

## AUTOMATED DIAGNOSIS OF RINGWORM INFECTION THROUGH A WEB APPLICATION

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### Abstract

Ringworm, a fungal infection caused by dermatophytes, is a common and contagious skin disorder that can cause significant discomfort and embarrassment. Traditional diagnosis of ringworm often involves visual inspection, microscopy and laboratory cultures, which may be time-consuming, resource-intensive, and subject to human error. This paper presents an innovative solution: an automated web application for diagnosing ringworm infections using machine learning and image processing techniques. The proposed system leverages a Convolutional Neural Network (CNN) to analyze clinical images of skin lesions, accurately identifying ringworm infections. The web-based platform is designed to be user-friendly, allowing both healthcare professionals and the general public to easily upload images for diagnosis. The model is trained on a diverse dataset of skin lesion images, using image preprocessing techniques to enhance quality and consistency. The results demonstrate high accuracy, precision and recall, indicating the potential of this approach to improve diagnostic speed and reliability. Future enhancements are discussed, including the expansion of the dataset, improvement of the model's accuracy and integration with telemedicine platforms.

**Keywords:** Machine Learning, Convolutional Neural Networks, Ringworm Infection, Skin Disease Detection.

### 1. Introduction

Ringworm, also known as tinea, is a common fungal infection that affects the skin, scalp and nails. It is caused by dermatophytes—fungi that thrive on keratin-rich tissues. The infection manifests

as itchy, red, circular patches with raised borders and it can be easily transmitted through direct contact with infected individuals, animals, or contaminated surfaces. Despite its high prevalence, the diagnosis of ringworm is still largely dependent on visual inspection, microscopy and fungal cultures. These diagnostic methods have limitations, including subjective interpretation and long processing times for culture results, which can delay appropriate treatment.

Given the high burden of dermatological diseases like ringworm and the growing need for accessible healthcare solutions, there is a strong case for developing automated diagnostic tools using technology. Machine learning, particularly Convolutional Neural Networks (CNNs), has shown great promise in automating the diagnosis of various skin conditions, such as melanoma and psoriasis, by analyzing dermatological images. However, the application of machine learning to diagnose fungal infections like ringworm is still an emerging area of research. This paper proposes the development of a web-based application that automatically diagnoses ringworm infections from skin lesion images using machine learning and image processing techniques. By automating the diagnostic process, this system aims to reduce diagnostic errors, enhance the speed of diagnosis and provide greater accessibility to dermatological care, especially in underserved areas.

## 2. Literature Review

Over the past decade, significant progress has been made in applying machine learning algorithms to the diagnosis of various dermatological conditions. Numerous studies have used image processing and machine learning techniques, such as CNNs, for skin cancer detection, particularly melanoma. These systems have demonstrated promising accuracy, often matching or surpassing the performance of dermatologists in detecting malignancies. However, the diagnosis of fungal infections, particularly ringworm, has received relatively less attention in the literature.

Studies like those by Esteva et al. (2017) and Han et al. (2018) highlight the effectiveness of deep learning in classifying skin conditions, including those that require visual identification of lesions. Most existing automated systems for dermatology focus on conditions like basal cell carcinoma, squamous cell carcinoma and melanoma, leaving fungal infections largely unexplored. Traditional methods for diagnosing fungal infections, such as direct microscopy and fungal culture, are still considered gold standards but are limited in their accessibility, speed and sensitivity. Some research has explored using machine learning for detecting dermatophyte infections from clinical images, but these methods have often relied on labor-intensive manual feature extraction or have not yet achieved the level of accuracy needed for real-world applications.

Recent work in dermatological image classification using CNNs and other advanced image processing techniques, such as segmentation and feature enhancement, shows potential for broader applications in diagnosing skin infections. However, few studies have specifically targeted ringworm infection. The absence of large-scale, labeled datasets for fungal infection diagnosis has posed a significant challenge, as many existing image datasets do not contain sufficient representations of fungal lesions. Despite these challenges, several successful case studies, such as the development of mobile applications for skin cancer detection, demonstrate the feasibility and potential impact of automated diagnosis.

### 3. Methodology

The development of the automated ringworm diagnosis system involves several key steps, from data collection and preprocessing to machine learning model training and web application deployment. This section outlines the methodology used to build the system.

#### 3.1 Data Collection and Preprocessing

The first step in the process was collecting a diverse dataset of clinical images of ringworm and other dermatological conditions. The dataset was sourced from public dermatology image repositories and dermatology clinics, ensuring that it included a wide variety of skin types, lesion types and severity levels. The dataset consisted of images taken under various lighting conditions and from patients of different age groups, genders and ethnicities. The dataset was labeled by dermatologists, with each image categorized as either ringworm (positive class) or non-ringworm (negative class).

The collected images were then subjected to preprocessing to enhance their quality and ensure consistency. This step involved resizing images to a standard size, normalizing pixel values to reduce lighting effects and applying noise reduction techniques to improve the quality of the images. Furthermore, image augmentation techniques were used, such as rotation, flipping and zooming, to artificially expand the dataset and help the model generalize better.

#### 3.2 Machine Learning Model

For the machine learning model, we selected a Convolutional Neural Network (CNN) due to its effectiveness in image classification tasks. CNNs are capable of automatically learning hierarchical features from images, making them ideal for dermatological image analysis. The model architecture consisted of several convolutional layers followed by pooling layers to reduce dimensionality, with fully connected layers at the end to output the class predictions.

The model was trained using a training dataset, with a validation dataset used to tune hyperparameters and prevent overfitting. We employed a categorical cross-entropy loss function and the Adam optimizer to train the model. The performance of the model was evaluated using metrics such as accuracy, precision, recall and F1-score, with a focus on minimizing false positives and false negatives.

#### 3.3 Web Application Development

The web application was developed using React.js for the front-end, ensuring that the interface was responsive and accessible across different devices. The back-end was built using Flask or Django, which provided a robust framework for handling image uploads, processing and serving predictions. The machine learning model, trained using TensorFlow/Keras, was integrated into the web application via a RESTful API.

Users can upload images of suspected ringworm lesions through the user interface, which then sends the image to the back-end for processing. The system preprocesses the image, runs it through the trained CNN model and returns a prediction along with the confidence score. The prediction is displayed visually, with the affected area of the lesion highlighted for easy identification.

### 4. Module Function

The web application includes several essential modules:

#### 4.1 Image Upload Module

This module allows users to upload images of skin lesions via a simple drag-and-drop interface or by selecting files from their device. The system accepts common image formats such as JPG, PNG, and TIFF.

#### 4.2 Image Preprocessing Module

Once an image is uploaded, it undergoes preprocessing to ensure that it meets the model's input requirements. This includes resizing, noise removal, color normalization to ensure consistency across the dataset.

#### 4.3 Model Prediction Module

The preprocessed image is passed through the CNN model, which classifies the lesion as either ringworm or non-ringworm. The system provides feedback in the form of a confidence score indicating the likelihood of the infection.

#### 4.4 Results Display Module

After processing the image, the results are displayed to the user with a clear indication of the predicted class (ringworm or non-ringworm) and the confidence score. If a ringworm infection is detected, the affected area is highlighted in the image.

### 5. Results

The model was evaluated on a test dataset consisting of clinical images that were not part of the training set. The system achieved an accuracy of 92%, with precision and recall values of 0.89 and 0.91, respectively. The F1-score was 0.90, indicating a balanced performance between false positives and false negatives. The web application was also tested for user experience and both healthcare professionals and lay users reported that the interface was intuitive and easy to use. In practice, the application successfully identified ringworm infections with high reliability and demonstrated its potential to streamline the diagnostic process.

### 6. Future Enhancements

While the current system demonstrates high accuracy, several enhancements are planned to further improve its performance:

1. **Larger and More Diverse Dataset:** The inclusion of more varied datasets from different regions and demographics will help improve the model's robustness and reduce biases related to skin types and lighting conditions.
2. **Improved Image Quality:** Further refinement of the image preprocessing pipeline to handle low-resolution images and lighting inconsistencies will improve the system's ability to analyze a broader range of images.
3. **Integration with Telemedicine:** The system could be integrated into telemedicine platforms, enabling healthcare professionals to remotely assess patients in areas with limited access to dermatologists.
4. **Real-Time Learning:** Incorporating user feedback and continuously updating the model with new data will help the system adapt and improve over time, ensuring that it remains accurate and reliable.

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