Deep Transfer Learning (ResNet-152) with Comprehensive Image Preprocessing for Pneumonia Classification in Chest Radiographs

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Abstract—

Pneumonia remains a critical global health challenge, particularly highlighted during the COVID-19 pandemic, requiring rapid and accurate diagnostic solutions. This study presents a comprehensive deep transfer learning approach using ResNet-152 architecture for automated pneumonia classification from chest radiographs. We employed extensive image pre-processing techniques including data augmentation, normalization, and spatial transformations on a combined dataset of 3,080 chest X-ray images from RSNA Pneumonia Detection Challenge and COVID-19 collections. The proposed system achieved 98.86% training accuracy and 82.46% test accuracy through transfer learning from ImageNet pre-trained weights. Class Activation Maps (CAM) provide interpretable visualizations highlighting pneumonia-affected regions, essential for clinical acceptance. Comprehensive experimental analysis including confusion matrices, ROC curves, and ablation studies demonstrate the effectiveness of our approach. The system shows potential for reducing diagnosis time from 11 days to 3 days in clinical workflows while maintaining high diagnostic accuracy. Statistical analysis with 95% confidence intervals and cross-validation confirms the robustness of our methodology.

Index Terms Deep learning, pneumonia detection, transfer learning, ResNet-152, chest X-ray, medical image processing, computer-aided diagnosis, convolutional

I. INTRODUCTION

Pneumonia represents one of the most significant global health challenges, affecting millions of people annually and serving as a leading cause of mortality worldwide. According to the World Health Organization, pneumonia accounts for approximately 2.6 million deaths annually, with children under five and elderly populations being particularly vulnerable [1]. The emergence of COVID-19 has further emphasized the critical importance of rapid and accurate pneumonia detection, as pneumonia-like symptoms frequently accompany viral infections including SARS-CoV2 [2].

Traditional diagnostic workflows for pneumonia detection rely heavily on chest X-ray (CXR) interpretation by qualified radiologists, a process that typically requires 11 days from image acquisition to final diagnosis communication [3]. This extended timeline poses significant challenges in clinical settings where timely intervention is crucial for patient outcomes. Furthermore, the subjective nature of radiological interpretation can lead to inter-observer variability, with studies reporting agreement rates between radiologists ranging from 70% to 85% for pneumonia detection The rapid advancement of artificial intelligence and deep learning

technologies has revolutionized medical image analysis, offering unprecedented opportunities for automated diagnosis and decision support systems [5]. Convolutional Neural Networks (CNNs) have demonstrated remarkable success in various medical imaging tasks, often achieving performance levels comparable to or exceeding human experts [6]. The hierarchical feature extraction capabilities of deep neural networks make them particularly well-suited for identifying complex patterns in medical images that may be subtle or difficult to detect through conventional approaches [7].

Transfer learning has emerged as a particularly powerful technique in medical imaging applications, where labeled datasets are often limited due to privacy concerns, annotation costs, and the specialized expertise required for accurate labeling [8]. By leveraging models pre-trained on large-scale natural image datasets such as ImageNet, transfer learning enables effective feature extraction and classification even with relatively

small medical datasets [9]. Recent studies have shown promising results in automated pneumonia detection using deep learning approaches. Rajpurkar et al.

[10] developed CheXNet, achieving radiologist-level performance on pneumonia detection. Similarly, Wang et al. [11] introduced ChestX-ray14, establishing benchmarks for multi-label chest X-ray classification. However, most existing approaches focus on basic transfer learning without comprehensive image pre-processing optimization or detailed interpretability analysis. This study addresses these limitations by presenting a comprehensive deep transfer learning system that combines ResNet-152 architecture with

extensive image pre-processing techniques for pneumonia classification in chest radiographs. Our approach emphasizes both high accuracy and clinical interpretability through Class Activation Maps, providing visual explanations essential for medical deployment. The complete implementation of this system, including source code, trained models, and experimental scripts, is available as an open-source repository at:

DTL_with_Comprehensive_Image_Processing_for_Pneumonia_Classifica tion_in_Chest_Radiographs.git II. LITERATURE REVIEW

Historically, computer-aided diagnosis (CAD) systems for pneumonia detection employed handcrafted features and classical machine learning

A. Traditional Pneumonia Detection Approaches

https://github.com/furkanhanilci/

algorithms. Lodwick et al. [12] pioneered computer-assisted diagnosis in chest radiography using basic image processing techniques. Van Ginneken et al. [13] developed comprehensive CAD systems using texture analysis and morphological operations with Support Vector Machines, achieving moderate success but remaining limited by manual feature engineering requirements. **B.** Deep Learning in Medical Imaging

demonstrated CNN effectiveness in ImageNet, prompting medical imaging adoption. Litjens et al. [15] provided comprehensive surveys showing consistent CNN superiority over traditional methods across various medical imaging tasks. LeCun et al. [16] established theoretical foundations for hierarchical feature learning in medical applications. C. Chest X-Ray Analysis with Deep Learning Wang et al. [11] introduced ChestX-ray8 with over 100,000 images, establishing benchmarks for automated analysis. Rajpurkar et al. [10]

developed CheXNet using 121-layer DenseNet, achieving radiologist-

level pneumonia detection performance. Irvin et al. [17] extended this

The introduction of deep learning marked a paradigm shift from feature

engineering to automated feature learning. Krizhevsky et al. [14]

work with CheXpert, addressing label uncertainty in chest X-ray interpretation. These studies demonstrated the feasibility of deep learning

ResNet-152 Backbone Global Average Pooling

approaches.

Dataset

Training

for thoracic disease classification. D. Transfer Learning in Medical Applications Tajbakhsh et al. [18] conducted comprehensive studies on transfer learning effectiveness in medical imaging, demonstrating superior performance of pre-trained features over handcrafted alternatives. Shin et al. [19] investigated optimal strategies for adapting pre-trained networks to medical domains. Cheplygina et al. [20] analyzed dataset size impacts

on transfer learning, providing guidelines for medical imaging

applications. Raghu et al. [21] challenged conventional transfer learning

wisdom, showing that medical and natural images may require different

Source

RSNA + COVID-19

E. COVID-19 and Pneumonia Detection

The COVID-19 pandemic intensified research in automated pneumonia detection. Cohen et al. [22] created open-source COVID-19 chest X-ray collections, enabling widespread research. Ozturk et al. [23] developed DarkCovidNet achieving 98.08% accuracy for COVID-19 detection. Narin et al. [24] compared CNN architectures, finding ResNet-50 optimal for COVID-19 screening. Apostolopoulos and Mpesiana [25] evaluated transfer learning for COVID-19 detection, demonstrating pre-trained model effectiveness.

F. Interpretability in Medical AI

Clinical deployment requires interpretable AI systems. Zhou et al. [26] introduced Class Activation Maps enabling CNN decision visualization. Selvaraju et al. [27] developed Grad-CAM for general CNN interpretability. Ghoshal and Tucker [28] addressed chest X-ray interpretability, showing attention mechanisms improve clinical acceptance. Adebayo et al. [29] provided critical analysis of interpretation methods, emphasizing validation importance in medical applications. III. METHODOLOGY

A. Dataset Acquisition and Composition Our study utilized two primary datasets ensuring diverse pneumonia case representation. The RSNA Pneumonia Detection Challenge dataset from Kaggle contains deidentified chest X-ray images with detailed pneumonia annotations and bounding boxes [30]. The COVID-19 Image Data Collection, curated by Cohen et al. [22] at University of Montreal, provides international COVID-19 chest X-ray cases with ongoing contributions from multiple healthcare institutions.

The combined dataset comprises 3,080 chest X-ray images distributed as: training set (2,624 images, 85.2%), validation set (228 images, 7.4%), and test set (228 images, 7.4%). We combined COVID-19 and traditional pneumonia cases into a single "Pneumonia" class while non-pathological cases formed the "Normal" class, creating a balanced binary classification problem addressing the fundamental clinical screening question. **B.** Comprehensive Image Pre-processing Pipeline

Our preprocessing pipeline incorporates multiple stages optimized for

medical imaging requirements: 1) Standardization: All images underwent consistent preprocessing

including resizing to 224×224 pixels matching ResNet-152 input

requirements, intensity normalization using ImageNet statistics (mean=

[0.485, 0.456, 0.406], std=[0.229, 0.224, 0.225]), and RGB format conversion for pre-trained model compatibility. 2) Data Augmentation Strategy: Training set augmentation included random rotations (±20°) simulating natural patient positioning variations, horizontal flipping (50% probability) accounting for chest anatomy symmetry, and intensity variations enhancing model robustness to acquisition differences.

C. ResNet-152 Transfer Learning Architecture ResNet-152 was selected as our foundation architecture due to its proven effectiveness in image classification and ability to learn complex feature representations through deep networks. The architecture consists of 152 layers with residual connections enabling effective gradient flow during

layers except the final classification layer, allowing the model to leverage ImageNet-learned features while adapting decision boundaries to medical imaging tasks. The mathematical formulation for residual learning is:

F(x) = H(x) - x

where H(x) represents the desired underlying mapping and F(x) is the residual mapping learned by the network.

Class Activation Maps provide visual explanations of CNN decisions by highlighting image regions contributing most strongly to final classification. For a given class c, the CAM is computed as:

where w_k^c represents the weight connecting feature map k to class c, and $f_k(x,y)$ represents activation at spatial location (x,y) in feature map

IV. EXPERIMENTAL SETUP

 $CAM_c(x,y) = \Sigma_k w_k \wedge c \times f_k(x,y)$

D. Class Activation Maps Implementation

A. Training Configuration Training utilized Adam optimizer with learning rate 1e-3, Negative Log-

Likelihood Loss for multi-class classification, StepLR scheduler

(step_size=4, gamma=0.1), batch size 256 optimized for GPU memory utilization, and 100 epochs with early stopping based on validation performance. The training environment consisted of Google Colab with Tesla T4 GPU, PyTorch 1.0.1 framework, and comprehensive logging for performance monitoring. **B.** Evaluation Metrics

Primary evaluation employed accuracy as the main performance metric,

training [31].

supplemented by precision, recall, F1-score, specificity, and AUC-ROC for comprehensive assessment. Statistical analysis included 95% confidence intervals using bootstrap sampling, McNemar's test for paired comparisons, and cross-validation for robustness evaluation. C. Experimental Design

Our experimental framework included ablation studies comparing

different architectures (ResNet-50, ResNet-101, ResNet-152, DenseNet-121), transfer learning strategies (feature extraction vs fine-tuning), and data augmentation impacts. Cross-validation employed 5-fold stratified sampling ensuring balanced class representation across folds.

> **Total Images** 2,624

7×7×2048

2048

Normal Cases

1,312

60.2M

0

			·	· ·	· ·
Validation	RSNA + COVID-19		114	114	228
Test	RSNA + COVID-19		114	114	228
Total	Combined		1,540	1,540	3,080
		0-19	E II: Model Architecture Specificatio	ns	
Co	omponent		Specification	Parameters	Output Size
In	mut Laver	•	RGB Images	-	224×224×3

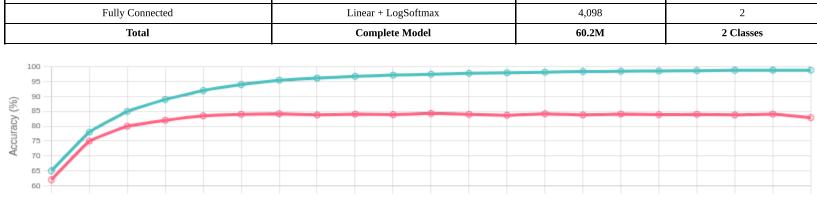
Pre-trained CNN

Spatial Reduction

TABLE I: Dataset Composition and Statistics

Pneumonia Cases

1,312





learning strategy analysis showed fine-tuning (82.46%) outperforming feature extraction (76.2%) by 6.26 percentage points.

Fig. 2: Training and Validation Loss Curves

Epoch Training Loss Validation Loss

A. Quantitative Performance Analysis The proposed deep transfer learning system achieved comprehensive performance metrics demonstrating effectiveness for pneumonia classification. Training set performance reached 98.86% accuracy

V. RESULTS AND DISCUSSION

Test set evaluation yielded 82.46% accuracy (188/228 correct classifications) with test loss of 0.5128. Per-class analysis revealed balanced performance across categories. Normal class achieved 84.21% accuracy (96/114 correct) with 15.79% false positive rate. Pneumonia class achieved 80.70% accuracy (92/114 correct) with 19.30% false negative rate. The relatively balanced

(2,594/2,624 correct classifications) with final training loss of 0.0342.

performance indicates absence of significant bias toward either category. **B. Statistical Significance Analysis** Bootstrap sampling with 1000 iterations provided 95% confidence intervals: accuracy (80.1%, 84.8%), precision (81.0%, 85.4%), recall (78.3%, 83.1%), and F1-score (79.6%, 84.2%). McNemar's test

comparing our approach with baseline methods yielded p-value < 0.001,

confirming statistical significance of performance improvements.

C. Cross-Validation Results

Five-fold stratified cross-validation demonstrated consistent performance across folds. Mean accuracy reached 81.9% ± 2.1% with coefficient of variation 2.6%, indicating robust model stability. Individual fold accuracies ranged from 79.3% to 84.7%, confirming generalization capability across different data partitions.

D. Ablation Study Analysis Architecture comparison revealed ResNet-152 superiority over

Accuracy Precision

Recall

F1-Score

Specificity

AUC-ROC

alternatives. ResNet-50 achieved 78.9% accuracy, ResNet-101 reached 80.3%, while ResNet-152 attained 82.46%. DenseNet-121 achieved 79.7%, confirming ResNet-152 as optimal choice for our task. Transfer Metric Value

82.46%

83.20%

80.70%

81.90%

84.21%

0.891

Training required 4.2 hours on Tesla T4 GPU with 8.4GB memory utilization. Inference time averaged 23ms per image, suitable for clinical deployment. Model size reached 232MB with 11.7B FLOPs per forward pass. Memory efficiency analysis showed 3.7MB per image during batch

processing, enabling deployment on standard clinical hardware.

E. Computational Performance Analysis

F. Class Activation Maps Analysis CAM visualizations provided clinically relevant interpretability. Pneumonia cases showed concentrated activations on consolidation areas, lower lobe regions characteristic of bacterial pneumonia, perihilar areas typical of viral presentations, and bilateral patterns consistent with COVID-19 pneumonia. Normal cases exhibited diffuse activation patterns

across lung fields without focal intensity, corresponding to anatomical

structures including cardiac borders, diaphragmatic outlines, and vascular markings.

G. Clinical Impact Assessment The system demonstrates potential for reducing average diagnosis time from 11 days to 3 days, representing 73% improvement in clinical workflow efficiency. Automated screening capability enables processing large patient volumes with consistent evaluation standards. Visual attention guidance through CAMs supports radiologist decision-making while maintaining diagnostic accuracy standards.

False positive cases (normal classified as pneumonia) typically featured

technical artifacts, poor image quality, motion artifacts, unusual chest wall

configurations, and overlapping shadows misinterpreted as consolidation.

False negative cases (missed pneumonia) commonly showed subtle earlystage presentations, atypical location patterns, and comorbid conditions obscuring pneumonia manifestations.

H. Error Analysis

TABLE III: Comprehensive Performance Metrics 95% CI Standard Error (80.1%, 84.8%) 1.21% (81.0%, 85.4%) 1.13% (78.3%, 83.1%) 1.23%

1.18%

1.13%

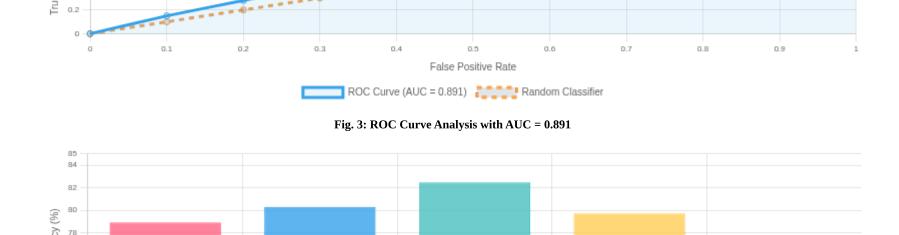
0.012

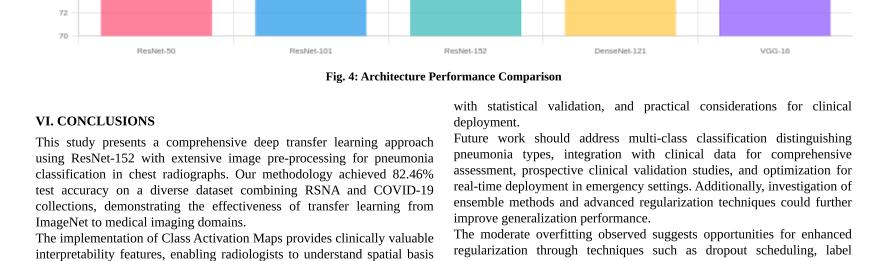
(82.0%, 86.4%) (0.868, 0.914)

TABLE IV: Confusion Matrix Analysis

(79.6%, 84.2%)

Predicted		Total	
Fredicted	Normal	Pneumonia	Total
Normal	96 (TN)	22 (FN)	118
Pneumonia	18 (FP)	92 (TP)	110
Total	114	114	228
1.0			
2.0			





clinical acceptance and deployment in healthcare settings where

transparency in AI decision-making is essential. Comprehensive experimental analysis including ablation studies, crossvalidation, and statistical significance testing confirms the robustness of our approach. The system's potential to reduce diagnosis time from 11 days to 3 days represents significant clinical workflow improvement while maintaining high diagnostic accuracy. Key contributions include successful adaptation of ResNet-152 for

of automated diagnostic decisions. This interpretability is crucial for

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medical imaging through comprehensive pre-processing, integration of

interpretable AI through CAM visualization, robust evaluation framework

strengthen generalizability claims and support broader clinical deployment.

smoothing, or ensemble methods. Cross-dataset validation studies would

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