

Comparative Analysis of Skin Disease in Different Deep Learning Models

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This study examines three CNN architectures—ConvNeXt, ResNet50, and DenseNet121—for skin lesion classification on the HAM10000 dataset. This research employs various techniques such as data augmentation, transfer learning and image segmentation to enhance each model's ability to handle variability in images. This study evaluates the three models on various parameters such as—accuracy, precision, recall, f1-score—to assess the strengths and limitations of each model. This evaluation will provide us with the model that provides the most effective approach and efficient results.

Keywords: SkinCancer, Dermoscopy, ArtificialIntelligence, Deeplearning, CNN

1. Introduction

Skin cancer is the most common type of cancer globally. The incidence rate of skin cancer has been increasing rapidly over the decades; hence the urgent need for effective prevention and detection measures. With the advancement in technologies, the survival rates have increased, however precise diagnosis remains crucial for better and optimal outcomes. Melanoma is one of the deadliest skin cancers. It is very challenging to treat this disease when diagnosed late or to limited access to dermatology specialists due to its high likelihood of metastasis. If the disease is not diagnosed early, the survival rate of an individual post-diagnosis will be a maximum of five years. However, detecting the disease at an early stage not only increases the survival rates but also offers nearly a 100% recovery probability. The slight difference across the skin lesion that can occur due to various factors such as the slight difference in the lighting, difference in skin types, age of an individual, and other environmental factors may lead to misdiagnosis despite the advancements in detection technologies. Dermoscopy has become an important tool that provides a more detailed and standardized view of skin lesions, improving the diagnostic accuracy for healthcare professionals.

The application of deep learning in medical images has been explored extensively, as highlighted by (1), Jiayi Wang (2023) in their survey on deep learning's impact on medical image analysis, which outlines the transformative role of CNNs in analyzing complex image data on skin disease classification. (2), Ismail Oztel (2023) developed a

smartphone-compatible deep learning model, emphasizing the importance of accessible and mobile solutions for diagnosis in remote or resource-limited settings. (3), Lubna Farhi (2022), applied Deep Belief Networks for dermoscopic image classification, demonstrating the flexibility and effectiveness of diverse neural architectures in distinguishing between benign and malignant lesions. (4), Cheng-Hong Yan (2021) introduced a deep hybrid CNN model specifically designed for melanoma lesion segmentation, illustrating the model's capability in enhancing segmentation accuracy for improved diagnosis. (5), Risheng Wang (2021) provided a comprehensive survey on the use of deep learning techniques, which has proven instrumental in tasks requiring precise identification of affected regions in medical images.

Artificial Intelligence (AI) has emerged as a powerful tool by providing insights on pattern recognition capabilities that support the doctors from identifying potential malignancies. Though AI algorithms have a significance in this context, clinicians still play a crucial role in validating these findings, incorporating the patient's history and lifestyle into the findings. This integration between AI and the clinical staff not only optimizes the diagnostic flow but also provides faster and data-supported insights to patients. Yet, several challenges persist in implementing effective AI-driven diagnostic tools. The need for vast, well-labelled datasets is paramount, which is one of the problems, because each cancer type has a different incidence number as well as image examples, leading to an imbalance between skin cancer classes, especially melanoma, leading to uneven data representation.

Our research addresses this problem by testing the model on a standardized skin disease dataset and evaluating the performance of each model based on its accuracy, precision, sensitivity, specificity, recall, f1-score and computation power to identify most suitable architecture for accurate and efficient skin disease classification.

2. Background Study

In (6), the study demonstrated that using wrapper-based feature selection, particularly the Grey Wolf Optimization (GWO) algorithm, yielded the highest classification performance, achieving an accuracy of 83.33% on the ISIC 2017 dataset and 93.50% on the ISIC 2018 dataset.

In (7), the study utilized a Deep Belief Learning (DBL) network architecture that achieved significant improvements in classification accuracy, especially on segmented dermoscopic images, with accuracy gains ranging from 8% to 47% over other models like AlexNet and dLeeNet. And segmented images classified with the DBL network also exhibited reduced error rates by 41.5% and faster processing times, focusing on weight distribution on the clustered regions rather than the whole image.

In (8), the study highlights that CNN-based models, such as ResNet and EfficientNet, provide robust results for skin lesion classification, with accuracy reaching up to 93.65%.

In (9), the proposed hybrid model (a combination of DeepLabV3+, MobileNetV2, EfficientNetB0, and DenseNet201) achieved an

accuracy of 94.42% and an F1score of 93.49% on the ISIC-2019 dataset, and an accuracy of 94.44% on the PH2 dataset, confirming its generalizability.

In(10),thethemodelusedareEfficientNetv2B0,Regnetx006,InceptionResnetv2.The EfficientNetv2 B0 model achieved the best overall performance across multiple classification tasks with an accuracy rate as high as 92.9% for specific binary classifications, enhancing both sensitivity and specificity.

In(11),thethemodelusedareResNet-18,EfficientNetb3,MobileNetv2.The study achieved a 74.27% classification accuracy on a seven-class skin disease dataset, including Monkeypox, using a TensorFlow Lite version of ResNet-18.

3. Methodology

The following sections provide a description of the dataset used, the techniques applied, the architecture of each model, and the structured workflow developed for training each model.

Dataset Collection: This project utilizes the HAM10000 dataset, chosen for its high-quality, dermatologist-reviewed images, which enhance the reliability of skin lesion classifications. It includes a diverse collection of 10,015 images, but only 7470 unique lesions, as duplicate images were removed to maintain data integrity and prevent biased training outcomes. The dataset consists of Actinic Keratoses (327 images), Basal cell carcinoma (514 images), Benign keratosis (1099 images), Dermatofibroma (115 images), Melanocytic nevi (6705 images), Melanoma (1113 images), and Vascular lesion (142 images), totalling to 10,015 images.

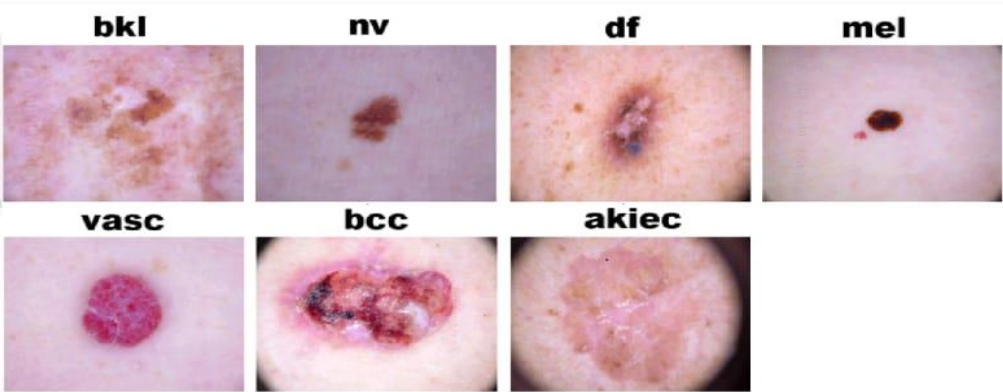


Fig.3.1.SevendistinctskinlesionsinHAM10000dataset.

- Preprocessing of Dataset:It is an important step forpreparingrawdatafordeeplearningmodels.
- Removing Duplicates: Duplicate images are removed toavoidbias.
- Resizing and Rescaling: Adjusting the dimensions of theimagesizeto224x224tofitthemodel.
- RemovingNoise:Identifyingandremovinganycorruptor irrelevant images that may affect the performance of themodel.

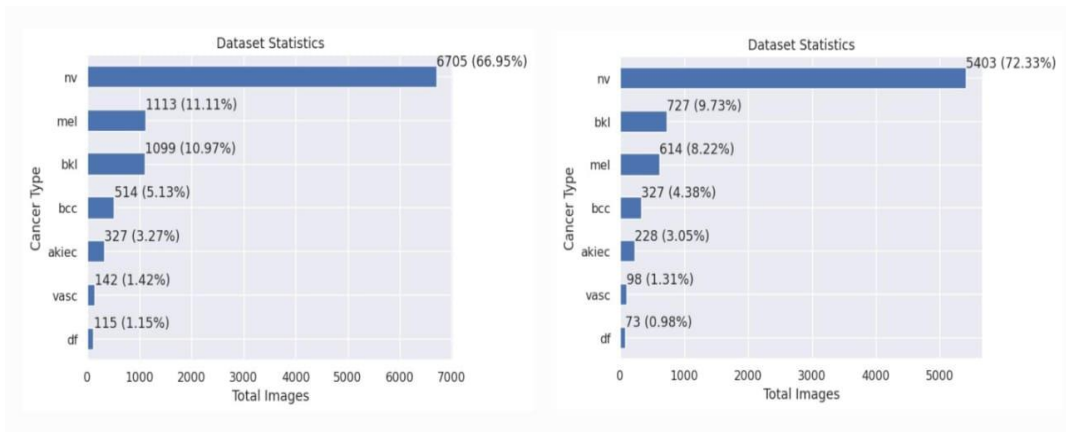


Fig.3.2.Dataset distribution classes, on the left side the graph of the raw dataset, and on the right there moved duplicate images.

Data Augmentation: It is the process of artificially expanding the size and diversity of a training dataset by applying various transformations to the images. The various data augmentation techniques used are:

Rotation: Rotating the image to specific angles.

2H-flip and V-flip: Mirroring the images horizontally and vertically.

Zooming: Randomly zooming the images and cropping the outer edges.

Shearing: Images being skewed or slanted either along the horizontal or the vertical axis.

WidthShift: Horizontal shift of the entire image along the x-axis.

HeightShift: Vertical shift of the entire image along the y-axis.

ChannelShift: Adjusting the intensity of the RGB color in the images.

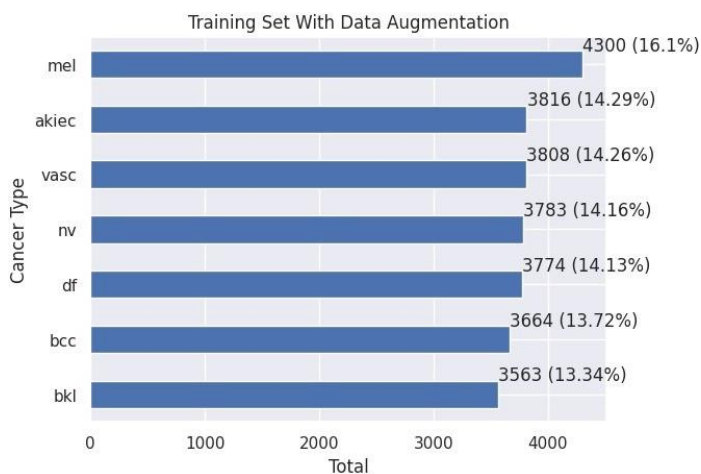


Fig.3.3.Dataset distribution after data augmentation

Transfer Learning: Transfer learning is a technique that solves related tasks using a pre-trained model. It helps in solving new problems with smaller datasets by using the knowledge of the model that has been trained on large and diverse datasets. By using these pre-trained weights, transfer learning speeds up the model training process and improves the accuracy. In this study, transfer learning is used to fine-tune the models where the original data serves as the foundation. Here, we allow the weights to be adjusted during the training process, thus ensuring that the model adapts itself to the varied representation of the medical images. This approach helps in improving the key performance metrics, providing more accurate and efficient diagnosis. The fine-tuning of the model is highly effective as it helps in capturing the disease-affected feature and enhancing the model's ability to recognize subtle differences in skin condition.

Model Architecture

This project evaluates the performance of three Convolutional Neural Network (CNN) architectures: ConvNeXt, ResNet50, and DenseNet121. The details on each of the architecture are provided below.

ConvNeXt: ConvNeXt was developed in 2020s, inspired by the design of Vision Transformer. The ConvNeXt architecture has 295 layers, out of which 36 are convolutional layers. This architecture uses approximately 89 million parameters which enhances the model's ability to capture complex patterns in images. ConvNeXt is designed for 224x224 pixel input images which makes the model more compatible with a wide range of pre-trained models. This architecture is very efficient when robust and high feature extraction is required. The model's ability to provide high accuracy like Vision Transformers and computation power and simplicity like CNN makes it an efficient model for computer vision.

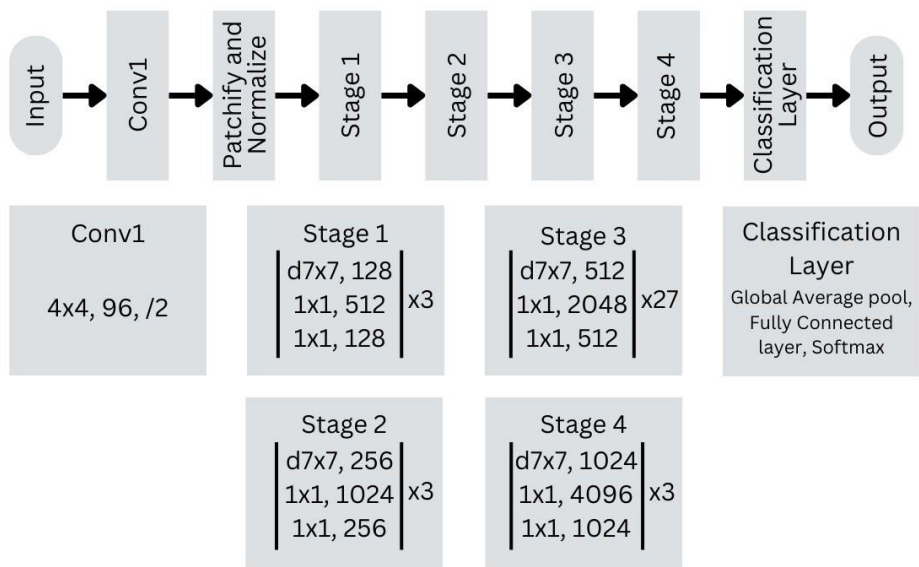


Fig.4.ConvNeXtModelArchitecture

ResNet50: ResNet50 is a variant of ResNet family. ResNet50 comprises of an initial convolutional and pooling layer followed by four stages of residual blocks, totalling to 50 layers. ResNet50 was developed to help mitigate the vanishing gradient which is a problem that occurs in deeper networks. In traditional deeper networks, as layers increase, the gradients decrease during backpropagation leading to difficulty in training. This problem is handled by the residual block of ResNet50 that incorporates shortcut connection to “skip” certain layers. This architecture uses approximately 25.5 million parameters and works efficiently with images at 224x224 pixel resolution. Its strong balance between depth and efficiency makes it an effective model for image recognition.

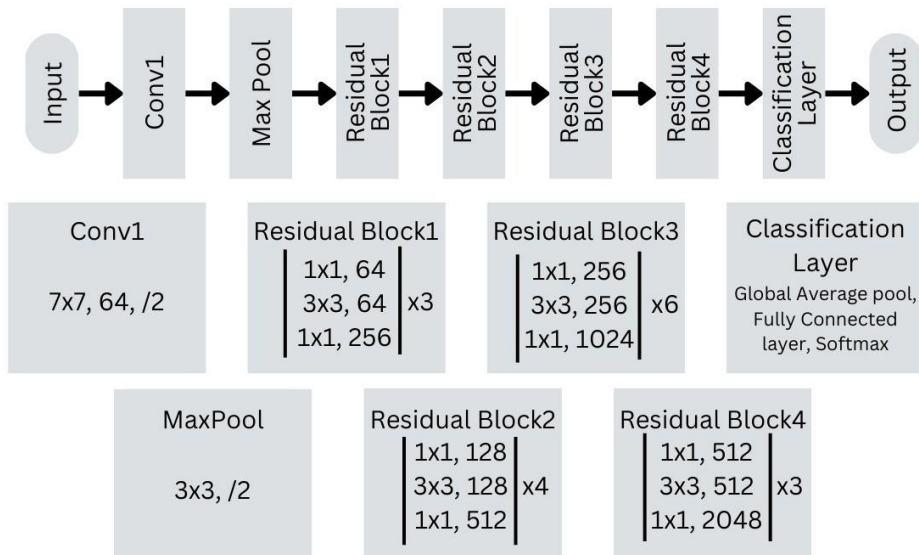
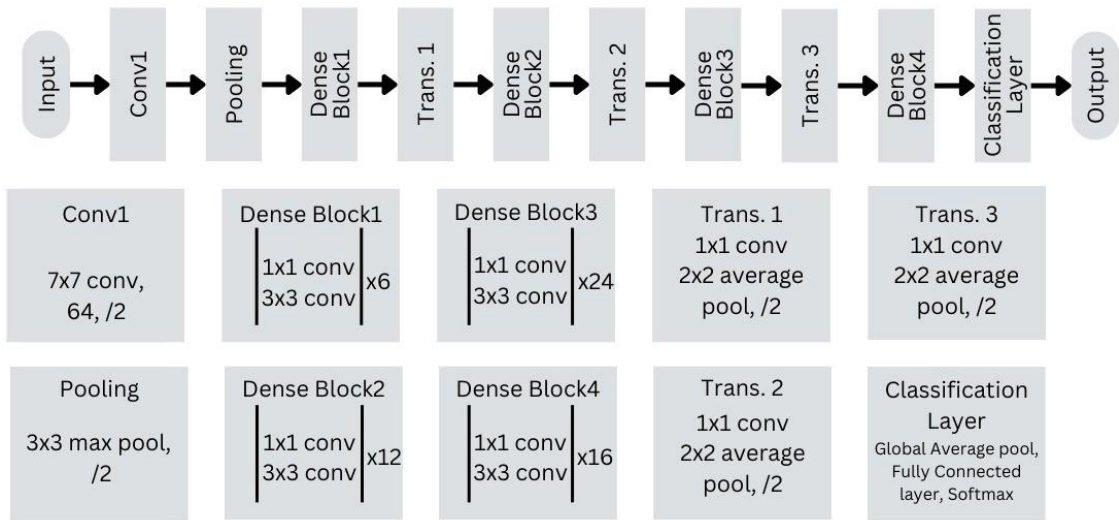


Fig.5. ResNet50 Model Architecture

DenseNet121: DenseNet121 is a variant of Dense Convolutional Network (DenseNet) architecture. DenseNet 121 is known for its innovative design where each layer connects to every subsequent layer, creating a “dense” connectivity pattern, unlike traditional CNN where each layer connects only to the previous and the next layer. DenseNet121 consists of an initial convolutional and pooling layer followed by four dense blocks and transition layers, totalling to 121 layers. This architecture uses approximately 8 million parameters and works efficiently with images of 224x224 pixel resolution. The innovative design of DenseNet121 allows better feature reuse and efficient gradient flow, thus making DenseNet121 an efficient and powerful model for image recognition tasks.



Note that each “conv” layer shown in the table corresponds to the sequence BN-ReLU-Conv

Fig.6.DenseNet121ModelArchitecture

Training:

A configurable notebook is created for this project, to centralize the main settings for each training model. Each model was tested with the dataset imported via Kaggle’s Dataset, using only the TPU to train the model, resulting sub-30 minutes of training time to conclude a single execution of the notebook. In this stage, the images are divided into three sets: training, validation, and testing with the respective rates: 70%, 10%, and 20%. The exact amount of images for each class is represented in the Fig.7:

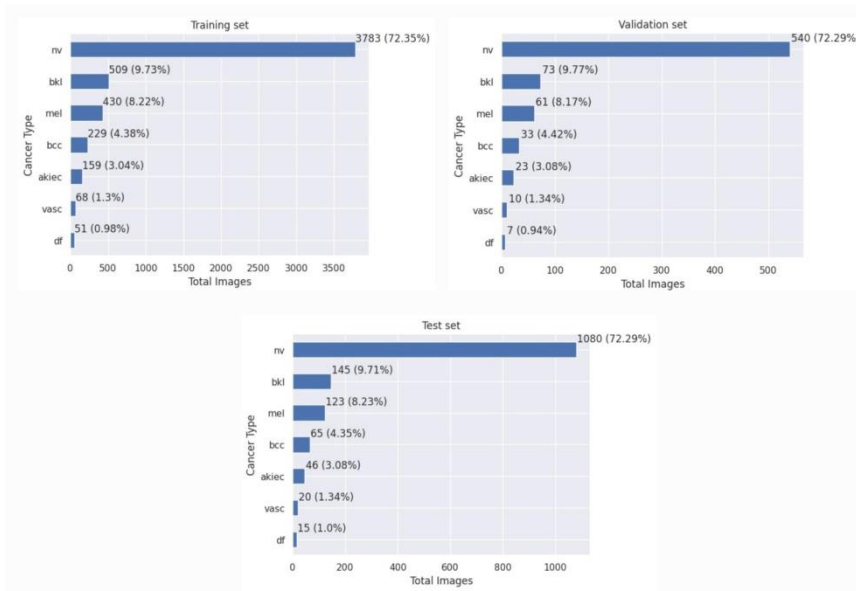


Fig.7.Dataset distribution between training, validation, and test sets

4. Results:

This study evaluates the performance of—ConvNeXt, ResNet50 and DenseNet121—on HAM1000 dataset using key metrics—accuracy, precision, recall, f1-score and specificity—after training on augmented and segmented images. All the three models had a variety of combination and the overall performance metrics are displayed in the Fig.

9. Only those combinations are taken that achieved an accuracy of more than 50%. The study revealed that ConvNext achieved the highest accuracy of 96.45% demonstrating robust feature extraction capabilities suitable for complex medical images, followed by ResNet50 with an accuracy of 96.03% with its residual block effectively vanishing the gradient problem, and DenseNet121 achieved an accuracy of 88.62% excelling in its innovative design allowing better feature reuse and efficient gradient flow. In conclusion, ConvNeXt was found to be the most suitable architecture for handling variability and complexities in skin lesions, making it the top-performing model in comparative analysis.

Models	Accuracy (%)	Precision (%)	Recall (%)	F1_score (%)	Specificity (%)	ROC_AUC (%)
convnextall	96.39	97.66	97.66	97.66	99.61	98.88
convnextbalancednormtransf	78.94	91.42	91.28	91.34	98.57	96.15
convnextdataaugmentationtransf	96.19	98.02	98.02	98.02	99.67	98.94
convnextdataaugmentationtransfbalanced	96.45	97.99	97.99	97.99	99.66	98.93
convnexttransf	89.27	95.57	95.57	95.57	99.26	97.85
convnexttransfbal	85.45	92.51	92.51	92.51	98.75	96.31
convnexttransfbalseg	85.78	92.35	92.35	92.35	98.72	95.89
densenet121all	80.16	81.61	81.61	81.61	96.93	90.28
densenet121balancednormtransf	74.18	84.84	84.71	84.77	97.47	92.61
densenet121dataaugmentation	79.15	81.58	81.55	81.56	96.93	90.24
densenet121dataaugmentationtrans	78.77	81.7	81.7	81.7	96.95	90.25
densenet121dataaugmentationtransfbalanced	79.76	80.77	80.74	80.75	96.79	89.97
densenet121dataaugtransfbalminmax	56.31	91.34	91.31	91.32	98.55	96.62
densenet121trans	88.62	86.24	86.24	86.24	97.7	92.64
densenet121transbal	84.08	68.5	68.5	68.5	94.75	82
densenet121transbalseg	84.71	71.41	71.41	71.41	95.23	83.81
resnet50all	95.57	96.77	96.77	96.77	99.46	98.41
resnet50dataaugmentationtransf	96.03	97.04	97.04	97.04	99.5	98.44
resnet50dataaugtransfbal	95.64	96.62	96.62	96.62	99.43	98.33
resnet50transf	94.71	97.09	97.09	97.09	99.51	98.66
resnet50transfbal	90.76	92.35	92.35	92.35	98.72	96.15
resnet50transfbalseg	90.43	90.06	90.06	90.06	98.34	94.38

Fig. 8. Performance metrics for different combinations of the three models.

5. Conclusion and Future Work

In summary, each model has their unique strengths in skin disease classification with ConvNeXt having the highest accuracy and leading in terms of overall performance. ConvNeXt is a very efficient model where high accuracy and speed are required. While DenseNet121 and ResNet50 also performed well, their architectures are more suited for environments where memory and computational limitations are less important. To improve further accuracy, future research can investigate on model ensembles that incorporates ConvNeXt, DenseNet121 and Nanotechnology Perceptions Vol. 20 No. S13 (2024)

ResNet50. It may also involve adding hybrid models that increase the sensitivity to subtle image features. Further broadening the study to a wide range of skin disease and more diverse dataset can increase the efficiency of the model.

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