lung tissue specimens.

The heat maps demonstrate sufficient accuracy and clinical utility to serve as an effective diagnostic aid when used under appropriate pathologist supervision.

Recommendation: APPROVED for clinical use as a supplementary diagnostic tool in lung pathology practice.

Limitations: This validation is based on the specific ease cohort reviewed. The AI system should be used as an adjunct to, not replacement for, standard pathological assessment. Continued monitoring and periadic revalidation are recommended.

LABORATO SE PARATO SE PROCESSIONALE SE P

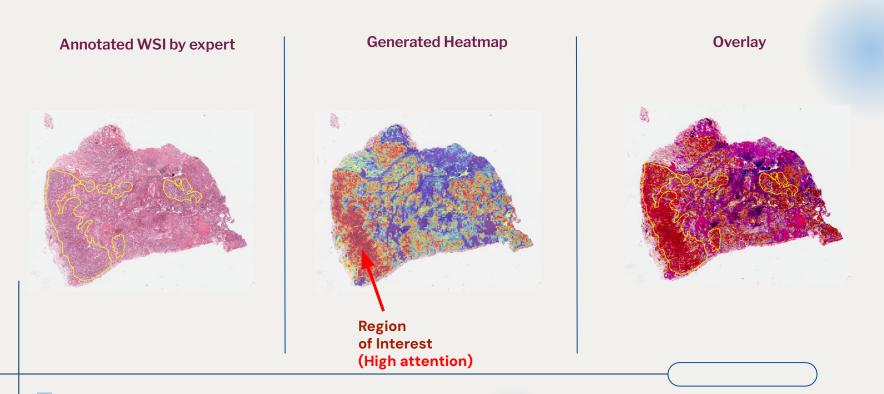
Dr. KHELIFI N Ep KARASAD

Date: 15/06/2025

MEDECIN SPECIALISTE EN ANATOMIE/PATHOLOGIQUE

1/1

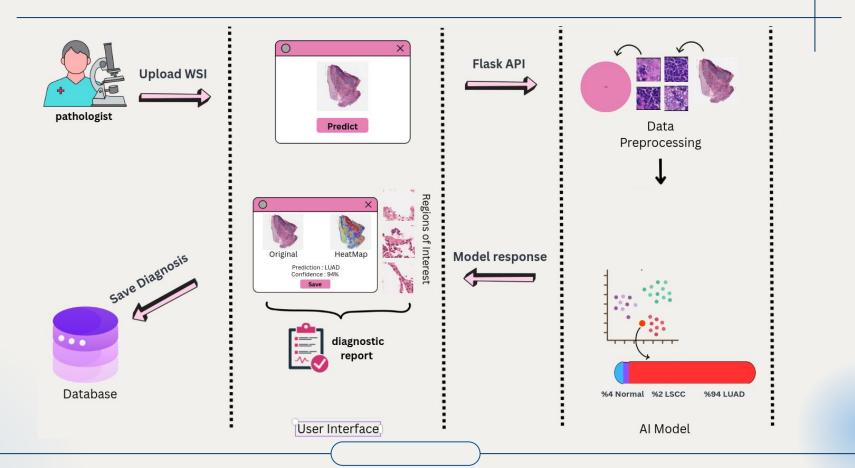
5- Explainability (XAI)



06

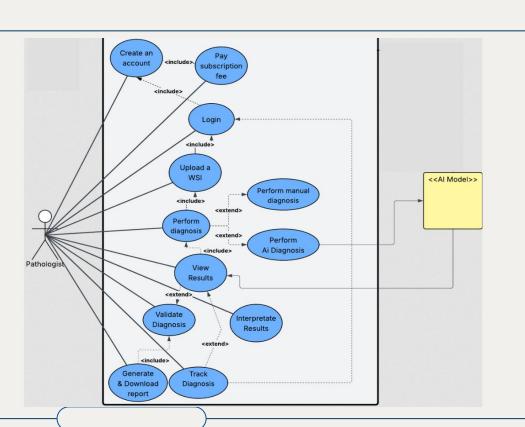
Platform Development

Platform Development | Design



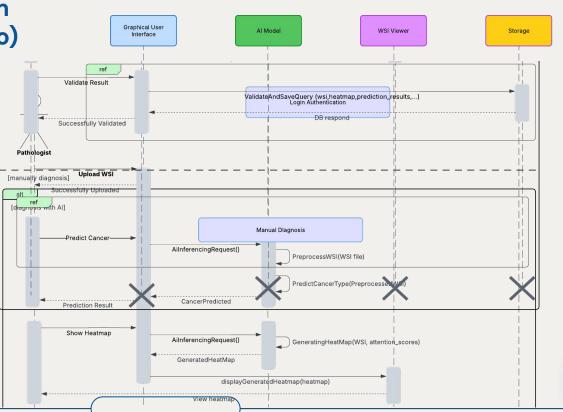
Platform Development | Design

1 - Use Case Diagram



Platform Development | Design

2 - Sequence Diagram (Al Diagnosis Scenario)



Platform Development | Tools







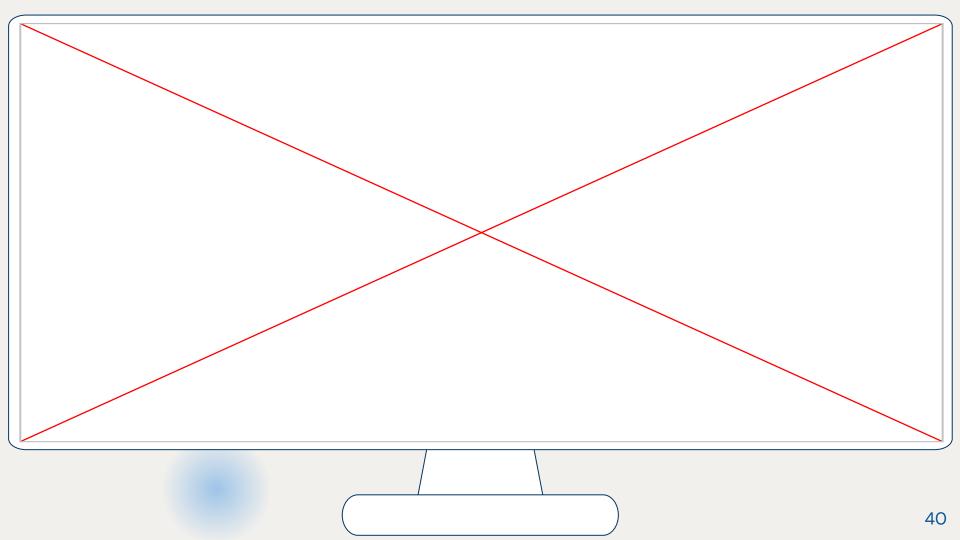












07

Conclusion

Conclusion



Leveraging deep learning techniques to enhance
Histopathology diagnosis accuracy by
training a model on WSIs for lung cancer subtype
classification



Provides clinical explainability through attention-based heatmaps



Integrate the validated model into a web application

Future Perspectives



Expanding the dataset from diverse institutions and regions to improve model generalizability



Include multi modal approach by Combining histopathological image features with clinical data



Implementing comprehensive multi-user collaboration features

Thank you

فكرة المشروع



تجاريا ، يقدم المشروع على شكل خدمة طبية تسوق للمخابر و المستشفيات بشكل أساسي .



سرطان الرئة مرض منتشر و خطير عالميا ومحليا ، حيث أن نسبة الوفيات من المرض عالية



إدماج خبرات الأطباء



منصة على الويب توفر خدمة تشخيص السرطان عن طريق صور الشرائح الكاملة (WSIs)

تحليل SWOT

نقاط القوة

- استخدام الذكاء الاصطناعي.
 - انشاء خرائط حرارية.
- منصة متكاملة سهلة الاستخدام
- عدم وجود منافسين مباشرين
 - دعم القرار الطبي.

S

نقاط الضعف

- محدودية الموارد المالية و البشرية
 - نقص الوعى المحلى.
- تحديات في الحصول على تراخيص رسمية
 - الحاجة إلى بيانات طبية.

W

الفرص

- الشراكة مع المؤسسات الصحية قلة المنافسة في السوق المحلية
 - از دياد الاهتمام بالتحول الرقمي.
 - دعم المشاريع المبتكرة والتكنولوجيا من قبل الحكومة
 - إمكانية التوسع

التهديدات

- دخول شركات أجنبية
 - التردد في اعتماد تقنيات الذكاء
 - الاصطناعي
- القيود القانونية المرتبطة باستخدام الذكاء الاصطناعي

نموذج العمل التجاري BMC

الشركاء	الأنشطة الرئيسية	القيمة المقدمة	العلاقات مع الزبائن	شرائح العملاء
- جامعة بسكرة - المصحات والمستشفيات - مراكز البحث - جمعيات و منظمات الأمراض والأورام	- البحث والتطوير - تصميم وتطوير نماذج الذكاء الإصطناعي - تقييم نتائج النموذج من خلال المخابر - تطوير تطبيق الويب الموارد الرئيسية - اشتراكات الخدمات السحابية - أدوات وبرامج تطوير البرمجيات - فريق التطوير	- تحديد نوع سرطان الرئة - الكشف عن المناطق المتضررة بالسرطان - استخدام خوارزميات التعلم العميق - واجهة مستخدم سهلة الاستعمال - تقليل وقت التشخيص بشكل ملحوظ أمان بيانات المرضى	- الشبكات الإجتماعية المشاركة التحديثات وتلقي المقترحات الموقع الإلكتروني الخاص بالخدمة حدم فني متخصص القنوات الطبية والمعارض التسويق بالمحتوى الشتراكات عبر الإنترنت حراكات مع المخابر المراكات مع المخابر المراكات مع المخابر	- أخصائي علم الأمراض (Pathologists) - مخابر التحاليل والتشخيص - المصحات - طالب الطب والباحثون
التكاليف			مصادر الإيرادات	
Ď Ď		- نفقات البحث والتطوير - التسويق - تكاليف الخدمات السحابية	- تحدیث النماذج	- الإشتراكات - نموذج Freemium - هجةود الخدمة و الصيانة