Final Project: Deep Learning for Breast Cancer Detection

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20 April 2025

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Abstract

- One-paragraph summary of the problem, dataset, methodology, and main findings.
- One-paragraph summary of the problem, dataset, approach, and key results.

1. Introduction

1.1 Background & Motivation

- Describe the clinical and societal significance of early breast cancer detection.
- Mention the NHS 2025 initiative and how AI fits into screening.

According to recent research[1], neural networks outperform …

1.2 Objectives

- Apply deep learning (CNNs) to mammography image classification.
- Evaluate performance vs. traditional methods/radiologists.

1.3 Scope

Briefly note focus on classification (benign vs malignant), dataset used, and evaluation metrics.

Background & Motivation

- Significance of early breast cancer detection.
- NHS 2025 initiative on DL for screening.

Project Objectives

- Build and evaluate CNN models using the DDSM/CBIS-DDSM dataset.
- Assess whether CNNs can match or exceed radiologist performance.

Scope

- Focus on classification (benign vs. malignant), with optional segmentation.
- Use curated public data for transparency and reproducibility.

2. Dataset

2.1 Dataset Description

- Dataset: CBIS-DDSM
- Number of cases: 753 calcifications, 891 masses
- Modalities: Mammograms with labels and ROI masks

2.2 Preprocessing

- Resizing, normalization, augmentation
- ROI extraction (if applied)

2.3 Splitting Strategy

- Training, validation, and test set proportions
- Use of predefined splits if applicable

Dataset Description

- Use of CBIS-DDSM (Lee et al., 2017) — curated version of DDSM.
- Number of images, classes (benign/malignant), calcifications vs. masses.

Preprocessing Steps

- ROI extraction, resizing, normalization.
- Augmentation techniques (flipping, rotation, etc.).

Train/Validation/Test Split

- Based on BI-RADS or predefined splits from the dataset.

3. Deep Learning Workflow

3.1 Problem Definition

Define input/output: - Input: X-ray mammogram or ROI - Output: Binary label (benign or malignant)

Define the supervised classification task: - Input: X-ray mammogram image (ROI or full view) - Output: Binary label (benign/malignant)

3.2 Data Preparation

- Preprocessing steps
  - Denoising, rescaling, grayscale conversion
  - Normalization [e.g., pixel range 0–1 or mean/std]
- Label encoding
- Data augmentation techniques: flips, rotations, zooms

3.3 Model Building

- Baseline model: custom CNN
- Advanced models:
  - Transfer learning (e.g., VGG16, ResNet50)
  - Optional segmentation with U-Net

3.4 Model Training

- Loss function: Binary Crossentropy
- Optimizer: Adam
- Metrics: Accuracy, AUC, Sensitivity, Specificity, F1-score
- Epochs, batch size, learning rate, early stopping, callbacks (e.g., early stopping, LR scheduler)

3.5 Evaluation

- Report performance on test set
  - Confusion matrix
  - ROC curve, AUC
  - Precision, Recall, F1-score

3.6 Model Improvement

- Regularization techniques: dropout, L2
- Data augmentation experiments
- Architecture tuning: more layers, batch norm

- Transfer learning comparisons
- Add dropout / L2 regularization
- Increase network depth
- Apply transfer learning
- Tune hyperparameters

4. Results

- Performance Tables: Accuracy, AUC, Sensitivity, Specificity per model
- Visualizations: ROC curve, training/validation loss curves
- Error Analysis: Misclassified cases, confusion matrix
- Example visualizations of predictions (e.g., Grad-CAM)

5. Discussion

- Compare results with literature benchmarks
  - Comparison with Radiologists (Wang 2024)
- Strengths and limitations of the model/approach
- Interpretability & practical deployment considerations
  - Grad-CAM (optional)

6. Conclusion

- Summary of findings
- Implications for clinical use: Whether deep learning improves screening performance
- Suggestions for future work: Recommendations for future research (ensemble models, multi-task learning)

7. References

- Wang L. (2024). *Frontiers in Oncology*
- Lee et al. (2017). *Scientific Data*
- Chollet, F. (2018). *Deep Learning with Python*
- TensorFlow/Keras documentation

Appendix

A. Code Snippets

- Add code snippets here later

B. Hyperparameter Table

- Add hyperparameter tables

C. Full Model Architecture

- Add full model architecture

D. Data Statistics

- Add any dataset distribution histograms or BI-RADS label breakdowns

E. Report Writing Tools

The writing process for this report was conducted using **Quarto**, a modern scientific and technical publishing system that integrates **Markdown**, **LaTeX**, and executable code within a single framework. The project uses the `manuscript` type configuration to generate both **PDF (via XeLaTeX)** and **HTML outputs** with consistent styling, numbered sections, and title-cased tables of contents. The directory follows a modular structure (`_quarto.yml`, `report.qmd`), with customizations for fonts, TOC titles, and citation formatting via `.bib` and `.csl` files. **Version control** was managed using **Git and GitHub**, enabling reproducible and collaborative manuscript development. Integrated with **VSCode** and **Zotero (via Better BibTeX)**, this setup provides a complete academic writing workflow—featuring live previews, citation support, and source-controlled outputs—crucial for high-quality, reproducible scientific communication.

References

1. Lee, R. S. *et al.* A curated mammography data set for use in computer-aided detection and diagnosis research. *Scientific Data* **4**, 170177 (2017).