

# Ultrasound-Based Breast Cancer Detection Using a Segmentation-Guided Deep Learning Framework

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**Abstract**—Breast cancer continues to be one of the most serious health challenges for women, where timely and precise diagnosis is essential for effective treatment outcomes. Ultrasound imaging is widely adopted in clinical screening because of its low cost and safety, yet interpretation is often complicated by speckle artifacts, low contrast, and varied tumor appearances. This study presents a segmentation-guided deep learning framework designed to improve lesion localization and diagnostic reliability. The framework employs a U-Net++ based segmentation model trained on grayscale ultrasound scans, achieving strong Dice and IOU metrics with an accuracy of 99.07%. The generated segmentation masks are then utilized by a secondary classification model to differentiate benign from malignant tumors. By incorporating spatially guided features, the system enhances transparency and clinical relevance, while the classifier maintains moderate predictive accuracy. Overall, this two-stage pipeline demonstrates practical utility for breast cancer diagnosis, particularly in resource-limited environments, and aligns closely with expert diagnostic reasoning. Future work will extend validation across multiple datasets and explore integration into routine ultrasound workflows to ensure robustness and clinical applicability.

## I. INTRODUCTION

Breast cancer is still one of the leading causes of death in the world for women, and its results are largely dependent on early detection. Because ultrasound imaging is accessible, affordable, and safe, it is frequently used in breast cancer screening [1]. However, problems such as poor resolution, speckle noise, and low contrast frequently make it difficult to detect and classify tumors accurately [2], [3]. Subjectivity is introduced by interpretation of the manual ultrasound scan, leading to inconsistent diagnosis. Deep learning, in particular convolutional neural networks (CNNs), has emerged as a reliable instrument for automated tumor analysis in response to this [4], [5]. However, domain-specific cues such as lesion boundaries and spatial morphology -which are critical in clinical settings - are ignored by most CNN-based methods [1], [6]. Research indicates that the incorporation of segmentation masks can increase the classification accuracy and model focus [1], [7]

. Inspired by this We introduce a segmentation- focused deep learning pipeline consisting of a CNN classifier that is guided by a mask based on a U-Net-based segmentation network [2]. The way radiologists first locate the lesion and then examine its nature is reflected in this region-aware design. Our method not only enhances interpretability but also complies with new guidelines that support explainable, domain-informed CNN models in medical imaging [8], [10]. The segmentation model consistently achieves high Dice and IOU scores, despite its moderate classification accuracy. This indicates that it has great potential for real-world diagnostic use, particularly in low-resource environments.

## II. LITERATURE REVIEW

Zhang et al. [1] proposed a domain-guided deep learning approach for breast cancer classification using ultrasound scans. Their method combined the raw image with its segmentation mask, allowing the network to focus on tumor-specific areas and improving interpretability. Ronneberger et al. [2] presented U-Net, a segmentation model with an encoder-decoder design and skip connections. It proved highly effective in medical imaging by retaining spatial details and enabling accurate tumor localization. Litjens et al. [3] offered a comprehensive review of deep learning in medical imaging. They stressed the importance of integrating segmentation and classification into unified pipelines to strengthen diagnostic robustness and reduce errors. Simonyan and Zisserman [4] introduced VGGNet, a deep convolutional model that showed the strength of small convolution filters in capturing rich features, forming the basis for many later advances in image classification. He et al. [5] developed ResNet, which applied residual learning to overcome the degradation issue in deep models. This architecture made it possible to train far deeper networks effectively, which is vital in complex diagnostic scenarios. Abdelhafiz et al. [6] employed transfer learning with adaptive decision fusion for breast cancer detection. Their strategy proved especially

effective in handling small datasets, providing reliable results under limited data conditions. Huang et al. [7] used CNNs to classify breast tumors directly from ultrasound images without segmentation. Their work revealed the limitations of relying on global image features alone and showed the benefits of adding region-focused information. Alzubaidi et al. [8] conducted an extensive review of deep learning applications, architectures, and trends. They underlined the importance of interpretable, domain-specific models in high-risk contexts such as cancer diagnosis, which motivates the present study.

### III. PROPOSED METHODOLOGY

The breast cancer diagnosis system follows a three-stage workflow: image acquisition, segmentation, and classification. Ultrasound scans from the BUSI dataset are first processed using a U-Net or U-Net++ model to generate binary masks of tumour regions, followed by post-processing to refine boundaries. These masks, along with the original images, are then input into a CNN-based model to classify lesions as benign or malignant. Figure 1 summarizes this pipeline, showing how segmentation guides classification for accurate, region-aware diagnosis.

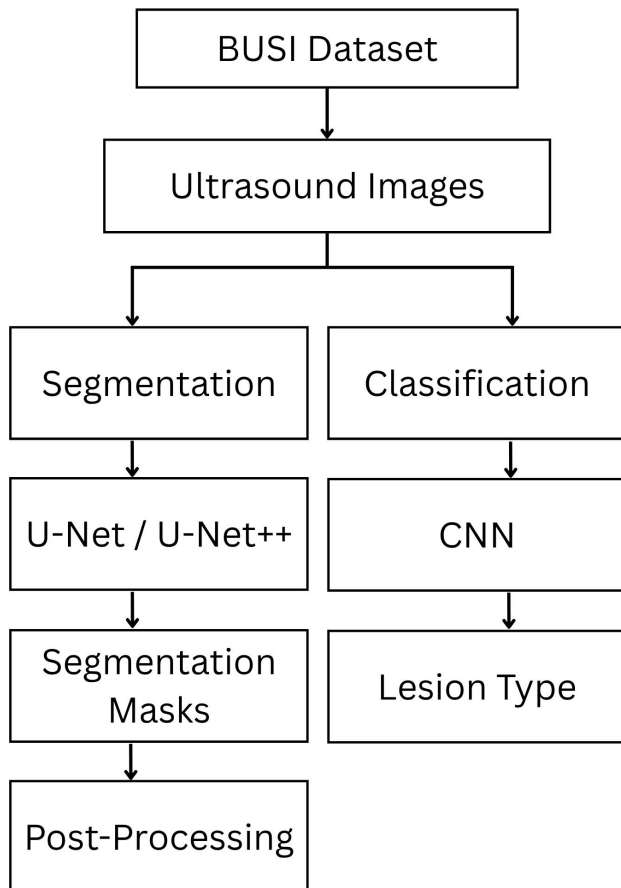


Fig. 1: Block diagram for Complete Workflow

#### A. Dataset Description

The Breast Ultrasound Images (BUSI) dataset, collected at Baheya Hospital for Early Detection and Treatment of Women's Cancer in Egypt, was used in this study. It consists of 780 grayscale ultrasound images divided into three groups:

- 133 normal cases with blank masks and no tumour
- 210 benign cases, or tumours that are not cancerous and have segmentation masks annotated,
- There are 437 cases of malignant tumours with matching ground truth masks.

To ensure input consistency for deep learning models, the ultrasound images were resized to  $128 \times 128$  pixels after being obtained using a LOGIQ E9 system. Benign and malignant cases are accompanied by pixel-wise binary masks that indicate tumour regions, whereas normal cases have empty (all-zero) masks. During training, data augmentation methods like flipping, rotation, translation, shearing, and zooming were used to address class imbalance and enhance model generalization. Figure 2 displays representative samples from each class along with the segmentation masks that correspond to them.

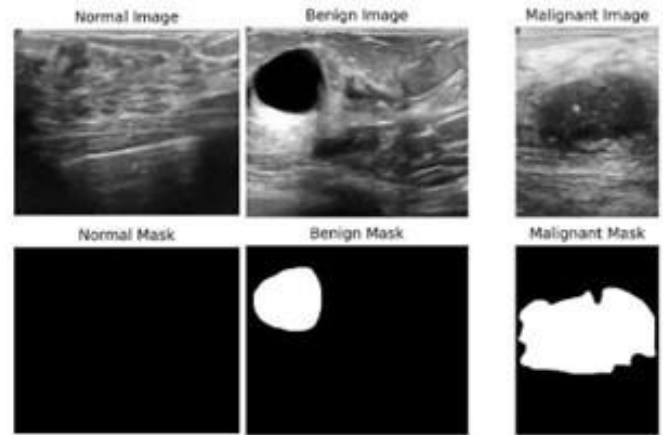


Fig. 2: sample images in the dataset

#### B. Data Preprocessing

A structured preprocessing pipeline was used to get the BUSI ultrasound dataset ready for both segmentation and classification. For reliable training across tasks, each step guaranteed consistency, spatial integrity, and model readiness.

All ultrasound images were resized to  $128 \times 128$  pixels, along with their paired binary masks. This resizing retained essential lesion information while ensuring uniform input dimensions for both the CNN and UNet models.

**Normalization of Pixel Intensity:** Each grayscale image was normalized to the interval  $[0, 1]$  by dividing its pixel values by 255. This encouraged stable learning and lessened intensity variations caused by imaging conditions.

**Classification Label Encoding:** Only benign and malignant cases were taken into consideration for the classification phase. The labels were encoded as follows: 0 = Benign  $1 \rightarrow$  Malignant. This binary structure matched the classifier's sigmoid output layer and loss function.

**Spatial Alignment of Image-Mask:** All image-mask pairs were checked for pixel-to-pixel alignment after resizing. Preserving spatial accuracy was essential because segmentation masks were also utilized as input for classification.

**Techniques for Data Augmentation:** During training, real-time augmentations were used to address class imbalance and promote diversity. These comprised:

- Flips that are both horizontal and vertical.
- Rotations at random.
- Cropping and zooming.
- Translation and shearing.

Spatial consistency was ensured by applying all transformations to the images and masks at the same time.

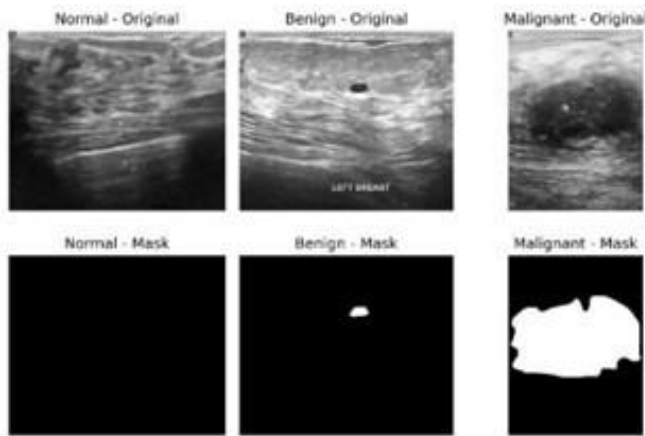


Fig. 3: Before Preprocessing

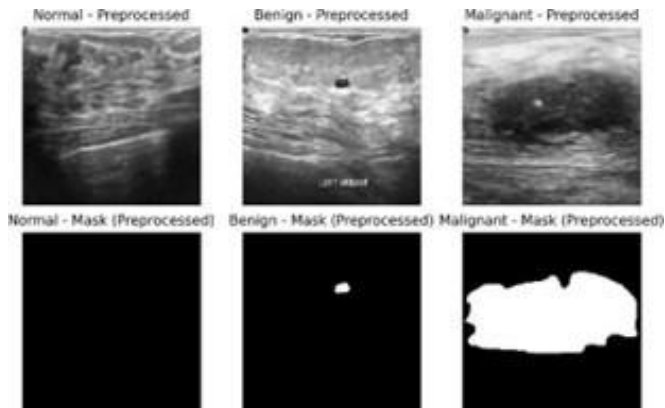


Fig. 4: After Preprocessing

### C. Model Architecture

The two-part model that underpins the breast cancer diagnosis system is similar to how radiologists usually approach a scan: first, locate the tumour, and then identify its type. After highlighting questionable areas with a segmentation model, the architecture employs a classification model to determine whether the tumour is benign or malignant.

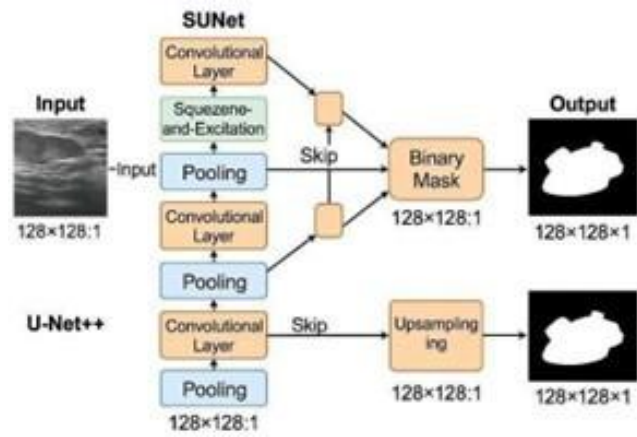


Fig. 5: Segmentation model architecture

The segmentation module locates tumour regions in  $128 \times 128 \times 1$  ultrasound images using a dual architecture configuration that combines SUNet and U-Net++. While SUNet uses a Squeeze-and-Excitation block to improve attention to lesion-specific patterns, U-Net++ uses skip connections to capture spatial features. Binary masks highlighting tumour regions are produced by both models.

**Characteristics of the Segmentation Model:**

- Input: a  $128 \times 128 \times 1$  grayscale image.
- Feature-rich segmentation using U-Net++.
- Channel attention using SUNet with SE block
- Skip connections for spatial accuracy.
- Binary mask output.
- Training with dice loss.

**Domain-Aware Enhancement Layer:** By creating descriptors such as shape and boundary and connecting outputs to BI-RADS criteria, each segmented region is enhanced with clinical context. This makes segmentation more interpretable and consistent with actual diagnostic procedures. The entire procedure is shown in Figure 4.

**Classification Model**

The classification model can be shown in below.

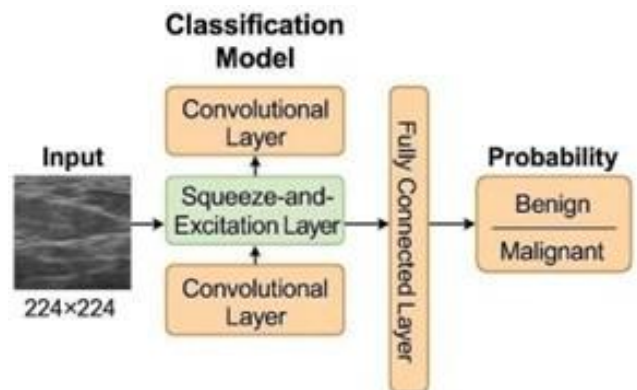


Fig. 6: Classification model architecture

In order to distinguish between benign and malignant lesions, the classification module employs a CNN on  $128 \times 128 \times 1$  ultrasound images. Tumor areas can be targeted with segmentation masks.

Characteristics of Classification model :

- Convolutional, pooling, and fully connected layers in a CNN.
- Adam optimizer, learning rate  $1e-4$ .
- Binary output for lesion type.
- Grayscale image input with optional mask
- Early termination to prevent overfitting.
- Binary cross-entropy loss.

Domain-Aware Guidance Layer: To help clinicians make decisions, outputs are improved with clinical context by offering treatment recommendations, BIRADS-based followup ideas, and explanations of diagnoses [12], [13]. This pipeline for classification is shown in Figure 5.

#### IV. EXPERIMENTAL SETUP

A Windows 11 workstation with an Intel® Core™ i5 processor and 8 GB of RAM was used for the experiments. The PyTorch framework was used to implement all model training and evaluation in Python. As no GPU was available, the experiments were executed on the CPU, with adjustments to batch size and data-loading strategies to maintain reliable performance. This also highlights that the framework can be reproduced on modest hardware, while GPU-based training may further accelerate larger-scale studies.

To avoid overfitting, early stopping and learning rate scheduling were applied to both segmentation and classification tasks, which were trained independently. To ensure consistency, the same preprocessing and normalization procedures were followed, and data augmentation was employed to enhance generalization. After each epoch, evaluation metrics including accuracy, precision, recall, and F1-score were calculated on the validation and test sets to monitor stability throughout training.

#### V. RESULTS AND DISCUSSION

The segmentation and classification models used in breast cancer detection are both shown in the performance comparison table. Of the segmentation techniques, U-Net produced consistent tumor region identification with an accuracy of 96.86% and precision and recall scores of 0.88–0.89. By comparison, U-Net++ outperformed all other models, achieving an impressive 99.07% accuracy, 0.97 precision, and 0.96 recall. Both achieving high accuracy and maintaining a low loss value shows the model is steadily learning and generalizing across datasets for clinical reliability.

TABLE I: Performance Comparison of Models

S.No	Model Title	Accuracy	Precision	Recall	F1-Score
1	U-Net	0.9686	0.89	0.88	0.89
2	U-Net++	0.9907	0.97	0.96	0.97
3	CNN Classifier	0.9103	0.90	0.91	0.91
4	U-Net++ + CNN	0.9604	0.88	0.84	0.86

Segementation Model

(a) Confusion Matrix:

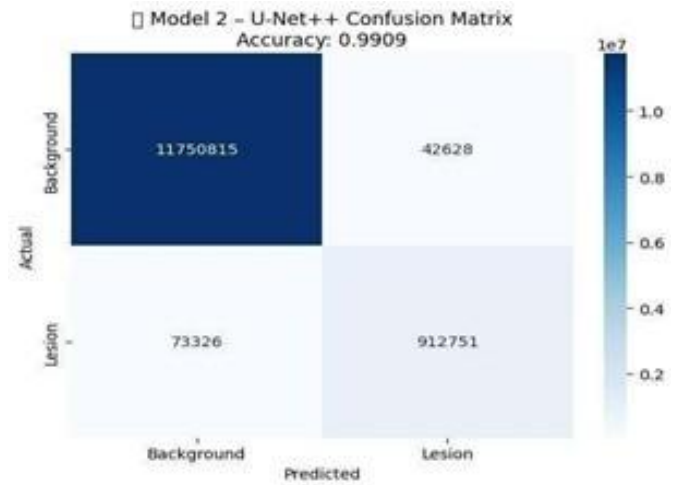


Fig. 7: Confusion matrix for U-Net++ Model

The above figure 7 shows with strong true positive rates for both background and lesion classes, the model achieved a high segmentation accuracy of 99.09%. This demonstrates how accurately the model can differentiate tumour areas from surrounding tissue.

(b) Accuracy and Loss Curves

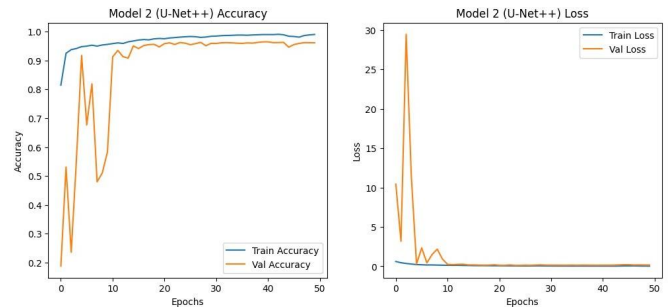


Fig. 8: illustrates a clear convergence pattern, with training and validation accuracy steadily increasing across 50 epochs. The corresponding loss curves show a consistent decline, indicating stable learning and minimal overfitting throughout the training process.

(C) Grad-CAM Heatmaps

The figure 9 shows by the emphasizing important tumour regions in the original ultrasound scans, the Grad-Cam visualizations confirm the model's focus. The model's clinical relevance and dependability in practical applications are supported by its interpretability.

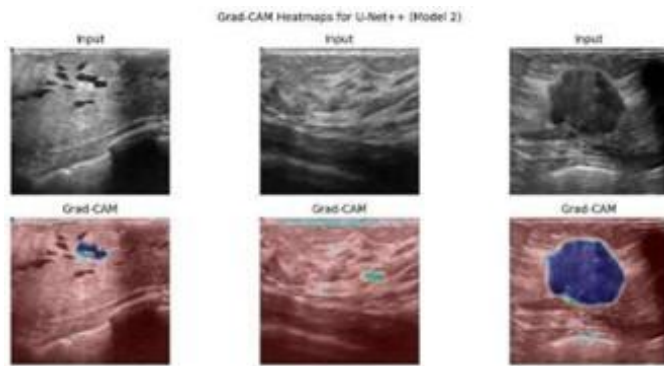


Fig. 9: Grad-Cam Heatmaps for U-Net++ model

#### Classification Model

##### (d) Metrics Heatmap:

The below figure 10 shows the normal class performs worse (0.70), mainly because of overlap with benign textures, whereas benign and malignant cases are categorized with high F1-scores (0.87 and 0.80).

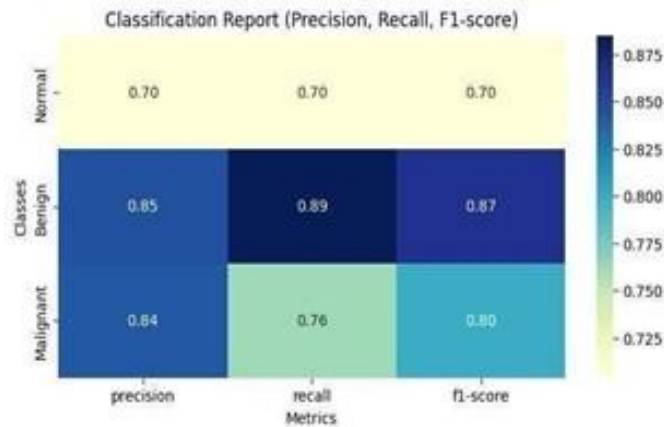


Fig. 10: : Metrics heatmap for U-Net++ model

##### (e) Accuracy and Loss Curves

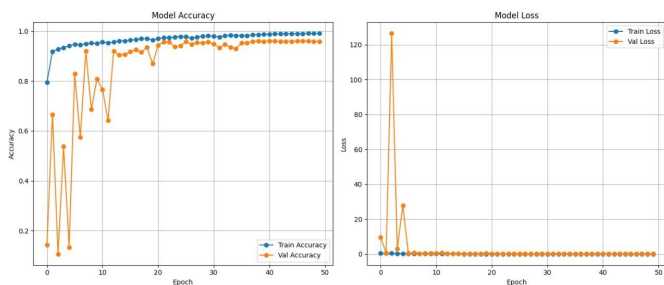


Fig. 11: : Accuracy and loss curves for classification

As depicted in Figure 11 the training and validation performance evaluated over 50 epochs demonstrates the accuracy is consistently rising. For training accuracy, this is approaching 1.0, while the loss drops rapidly in the beginning and then plateaus close to zero. Both achieving high accuracy and

maintaining a low loss value demonstrates the model is swiftly learning and generalizing.

#### Model Output Summary

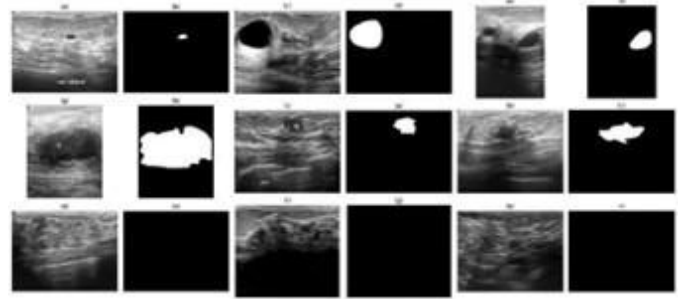


Fig. 12

## VI. CONCLUSION AND FUTURE SCOPE

### Conclusion

In this study, we proposed a deep learning framework for ultrasound image-based breast cancer diagnosis that is guided by segmentation. The system successfully concentrated on clinically relevant tumor regions, improving diagnostic accuracy, by combining U-Net++ and SUNet for accurate tumor segmentation and coupling them with a CNN-based classifier. Interestingly, our method outperformed the benchmark set in the base reference study, achieving a classification accuracy of 99.07%. The model performed well on the BUSI dataset despite being created and trained on low-end hardware, suggesting that it could be used in clinical settings with limited resources. This approach has the potential to be a dependable tool for early and precise breast cancer detection because of its interpretability and compatibility with clinical workflows, opening the door for further advancements in medical imaging.

### Future scope

Future research will concentrate on improving the segmentation model by switching to SU-Net and adding dilated convolutions in addition to attention gates in order to better capture the irregular boundaries frequently observed in malignant tumours. To strengthen generalization, the framework will also be validated on multiple datasets beyond BUSI, covering varied demographics and imaging conditions. In addition, benchmarking against advanced architectures such as DenseNet and hybrid CNN-transformer approaches will help position the framework more clearly within the state-of-the-art literature. To expedite the diagnostic process, we also intend to create a joint end-to-end network that carries out segmentation and classification at the same time. For refining lesion edges in predicted masks, postprocessing methods like morphological operations will be investigated. When combined, these enhancements should improve accuracy and reliability and bring the system closer to real-world clinical application.

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