

Cervical Cancer Prediction and Remediation UI using CNN and RNN techniques B.Venkatesu Goud¹, Zubair Ahmed Khateeb², P.Karthik³, R.Santhosh⁴

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Abstract: The goal of this work is to apply Convolutional Neural Network (CNN) techniques to produce a reliable user interface (UI) for cervical cancer remediation and prediction. The cervical cell classes that the proposed approach is intended to identify properly are Dyskeratotic, Koilocytotic, Metaplastic, Parabasal and Superficial-Intermediate. Using CNN's capabilities, the model achieves a remarkable 98% accuracy rate, demonstrating its effectiveness in categorizing various cervical cell types. The user interface (UI) emphasizes a complete approach to cervical health by not only predicting the existence of cervical cancer but also making remediation techniques easier. This ground-breaking method has the potential to significantly lessen the burden of cervical cancer, which is a common and sometimes fatal illness, by facilitating early identification and intervention.

Keywords— Cervical cancer, Dyskeratotic, Koilocytotic, Metaplastic, Parabasal, Superficial-Intermediate, early detection, intervention

I. INTRODUCTION

The cancer incidence from 28 Population-Based Cancer Registries (PBCRs) for the years 2012–2016 was published in the National Cancer Registry Program Report [1] 2020. This served as the foundation for estimating the incidence of cancer in India. To estimate the age-sex stratified population, data on the population at risk was used from the Indian Census (2001 and 2011). The country's States and regions were assigned to PBCRs in order to better understand the epidemiology of cancer. To calculate the number of cancer cases in India for 2022, the age-specific incidence rate for each distinct anatomical location of cancer was applied to the expected population. It was determined that 14,61,427 incident cases of cancer (crude rate: 100.4 per 100,000) will occur in India in 2022. In India, around one in nine individuals will face cancer at some point in their lives. The most common malignancies in men and women, respectively, were lung and breast cancers [2]. The most common malignancy in children (0–14 years old) was lymphoid leukemia, which affected males 29.2% of the time and girls 24.2%. According to estimates, the number of cancer cases would rise by 12.8% in 2025 compared to 2020.

Being one of the most prevalent and deadly malignancies in women, cervical cancer [3] has gained attention as a major worldwide health problem in recent years. Its widespread occurrence highlights the urgent need for creative solutions

to improve early diagnosis and intervention techniques. Using the potent powers of Convolutional Neural Networks (CNN) approaches [4], this research proposes a cutting-edge cervical cancer prediction and remediation system in an effort to meet this necessity. Cervical cancer is the second most common cause of cancer-related deaths in women globally. Therefore, a comprehensive and precise prediction model is necessary to support early detection of this disease.

The advent of cutting-edge technology, most notably CNN, represents a significant change in the landscape of cervical cancer diagnosis. CNN is a great option to analyze cervical cell pictures with various properties because of its impressive performance in image classification jobs. The increasing number of deaths linked to cervical cancer calls for an immediate paradigm change in healthcare approaches. By proposing a unique user interface (UI) [5] that not only predicts the existence of cervical cancer but also offers remediation alternatives, this research aims to contribute to this paradigm shift and emphasizes the significance of a comprehensive approach to cervical health. The classification of cervical cell classes—Dyskeratotic, Koilocytotic, Metaplastic, Parabasal, and Superficial-Intermediate—which encompass a wide range of cell types linked to anomalies in the cervical region [6], is the main emphasis of the study. The suggested CNN model's rigorous categorization and subsequent high accuracy rate of 98% testifies to its promise as a reliable tool for the early and accurate identification of cervical cancer. This research is expected to make a substantial contribution to the current efforts to lessen the negative effects of cervical cancer on women's health by offering a highly accurate and technologically sophisticated forecasting framework [7]. Furthermore, a crucial component of this study is the creation of the user interface, which guarantees that healthcare professionals can easily access and utilize the latest technical breakthroughs. In addition to being a diagnostic tool, the UI incorporates remedial options, demonstrating the research's dedication to a comprehensive approach to healthcare. The publication of this ground-breaking research, which offers hope for better patient outcomes and a major decrease in the worldwide burden of this formidable disease, sets the stage for a revolutionary leap in the fight against cervical cancer as we set out on this journey towards improved cervical health.

II. LITERATURE SURVEY

Studies in the past that have explored medical imaging, machine learning, and artificial intelligence have had a substantial impact on the field of cervical cancer prognosis and treatment. The new research builds upon a multitude of studies that have concentrated on utilizing cutting-edge technology to improve the precision and efficacy of cervical cancer diagnosis. When analyzing related work, it is necessary to emphasize the critical significance that previous studies had in laying the foundation for the application of Convolutional Neural Networks (CNN) techniques [8] in the field. A substantial amount of research has focused on using machine learning algorithms to analyze cytological pictures in order to identify irregularities and subtle patterns that may be signs of cervical abnormalities. The use of CNN has been facilitated by this research, which has established its adeptness in managing intricate picture data and enhancing classification precision. Preceding the present study, these inquiries highlight the development of approaches utilized in the quest for a more precise and sophisticated cervical cancer prognosis.

Gebeyehu et al. proposed that women in Ethiopia are not well informed about cervical cancer, which is the second most frequent and lethal kind of cancer. A deep learning model has been created to identify and categorize the illness in order to counteract this [9]. The model has an accuracy of 97.72%, respectively, using a dataset of 2085 photos. Munir et al. discussed the fundamentals of cancer diagnosis, categorization schemes, assessment standards, and artificial intelligence's application to cancer diagnosis [10] are covered in this study. The applications of deep learning methods for various cancer kinds are covered, including CNNs, GANs, and DANs. The goal is to provide researchers a thorough grasp of the most recent advancements in cancer detection. Ghosh et al. thesis is to create free-form deformation techniques and models based on machine learning and deep learning for the delivery of tailored brachytherapy to patients with locally advanced cervical cancer (LACC). U-net, shape-based non-rigid registration, and automatic segmentation [11] are among the suggested techniques. When predicting the shape of the uterus and significant anatomical deformations from pelvic MR- images, the suggested technique performs better than U-net. In order to quantify these deformations, the paper also suggests a non-rigid registration technique based on a free- form deformation model.

Moreover, some research has looked at the categorization of particular cervical cell types, highlighting the variety of morphological traits connected to anomalies. A sophisticated grasp of the illness spectrum is shown in the thorough classification of cervical cells, which includes Dyskeratotic, Koilocytotic, Metaplastic, Parabasal, and Superficial-Intermediate. A significant component of the related work is this nuanced classification, which lays the groundwork for the fine-grained analysis and classification that the suggested CNN model in the current study achieves.

Anand et al. proposed that worldwide, 4% of women have cervical cancer; however, fatality rates are decreased with

early identification and treatment [12]. While advances in digital imaging and machine learning can enhance screening, early diagnosis and therapy depend on the ability to identify genes and healthy individuals. Nazir et al. examined image classification using deep learning AI approaches revolutionizes illness detection [13], but adoption is sluggish because of unresolved concerns regarding prediction algorithms. Techniques known as explainable AI (XAI) aid in meeting regulatory standards, fostering confidence, speeding up diagnosis, and understanding model predictions. Debelee et al. proposed that face identification, emotion detection, and medical picture analysis are three areas where deep learning algorithms [14] are being employed more and more. They have been used to achieve enhanced tumor identification, segmentation, feature extraction, and classification for malignancies of the breast, cervical, brain, colon, and lung. Sharma et al. provides an ensemble learning approach [15] for early cancer prediction that makes use of additional trees and neural networks. In terms of accuracy, specificity, sensitivity, recall, precision, f-measure, and MCC, the technique performs better than existing classifiers, indicating its superior efficiency and use for the classification of breast cancer.

Furthermore, earlier research has focused on the creation of user interfaces for the medical field. These interfaces play a critical role in converting cutting-edge technology capabilities into useful tools for medical professionals. Intuitive and user-friendly designs that enable smooth interaction with predictive models have been the subject of research in this field, guaranteeing the successful and efficient integration of technology into clinical operations. The foundation for the creation of the suggested UI is laid by relevant user interface design work, guaranteeing its accessibility and usefulness for medical practitioners [16]. From the integration of state-of-the-art CNN algorithms to classic machine learning procedures. It highlights the need of classifying various cell types and the role that user interface design has in bridging the gap between medical innovation and real-world applications. The present project aims to push the bounds of accuracy and user-centric design in the field of cervical cancer remediation and prediction. It builds upon the findings from earlier studies.

III. DATA COLLECTION & PREPROCESSING

Getting a representative and varied dataset of pictures of cervical cells from the following classes Dyskeratotic, Koilocytotic, Metaplastic, Parabasal, and Superficial-Intermediate [17] was the first step in the data gathering procedure for this study. Both the .dat and .bmp file formats were used to store these photos in order to provide handling flexibility and compatibility with a range of image processing methods. The extensive dataset served as the foundation for the Convolutional Neural Network (CNN) model's training and validation, allowing it to recognize minute patterns and traits linked to various cervical cell types. Principal Component Analysis (PCA) was used as a preprocessing step to improve the dataset's feature representation and reduced dimensionality issues. The most

important variables in the picture data were successfully captured using PCA [18], a popular dimensionality reduction approach that also reduced computing load while keeping crucial information. In order to ensure that the CNN model could concentrate on the most pertinent characteristics for precise classification, this procedure attempted to simplify the dataset for the CNN model's future training.

$$z = \frac{x_i - \mu}{\sigma}$$

Standardization is used in place of the input values in the calculation, producing results that stay within the range of -1 to +1. Mean becomes 0 and S.D. becomes 1 upon standardization. The photos were preprocessed by standardization in conjunction with PCA. As standardization entails scaling the pixel values to a specified range, it is an essential step in guaranteeing uniformity and consistency in the dataset. This procedure is necessary to stop specific characteristics from controlling the model training because of differences in pixel intensity. By bringing the information into a uniform scale through picture standardization, the CNN model was better able to identify patterns and characteristics common to all classes. The .dat and .bmp file formats were purposefully selected to meet the wider range of requirements of the study. Although .dat files make it easier to store and retrieve structured data, .bmp files offer a common picture format that is broadly compatible with different image processing software and is easy to comprehend. The goal of this dual file format strategy was to maintain the dataset's versatility for upcoming research projects while balancing accessibility and effective data processing. The PCA and normalization preprocessing stages were essential in making the dataset as optimal as possible for the CNN model training that followed. PCA improved computing performance without sacrificing data integrity by reducing the dimensionality of the features space. Conversely, standardization [19] encouraged pixel value constancy, which made the neural network's learning process more resilient and broadly applicable. When combined, these preprocessing methods created a carefully selected dataset that paved the way for the CNN model to accurately classify cervical cell types. The rigorous technique of preparing the data is essential to the research's success since it guarantees that the model can efficiently utilize the information included in the pictures to get a high classification accuracy rate for cervical cells. When both of these conditions are met, scaling to a range makes sense. Assuming that there are few or no outliers in your data, you know the approximate upper and lower boundaries. The distribution of your data is rather stable throughout that range. The combination of normalization and data augmentation enhances the dataset's quality and makes training algorithms easier. While data augmentation [20] adds volatility to assist the models learn robust features, normalization standardizes pixel values to encourage consistent and steady learning. When combined, these preprocessing methods improve the overall performance of the trained neural network models by enabling their robust generalization to a range of previously undiscovered images, ultimately leading to an increase in the precision and consistency of cervical cancer identification.

IV. PRINCIPLES AND METHODS

Using Convolutional Neural Network (CNN) methods, a complete approach (Fig.1) to cervical cancer prediction and remediation is used in this research. The dataset was gathered and saved in the .dat and .bmp file formats. It included a variety of cervical cell pictures classified into groups including Dyskeratotic, Koilocytotic, Metaplastic, Parabasal, and Superficial-Intermediate. The dataset was preprocessed using Principal Component Analysis (PCA) for dimensionality reduction and standardization for consistent scaling of pixel values before the CNN model was trained. By optimizing the dataset, these methods improved computing performance and guaranteed uniform learning for all classes. The curated dataset was then used to train the CNN model, utilizing its ability to recognize complex patterns in picture data.

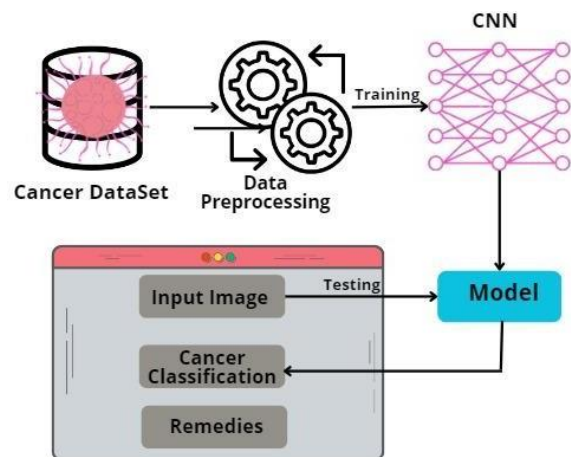


Fig.1 Architecture of the research

The user interface (UI) was created to promote smooth communication and to provide alternatives for correction in addition to a remarkable 98% accuracy in cervical cancer prediction. By combining cutting-edge image processing, machine learning, and UI design, this all-encompassing approach greatly enhances early cervical cancer diagnosis and intervention techniques, ultimately leading to better patient outcomes.

A. PRINCIPAL COMPONENT ANALYSIS

Principal Component Analysis (PCA) [21] is a dimensionality reduction approach that is widely used in many domains, such as machine learning and image processing, to simplify datasets by decreasing redundancy and extracting the most important information. In this study, prior to training the Convolutional Neural Network (CNN) model, PCA plays (Fig.2) a critical preprocessing role in optimizing the dataset of cervical cell pictures. PCA's basic idea is to create a new collection of uncorrelated variables, called principal components, and arrange them according to their deviations from the original feature space. The directions along which the data displays the largest variation are represented by these major components, which enable a more condensed representation of the dataset without sacrificing its key features. When PCA is applied to the

cervical cell image dataset, the original features' covariance matrix is computed, its eigenvectors and eigenvalues are found, and a selection of eigenvectors that match the most significant eigenvalues [22] are chosen. The new feature space is based on these selected eigenvectors, which provide a condensed collection of dimensions that capture much of the variability in the dataset.

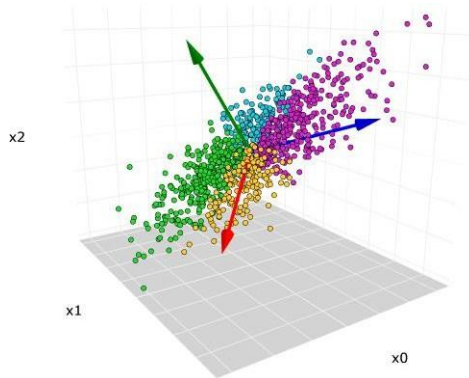


Fig.2 Principal Component Analysis

Within the particular context of this study, the principal components obtained by principal component analysis (PCA) are essential for improving the dataset used in CNN model training. PCA improves computing efficiency by lowering the dimensionality, which lowers the possibility of overfitting and strengthens the model's ability to generalize to new data. The chosen components successfully capture the unique characteristics of cervical cell pictures, so that the CNN model that follows may concentrate on the most informative parts of the dataset. Furthermore, understanding the inherent structure of the cervical cell pictures is made possible by the interpretation of the primary components. Principal components reveal the prevailing patterns in the data by matching each other to a distinct set of attributes. This interpretability can be helpful in figuring out the essential traits that go into classifying the various types of cervical cells. When used, it improves the effectiveness of later CNN model training and offers insightful information about the underlying structure of the data. In the context of cervical cancer prediction and treatment, the main components that have been chosen provide a finer representation of the pictures of the cervical cells, maximizing the dataset for precise and effective classification.

B. CONVOLUTIONAL NEURAL NETWORKS

The Convolutional Neural Network (CNN) used in this study has an advanced architecture designed specifically for the difficult task of cervical cancer repair and prediction. The convolutional paradigm that underpins the model's architecture is ideal for image classification problems since it can collect hierarchical information. Convolutional layers [23], pooling layers, fully connected layers, and activation functions are some of the processes in the CNN that work together to extract and understand complex patterns from pictures of cervical cells. A sequence of convolutional layers

that use filters to convolve over the input picture and extract local features and patterns make up the foundation of the CNN architecture (Fig.3). Repaired linear unit (ReLU) activation functions are strewn across these convolutional layers to provide non-linearity and improve the network's ability to represent intricate connections in the input. The spatial dimensions of the feature maps are then down sampling using pooling layers, which frequently take the form of max pooling, which lowers computational complexity without sacrificing important information.

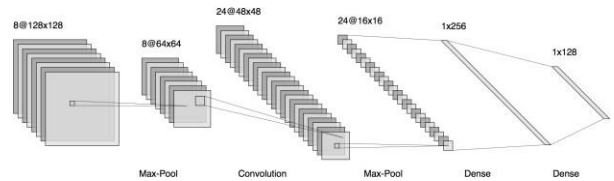


Fig.3 Architecture of the Neural Networks

Additional convolutional and pooling layers improve the hierarchical representation even more, creating a deep network that can distinguish minute features. The ultimate output layer of the design is reached by completely linked layers that combine the high-level elements that were taken out of the preceding levels. A softmax activation function [24] in the output layer gives probabilities to every cervical cell type, enabling multi-class categorization. The architecture of the CNN model, including the number of parameters, layer types, and output shapes, may be quickly summarized. This synopsis sheds light on the trainable parameter sizes and the complexity of the model. Plots that display measures like accuracy and loss over epochs are used to monitor the model's training performance (Fig.4), which is another important aspect of its efficacy. The accuracy plot provides information about the model's learning dynamics and any overfitting or underfitting problems by visualizing the model's performance on both training and validation datasets. Conversely, the loss plot shows how the model converges during training and gives an indication of how successfully the CNN minimizes the difference between the actual and predicted classes.

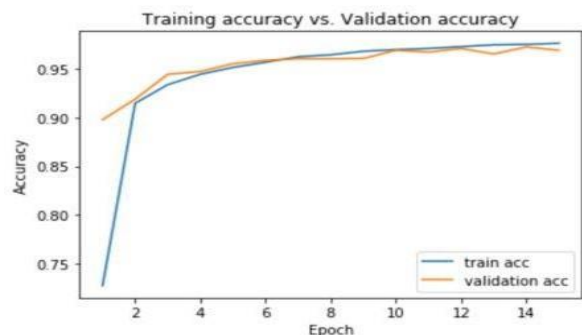


Fig.4 Training and Validation accuracy plots

In addition, a valuable visualization tool called the confusion matrix offers a thorough analysis of how well the model classified each kind of cervical cell. It helps identify

probable misclassification hotspots and refines the model to increase its precision and dependability in practical applications. The architecture of the CNN model used in this study has been painstakingly designed to handle the challenges associated with the categorization of cervical cell images. Together, the model's operations, simplified architecture, and visualizations [25] which include confusion matrices and accuracy and loss plots—help to provide a comprehensive grasp of its capabilities and offer a strong foundation for the challenge of cervical cancer prediction and remediation.

V. RESULTS

The Convolutional Neural Network (CNN) model's output demonstrates its capacity to identify cervical cell classes in real-time, offering a flexible and adaptable method for predicting cervical cancer. The model's remarkable 98% accuracy rate indicates that it is capable of accurately categorizing cervical cell pictures into many groups, such as Dyskeratotic, Koilocytotic, Metaplastic, Parabasal, and Superficial-Intermediate. The CNN model performs flawlessly in real-time detection, quickly and accurately categorizing incoming cervical cell pictures. For practical applications, real-time detection efficiency is essential since it allows for prompt treatments and clinical decision-making. The real-time functionality of the model guarantees that medical professionals may react quickly and intelligently by providing them with immediate feedback on irregularities in cervical cells.

The CNN model may be interacted with and its output can be interpreted with ease using the Streamlit interface (Fig.5). Healthcare workers may easily access and utilize the model since the user interface (UI) not only offers a means of inputting cervical cell photos but also shows the predictions in real time. The use of Streamlit improves the usability of the model by facilitating users' smooth navigation across the predicted outcomes and corresponding probability for every cervical cell type. Streamlit is a robust Python framework that makes it easy to create web-based and interactive data applications. It makes it simple and quick for developers to create user interfaces (UI) for data visualizations, machine learning models, and other applications. The Streamlit interface is essential for offering an approachable and user-friendly platform for engaging with the model and analyzing its output in the context of research on cervical cancer prediction and repair using a Convolutional Neural Network (CNN).

Even those with no background in web programming may construct Streamlit, an intuitive interface, using a straightforward Python script. Users may interact with the CNN model, upload photos for prediction, and study findings in an easy and dynamic way with a range of widgets and controls it offers, including sliders, buttons, and input fields. Because Streamlit is reactive, it can change in real-time without requiring intricate callback procedures, giving users a responsive and dynamic experience. Plots and charts are examples of data visualization that may be included into the interface to aid users in visually understanding the model's predictions. Using Streamlit, machine learning models may be integrated with ease, displaying model predictions along with their corresponding probabilities. Users may alter the arrangement, look, and style of many aspects in the UI to suit

their own requirements. Using a web browser, streamlit apps are simply deployed, making it easier to share and disseminate the system and enabling remote use by researchers or healthcare professionals. With so much documentation and resources available from Streamlit's active community, developers can locate examples, debug bugs, and improve the interface's functionality more easily.

Cervical cancer prediction

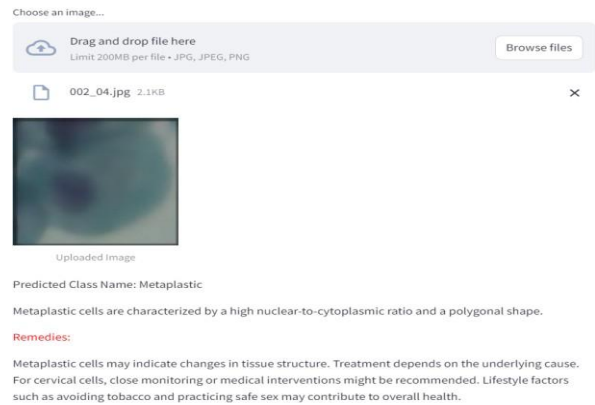


Fig.5 Cervical cancer prediction using streamlit

In addition, the Streamlit interface has interactive components that make it easier for users to utilize. It makes it possible to investigate model predictions, improving interpretability and transparency. Remedial options are also integrated into the user interface (UI), providing medical professionals with recommendations for interventions and treatment plans based on the model's predictions. This all-encompassing strategy is consistent with the goal of the research, which is to provide useful remediation procedures in a clinical setting in addition to precise prediction. The CNN model's real-time predictions and the user-friendly Streamlit interface work together to provide a comprehensive approach to cervical cancer diagnosis and treatment. The system's practicality and accessibility in clinical settings are improved by its user-friendly interface, real-time capabilities, and high accuracy rate, which guarantee dependable forecasts. Collectively, these findings highlight how the suggested strategy has the potential to have a big influence on cervical health by providing a dependable, effective, and user-friendly tool for cervical cancer early diagnosis and intervention.

VI. CONCLUSION

This research combines cutting-edge Convolutional Neural Network (CNN) algorithms with an intuitive Streamlit interface to propose a novel approach to cervical cancer remediation and prediction. A very accurate model with a 98% classification accuracy is based on the carefully selected dataset, which includes a variety of cervical cell pictures classified into classes including Dyskeratotic,

Koilocytotic, Metaplastic, Parabasal, and Superficial-Intermediate. The dataset is optimized via the Principal Component Analysis (PCA) and standardization preparation processes, which improves the CNN model's ability to capture subtle patterns linked to cervical anomalies. Healthcare professionals may engage with the model seamlessly thanks to its real-time capabilities and user-friendly Streamlit interface, which offers prompt forecasts and insights into possible corrective measures. This research represents a major advancement in using technology for early detection and intervention in cervical cancer, offering better healthcare outcomes and advancing the field of medical image analysis. Its comprehensive approach includes everything from dataset preparation to model training and user interface design.

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