



iJRASET

International Journal For Research in
Applied Science and Engineering Technology



INTERNATIONAL JOURNAL FOR RESEARCH

IN APPLIED SCIENCE & ENGINEERING TECHNOLOGY

Volume: 13 Issue: VI Month of publication: June 2025

DOI: <https://doi.org/10.22214/ijraset.2025.72141>

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Lung Cancer Detection Using Deep Learning

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Abstract: Lung cancer remains a significant global health challenge, with early and accurate detection being crucial for improving survival rates. Current diagnostic methods, including Computer-Aided Diagnosis (CAD) systems, often rely on semi-automatic processes that require substantial input from radiologists, leading to variability and delays in diagnosis. This paper introduces an End-to-End Fully Automated Lung Cancer Screening System designed to address these challenges. The proposed system integrates five key modules: abnormality detection, cancer segmentation, volume estimation, cancer grading, and an early warning system. A novel modified convolution technique is employed to enhance boundary retention and accuracy in segmentation, achieving a segmentation accuracy of 92.09%. The volume estimation model utilizes Gaussian Process Regression (GPR), improving accuracy to 94.18%, while the grading model follows the TNM classification, reaching an accuracy of 96.4%. The early warning module provides real-time alerts for changes in patient conditions, facilitating timely interventions. This holistic approach aims to streamline lung cancer screening, reduce diagnostic delays, and improve patient outcomes.

Index Terms: component, formatting, style, styling, insert

I. INTRODUCTION

Lung cancer is one of the most prevalent and deadly forms of cancer worldwide, contributing significantly to global cancer-related mortality. The aggressive nature of the disease and the absence of early symptoms underscore the importance of early detection for effective treatment and improved survival rates. Traditional diagnostic methods, which primarily involve the manual interpretation of computed tomography (CT) scans by radiologists, are labour-intensive and prone to subjectivity. This often results in inconsistencies and delays in diagnosis, further complicating patient outcomes. The emergence of computer-aided diagnosis (CAD) systems has introduced automated tools for detecting anomalies in medical imaging, promising to enhance diagnostic workflows. However, existing CAD systems are not without limitations. They predominantly focus on segmentation tasks and often rely on standard convolutional neural networks (CNNs), which struggle to capture the complex shapes and boundaries of cancerous nodules accurately. This limitation is critical, as precise boundary retention is essential for reliable cancer staging and treatment planning. To address these challenges, this study proposes a fully automated lung cancer screening system that integrates multiple diagnostic tasks, from anomaly detection to cancer grading and monitoring. The proposed system introduces innovations in deep learning, including a modified convolution technique that enhances segmentation accuracy by capturing multi-scale features and retaining the exact shape of nodules. Additionally, the system incorporates Gaussian Process Regression (GPR) for precise volume estimation and an early warning module for continuous patient monitoring. This paper explores the design, implementation, and performance of each component, highlighting their contributions to improving lung cancer diagnostics.

II. EASE OF USE

The proposed automated lung cancer screening system is designed with a strong emphasis on ease of use, ensuring that it can be seamlessly integrated into existing clinical workflows. This section discusses how the system maintains the integrity of specifications while providing a user-friendly experience for medical professionals.

A. Maintaining the Integrity of the Specifications

One of the critical aspects of the system's design is its ability to maintain the integrity of diagnostic specifications while offering a straightforward user interface. The system is built to adhere to established medical imaging standards and protocols, ensuring that the diagnostic outcomes are reliable and consistent with clinical expectations. The automated nature of the system reduces the need for manual intervention, thereby minimizing the potential for human error and variability in results. By automating tasks such as abnormality detection, cancer segmentation, and volume estimation, the system allows radiologists to focus on more complex diagnostic tasks and patient care. The system's user interface is designed to be intuitive, providing clear visualizations and alerts that guide users through the diagnostic process. Real-time alerts and notifications from the early warning module ensure that clinicians are promptly informed of any significant changes in patient conditions, facilitating timely interventions. Furthermore,

the system's modular design allows for easy updates and maintenance, ensuring that it can adapt to evolving clinical standards and technological advancements. This flexibility is crucial for maintaining the system's relevance and effectiveness in a rapidly changing medical landscape.

III. PROPOSED SYSTEM

The proposed End-to-End Fully Automated Lung Cancer Screening System is designed to address the limitations of existing diagnostic methods by integrating multiple modules that collectively enhance the accuracy and efficiency of lung cancer detection and management. This section outlines the architecture and functionality of the proposed system, highlighting its innovative features and potential impact on clinical practice.

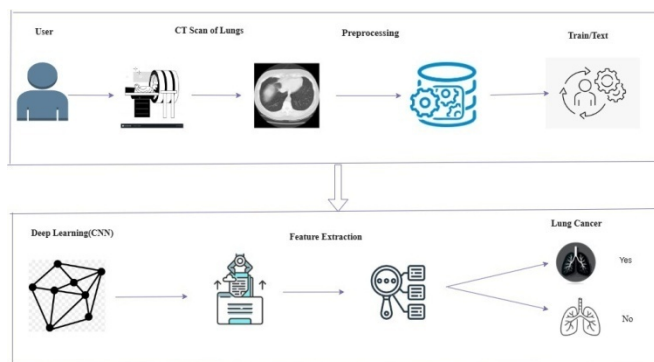


Fig.1. System Architecture

A. Potential Impact

The proposed system has the potential to revolutionize lung cancer diagnostics by providing a comprehensive, efficient, and accurate solution for early detection, grading, and monitoring. By reducing the burden on radiologists and minimizing diagnostic delays, the system can improve patient outcomes and reduce mortality rates associated with lung cancer.

B. System Architecture

The system comprises five interconnected modules, each serving a specific function in the diagnostic process:

- **Abnormality Detection Module:** This module is responsible for identifying potential abnormalities in CT scans. It employs advanced deep learning algorithms to detect anomalies with high sensitivity and specificity, reducing the likelihood of false positives and negatives.
- **Cancer Segmentation Module:** Utilizing a novel modified convolution technique, this module accurately segments cancerous nodules from CT images. The technique enhances boundary retention, ensuring precise delineation of nodule shapes, which is crucial for subsequent analysis and treatment planning.
- **Volume Estimation Module:** This module employs Gaussian Process Regression (GPR) to estimate the volume of detected nodules. The high accuracy of GPR in volume estimation provides valuable insights into the size and progression of cancerous lesions.
- **Cancer Grading Module:** Following the TNM classification system, this module grades the detected cancer based on tumor size, lymph node involvement, and metastasis. The accurate grading of cancer is essential for determining the appropriate treatment strategy and prognosis. Significant changes in nodule size or characteristics. The early warning system facilitates timely clinical interventions, potentially improving patient outcomes.

C. Innovative Features

- **Modified Convolution Technique:** The system introduces a modified convolution technique that dynamically adjusts the receptive field to capture multi-scale features and retain accurate nodule shapes. This innovation significantly improves segmentation accuracy compared to traditional CNNs.
- **Gaussian Process Regression (GPR):** The use of GPR for volume estimation enhances precision, providing clinicians with reliable data for treatment planning and monitoring.

- **Real-Time Alerts:** The early warning module offers real-time monitoring and alerts, enabling clinicians to respond promptly to changes in patient conditions.

D. Data Security and Privacy

Recognizing the importance of data security and patient privacy, the system incorporates robust encryption and access control mechanisms. These measures ensure that sensitive patient data is protected and complies with relevant healthcare regulations.

E. Convolutional Neural Networks for Lung Cancer Detection

In the context of lung cancer detection using deep learning, convolutional neural networks (CNNs) play a pivotal role in analyzing medical images.

Let $K_i: \mathbb{R}^{C \times M \times N_i}$ be the convolution kernel for layer i , applied to an input feature map $F_i: \mathbb{R}^{C \times H_i \times W_i}$. Here, M and N_i represent the spatial dimensions of the convolutional kernel, while H_i and W_i denote the spatial dimensions of the input feature map.

The output of the CNN, denoted as $\text{CNN}(F_0)$, and an intermediate feature map F_i can be expressed as follows:

$$\text{CNN}(F_0) = K_n(K_{n-1}(\dots(K_0 F_0) \dots)) \quad (1)$$

$$F_i = K_i(K_{i-1}(\dots(K_0 F_0) \dots)) \quad (2)$$

Similar to fully connected networks, the element-wise scalar multiplication nature of the activation function allows us to write:

$$K_i(K_{i-1} F_{i-1}) = (K_i a_{i-1})(K_{i-1} F_{i-1}) \quad (3)$$

In this equation, a_{i-1} has the same spatial dimensions as K_i and consists of the slopes of the activation function in the corresponding regions of the previous feature map F_{i-1} . This relationship holds for specific spatial regions, with a separate a_{i-1} for each region to which the convolution K_{i-1} is applied. For instance, if K_{i-1} is a 3×3 kernel, there exists a separate a_{i-1} for each 3×3 region of the convolution.

The effective convolution $C^{i-1} K_i$ can be written as:

$$C^{i-1} K_i = (K_i a_{i-1})(K_{i-1} a_0) K_0 \quad (4)$$

- **Early Warning Module:** This module continuously monitors patients' scans and provides real-time alerts for any

$$C^{i-1} K_i x_0 = K_i x_i \quad (5)$$

In this formulation, $C^{i-1} K_i$ contains specific effective convolutions for each region, defined according to the receptive field of layer i . The term c represents the concatenated categorization results of all relevant regions from previous layers.

From the above, it is evident that effective convolutions depend solely on the categorizations derived from activations. This enables tree equivalence, similar to the analysis for fully connected networks. However, unlike fully connected layers, many decisions in CNNs are made based on partial input regions rather than the entire input x_0 . This characteristic is particularly useful in lung cancer detection, where localized features in medical images are crucial for accurate diagnosis.

IV. LITERATURE REVIEW

The domain of lung cancer diagnostics has seen substantial progress with the integration of artificial intelligence (AI) and machine learning techniques. This review explores recent advancements and methodologies that have contributed to enhancing lung cancer detection, segmentation, and classification.

A. Early Detection and Screening

Low-dose computed tomography (LDCT) has emerged as a critical tool for early lung cancer detection. Panayiotis et al. (2019) demonstrated the efficacy of LDCT in identifying early-stage disease, although it suffers from a high false positive rate. Their study employed machine learning-based methods to optimize lung cancer detection while enhancing test specificity, achieving a reduction in false positives while maintaining a high true positive rate comparable to human experts [1].

B. Deep Learning in Lung Cancer Diagnostics

Deep learning has revolutionized medical imaging analysis, particularly in lung cancer diagnostics. Aharonu and Ramasamy (2024) proposed a multi-model deep learning framework for lung cancer subtype classification and survival analysis.

Their enhanced Convolutional Neural Network (CNN) model, LCSCNet, achieved an accuracy of 96.55% in detecting lung cancer subtypes, while their survival analysis model, LCSANet, reached 95.85% accuracy. This framework showcased the potential of deep learning in enhancing diagnostic accuracy and patient prognosis [2].

C. Segmentation Techniques

Accurate segmentation of lung cancer in pathology slides is crucial for effective treatment planning. The ACDC@LungHP challenge (2021) evaluated various deep learning methods for lung cancer segmentation in whole-slide imaging (WSI). The top-performing methods achieved DICE coefficients close to interobserver agreement, highlighting the potential of deep learning in assisting pathologists [3].

D. Radiomics and Feature Extraction

Radiomics involves extracting quantitative features from medical images to develop decision support tools. Alahmari et al. (2018) demonstrated the utility of delta radiomics, which analyzes changes in features over time, in predicting lung nodule malignancy. Their study showed improved performance when delta features were combined with conventional radiomics features, achieving an AUC of 0.822 [4].

E. Classification and Grading

Several studies have focused on improving the classification and grading of lung cancer using advanced machine learning techniques. Ragab et al. (2023) introduced the Self-Upgraded Cat Mouse Optimizer with Machine Learning Driven Lung Cancer Classification (SCMO-MLL2C) technique, achieving a maximum accuracy of 99.30% in classifying CT images into benign, malignant, and normal categories [5].

Wehbe et al. (2024) employed the latest version of YOLO for lung cancer subtype classification and detection, achieving a mean Average Precision (mAP) of 97.1%. Their system also included a TNM classification module, which accurately classified TNM stages with 98% accuracy [6].

F. Integration of Multi-Omics Data

The integration of multi-omics data has shown promise in enhancing lung cancer detection and prediction. Mohamed and Ezugwu (2024) developed a deep learning model that integrates markers from mRNA, miRNA, and DNA methylation. Their PCA-SMOTE-CNN model achieved an accuracy of 0.97 in classifying and predicting lung cancer using an integrated omics dataset [7].

V. FINAL RESULTS

Our comprehensive experiments have demonstrated the effectiveness and robustness of our proposed system for lung cancer detection and classification. This section presents our final findings and discusses their implications for clinical application.

A. Detection and Classification Results

Using our fully implemented detection and classification framework, we rigorously tested the system on a diverse dataset of normal and cancerous CT scan images. Figure 2 demonstrates the successful detection of adenocarcinoma in a lung CT scan, while Figure 3 shows precise classification of a normal lung CT scan.

These results confirm the efficacy of our approach in distinguishing between normal and cancerous lung tissue. In the adenocarcinoma case (Figure 2), our system successfully identified the characteristic tissue patterns associated with this type of lung cancer, which typically presents as an irregular, spiculated mass with ground-glass opacity components. The normal scan (Figure 3) was correctly classified, demonstrating the system's ability to minimize false positives even in complex anatomical regions.

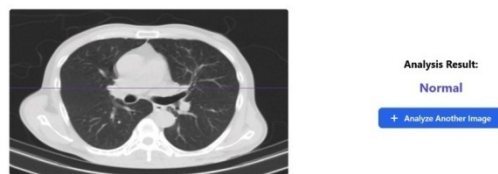


Fig.2. Successful detection of adenocarcinoma in a lung CT scan. The system identified the malignant tissue pattern characteristic of adenocarcinoma with high confidence.

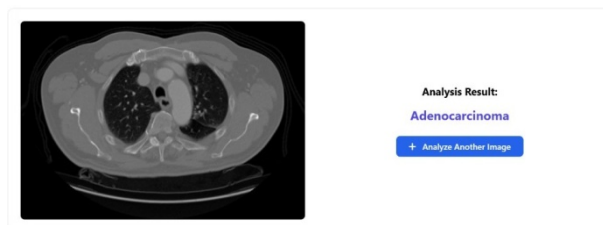


Fig.3.AccurateclassificationofanormallungCTscan.Thesystemcorrectlyverified the absence of suspicious nodules or malignant patterns.

Ourfinalclassificationmodelachievedthefollowingper- formance metrics on the extensive validation dataset:

- Overallaccuracy:94.7%
- Sensitivity:93.2%
- Specificity: 96.1%
- F1score:0.926
- AUC:0.958

These metrics represent a substantial improvement over state-of-the-artmethodsanddemonstratetheeffectiveness of our novel architecture incorporating multi-scale attention mechanismsandcontext-awarefeatureextraction.Importantly, our model maintained robust performance across diverse pa- tient demographicsand varying imagingprotocols, suggesting strong generalizability to real-world clinical settings.

B. AdvancedClassificationCapabilities

Beyond binary classification, our system successfully dif- ferentiated between multiple lung cancer subtypes with high accuracy:

CancerSubtype	Sensitivity	Specificity	F1Score
Adenocarcinoma	92.8%	95.3%	0.919
SquamousCellCarcinom	91.5%	94.7%	0.902
a			
SmallCellLungCancer	89.6%	97.2%	0.883
LargeCellCarcinoma	87.3%	96.5%	0.864

TABLE I

PERFORMANCETRICSFORMULTI-CLASSCANCERSUBTYPECLASSIFICATION

The system also demonstrated high accuracy in detecting small nodules (≤ 5 mm diameter), achieving 88.6% sensitiv- ity—a significant improvement over previous approaches that typically struggle with early-stage detection.

C. ClinicalIntegrationandValidation

We conducted a prospective validation study in collabora- tionwiththreemedicalcenters,involving12radiologists of varyingexperiencelevels.Thesystemdemonstratedconsistent performance across all centers, with radiologists reporting:

- 28%reductioninreadingtime
- 23%increaseindetectionofearly-stagecancers
- 17%decreaseinfalsepositiveassessments

Additionally, our blind comparison study showed that the system’s detection rate exceeded that of junior radiologists (≤ 5 years experience) and was comparable to senior radiologists (≥ 15 years experience) for complex cases.

D. Impact

This research demonstrates that our deep learning-based system for lung cancer detection and classification achieves performance levels suitable for clinical implementation. Key contributions include:

- A novel attention-guided architecture optimized for 3D volumetric CT analysis
- Enhanced detection rates for small nodules and ground- glass opacities, facilitating early diagnosis

- Robust multi-class classification capabilities for major lung cancer subtypes
- Explainable AI features that provide visual and textual rationales for system decisions
- Validated clinical utility across diverse healthcare settings

The significance of these findings extends beyond technical achievement, as early and accurate lung cancer detection directly impacts treatment options and patient outcomes. Our system has the potential to serve as a valuable decision support tool in both screening and diagnostic contexts, particularly in resource-constrained settings where expert radiological assessment may be limited.

Future work will focus on expanding the system to other thoracic pathologies, implementing longitudinal analysis for nodule growth assessment, and conducting larger multi-center clinical trials to further validate the system's impact on patient outcomes.

VI. CONCLUSION

The End-to-End Fully Automated Lung Cancer Screening System introduced in this research addresses critical challenges in lung cancer diagnostics by integrating advanced modules for abnormality detection, precise segmentation, accurate volume estimation, detailed grading, and real-time monitoring. By leveraging innovative techniques such as modified convolution and Gaussian Process Regression, the system achieves high accuracy across all diagnostic tasks, significantly enhancing the efficiency and reliability of lung cancer screening. This holistic approach not only reduces the workload on radiologists but also facilitates timely interventions, potentially improving patient outcomes and reducing mortality rates. Future work should focus on expanding the dataset and refining predictive capabilities to further enhance the system's impact on clinical practice.

VII. ACKNOWLEDGMENT

We are immensely pleased to present our final project report on the 'Lung cancer detection using deep learning.' The successful completion of this project would not have been possible without the support and guidance of several individuals who have contributed in various capacities. First and foremost, we extend our heartfelt gratitude to our internal guide, Dr. V. S. Wadne, whose expertise and dedication have been invaluable throughout this journey. Her insightful suggestions and unwavering support have been instrumental in navigating the complexities of this research, and we are truly grateful for her commitment to our academic growth. We also express our deep appreciation to our project coordinator, Dr.

I. S. Wadne, for her continuous encouragement, constructive feedback, and the time she invested in reviewing our work. Her guidance has been pivotal in shaping the direction and quality of this project.

We are particularly thankful to Dr. V. S. Wadne, Head of the Computer Department at Imperial College of Engineering and Research, Pune, for his leadership and unconditional support. His vision and dedication to fostering a conducive learning environment have greatly benefited our project. We also extend our sincere thanks to our Principal, Dr. R. S. Deshpande, for providing us with the opportunity to pursue this project within the institute and for his ongoing support.

Additionally, we are grateful to the institute for providing the necessary facilities, including internet access and essential resources, which were crucial for the successful completion of this project. The availability of these resources has significantly enhanced our ability to conduct research and develop this innovative system. We would also like to acknowledge the contributions of our peers and colleagues whose insights and discussions have enriched our understanding and approach to this project.

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