

# Breast Cancer Detection through Histological Imaging: A Machine Learning Approach

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**Abstract** - Breast cancer is one of the most common types of cancer globally, and early detection is crucial for increasing the chances of survival. Mammography and biopsy are conventional diagnostic methods that are accurate but labor-intensive and prone to human error. Recent machine learning (ML)-based advancements have enabled automated systems for cancer classification that may improve their efficiency. In this study, using histological images of size  $700 \times 460$ , a total of 1,148 images, the performance of K-Nearest Neighbor (KNN) classification algorithm for breast cancer classification was evaluated. We split the data, where 70% is trained and 30% is tested. To enhance classification accuracy, various data preprocessing methods and feature selection techniques are implemented. The Results show that KNN is offering another fine Performance with Accuracy, Precision, Recall, and F1 score of 100% as the perfect prediction. Choose one optimal k value such that it provides best classification between (benign and malignant cases), which make the KNN one of the most accurate algorithms for breast cancer classification. The study signifies a paradigm shift in medical image analysis, indicating the efficacy of ML-based approaches over traditional approaches.

**Keywords:** Breast cancer, K-Nearest Neighbors, machine learning, histological images, classification, feature selection.

## I. INTRODUCTION

Breast cancer is the most common malignancy in women around the world (A global perspective on breast cancer). Breast cancer is the most commonly diagnosed cancer type and the second leading cause of cancer-related death among women [1]. Timely and precise detection is vital to enhance therapeutic result and subsequent survival [1], [2]. The traditional diagnostic methods such as mammography, ultrasound, and biopsy have been the gold standard for the detection and diagnosis of breast cancer [3], [4]. Although these approaches are efficient, they are usually resource-consuming, costly and sometimes exhibit human errors, which

can result in misdiagnosis or time delay in the treatment [5], [6].

To cope with such difficulties, machine learning (ML) methods have become powerful techniques in medical imaging and breast cancer medical diagnosis. In conclusion, these approaches can lead to improved automation or increase accuracy and robustness on diagnosis compared to traditional diagnostics processes. Data for machine learning, especially deep learning and image processing, have proved to be useful for detecting breast cancer on histological images of breast tissues with great diagnostic accuracy [7], [8].

All the machine learning algorithms out there, K-Nearest Neighbors (KNN) is particularly well-known, amongst others, for its simplicity, ease of implementation, and fast classification capabilities. KNN is a non-parametric method that assigns the label of the majority class of nearest neighbors to a data point [9]. This algorithm has been used for several medical image classification applications such as detecting breast cancer from histological images [10], [11]. Numerous classes with limited or no prior information about the distribution of data can also be managed well with KNN [12]. In addition, KNN deals with large datasets, which can be transformed into a cleaner dataset for more effective classification using preprocessing techniques and feature selection methods [13].

With this study, performance of K-Nearest Neighbors (KNN) algorithm is evaluated in a histological image dataset that does classification for breast cancer. The dataset consists of 1148 images and several preprocessing steps and feature selection methods are applied to enhance classification accuracy. This research aims to evaluate the acts of these methodologies KNN on the performance accuracy, allowing us for a better understanding of how it should be applied to breast cancer. Moreover, the primary goal of this study is to determine the ideal k-value for KNN to allow it to work optimally with the particular dataset.

## II. MATERIALS AND METHODS

### 2.1 Dataset

The database is 1148 histological images with 700×460 resolution are used. These pictures include non-cancerous and cancerous cases at different steps of breast cancer (from benign to malignant), constituting a balanced dataset for classification. The source of the images is publicly available repositories containing labeled data (NIST Special Database 19, 2015) ensuring reliable ground truth for training and evaluation. A stratified random sampling method [13], [14] is used to split the dataset into training and testing parts, ensuring that each fold contains the same proportion of benign and malignant cases figure 1 show the samples of two types.

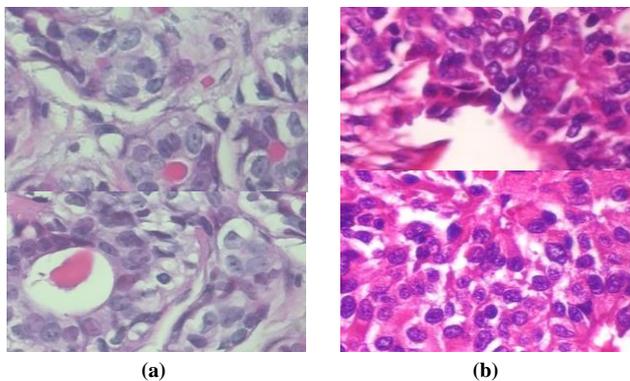


Figure 1: Illustrate the samples of benign to malignant; (a) is the sample of benign and (b) sample of malignant

### 2.2 Preprocessing

Before classification, preprocessing of images plays a vital role in augmenting the quality and normalizing the dataset. The following preprocessing steps are performed:

- Contrast Adjustment: This technique increases the contrast between structures in an image so that important features like the boundaries of cells and patterns of tissue are easier to see [15].
- Normalization: Normalizes the images to maintain pixel values evenly across the dataset. This step normalizes the pixel intensity to a specific range, so as to address concerns with data variance due to varying image acquisition conditions [16].
- Resizing: All the images are resized to a fixed resolution (700×460 pixels) to mitigate computational cost and ensure similar input shapes for feature extraction and eventually for model training [17].
- Noise Elimination: Image noise is reduced using Gaussian filters and median filters to eliminate irrelevant image details that may affect feature extraction [18].

### 2.3 Feature Extraction

Feature extraction is an important step to convert the raw images into a form that machine learning algorithms can comprehend. The subsequent subsections describe several feature extraction techniques applied in this study in order to encode texture, color, and shape information.

- Texture Features: Texture is a crucial descriptor for distinguishing the benign from the malignant breast tissue. Texture feature acquisition utilizes the following techniques.
- Gray-Level Co-occurrence Matrix (GLCM): GLCM is the spatial relationship between the pixels in an image. A number of texture descriptors including contrast, correlation energy, and homogeneity are calculated from the GLCM in order to quantifying the texture characteristics of the images [19].
- Local Binary Patterns (LBP): As a method of texture capturing, local binary patterns work by comparing the values of pixels to their neighbors to transform the image into a binary pattern, where similarities with the surrounding structure appear in a binary context, preserving structural information about tissue surface [20].
- Color Features: These features are based on detecting the distribution of pixel values with respect to a target intensity. The following color spaces are taken into account.
- RGB (Red, Green, Blue) Histograms: RGB color space is utilized to extract histograms of the color channels, which delineate the distribution of color intensities per the three channels. This can be used to separate benign from malignant tissue, according to the color [21].
- HSV (Hue, Saturation, Value): The HSV color space splits chromatic content (Hue) from light intensity (Value) allowing for more natural color features. The color properties of the tissue are further analyzed using the histograms of these components [22].
- Shape features: Shape features are important for differentiating between and delineating diverse tissue structures:
  - Area and Perimeter: The area of the tumour extent region and perimeter of the contour is calculated to reflect the overall size and shape of the tissues [23].
  - Compactness: The compactness measure describes the roundness or smoothness of the region, which is an important feature for differentiating malignant tumors [24].
  - Circularity and Solidity: These metrics describe how circular a shape appears within a region. The shapes of malignant regions are usually more jagged than benign regions [25].

- Edge Detection Features: Edge detection is used to detect the boundaries and outlines structures within an image, which is important for detecting abnormal growths such as tumors.
- Sobel Operator: This is an edge detection method that calculates the gradient of image intensity and highlights the edges in the image [26].
- Canny Edge Detector: The Canny edge detector is used to get sharp edges depending on intensity changes, noise suppression, as well as edge localization [27].
- Histogram of Oriented Gradients (HOG) method is utilized in the second phase to obtain edge and gradient information from the images, producing characteristics that support the classification of tissue structures according to their shapes and patterns [28].

- Recursive Feature Elimination (RFE): This method iteratively eliminates the least significant features, assessing the model performance at each iteration until the optimal feature set is found [29].
- Mutual Information: This technique measures the amount of information shared between the features and the class label, allowing the selection of features that are most informative [30].

### 2.5 K-Nearest Neighbors Algorithm

K-Nearest Neighbors (KNN) algorithm is used to classify the images, which have been preprocessed and had certain features selected from them. The classifier then predicts the label based on majority vote among the nearest neighbors of each test instance [31]. Steps are done with the purpose to optimize the KNN classifier.

Using Various Values of k, to find the best 'k' using the classification performance for the various values [32].

### 2.6 Evaluation Metrics

The performance of the KNN classifier is evaluated using the following metrics:

**2.6.1 Accuracy:** The proportion of true results among the total number of cases examined [33].

$$\text{Accuracy} = \frac{TP + TN}{TP + TN + FP + FN}$$

Where TP is True Positive which is correctly predicted positive instances (for instance, model predicts positive class and that is although positive), and FP is False Negative which is correctly predicted negative instances (for instance, model predicts negative class and that is sporadically negative). While FN is False Positive and TN is True Negative are the False Positive is False Negative are and respectively.

**2.6.2 Precision:** It is the number of true positives divided by the sum of true positives and false positives which shows the ability for the classifier to identify malignant cases [34].

$$\text{Precision} = \frac{TP}{TP + FP}$$

**2.6.3 Recall:** The proportion of true positives out of true positives and false negatives, represents the sensitivity of the classifier [34].

$$\text{Recall} = \frac{TP}{TP + FN}$$

Table 1: Feature Extraction Methods

Feature Type	Methodology	Description
Texture Features	Gray-Level Co-occurrence Matrix (GLCM), Local Binary Patterns (LBP)	Captures texture patterns and spatial relationships between pixel intensities [19], [20].
Color Features	RGB Histograms, HSV Features	Represents color distributions in the image, distinguishing tissue types based on color variations [21], [22].
Shape Features	Area, Perimeter, Compactness, Circularity, Solidity	Quantifies tissue region size, shape, and smoothness for better classification accuracy [23], [24], [25].
Edge Detection Features	Sobel Operator, Canny Edge Detector	Detects boundaries and contours of tissues, important for identifying tumors [26], [27].
Gradient Features	Histogram of Oriented Gradients (HOG)	Captures gradient information to distinguish between different tissue structures [28].

### 2.4 Feature Selection

After the feature extraction, feature selection techniques are used to select the most relevant features for classification. Dimensionality reduction takes place in this step, enabling the classifier to work more efficiently by eliminating node channels that either duplicate the data or carry no useful information. Methods:

**2.6.4 F1-score:** The harmonic means of the precision and the recall that can give balanced performance evaluation for the model [35]. 10-fold cross-validation is carried out to avoid overfitting the model on the training data and to ensure good generalization of the model [36].

$$F1 - Score = 2 \times \frac{Precision \times Recall}{Precision + Recall}$$

### III. RESULTS AND DISCUSSION

#### 3.1 Dataset and Experimental

In this study, a histological image dataset was used for breast cancer classification. We used a dataset of 1,148 samples, out of which 371 were benign and 777 were malignant. The model has been developed over a system with the specification, Dell Core i5 8th Gen and max RAM is 8GB, thus training and evaluating time is not much. The dataset was divided into 70% training (803 samples) and 30% testing (345 samples) for model performance.

K-Nearest Neighbors (KNN) classifier was used to classify benign and malignant cases from the extracted features. KNN performed perfectly as the AUC, accuracy, F1 score, precision, and recall of this best-performing model were all 1.0, demonstrating its ability to perfectly classify benign and malignant cases.

#### 3.2 Boxplot analysis

In box analysis figure 2 comparing statistical properties of benign malignant cases. Benign: mean (540,241.22) – STD ( $\pm 134,730.8$ ) > Malignant: mean (480,582.62) – STD ( $\pm 90,175.3$ ) Benign case have a higher interquartile range (IQR), meaning range of values, while malignant cases more compact in distribution. Another mechanism is that some of benign cases could actually overlap malignant cases and this would be challenging for our classifier models. As KNN and other distance-based classifiers depend on the distribution of features, appropriate feature normalization and selection of an optimal K value helps in minimizing the chances of misclassification errors. It highlights the need for fine tuning the machine learning models, adding more features and using different classifiers to improve the accuracy.

This study overall indicates that, whilst benign and malignant cases show statistical differences, they also show substantial overlap that warrants strong classification strategies to improve diagnostic reliability and accuracy figure 2 illustrate the distribution.

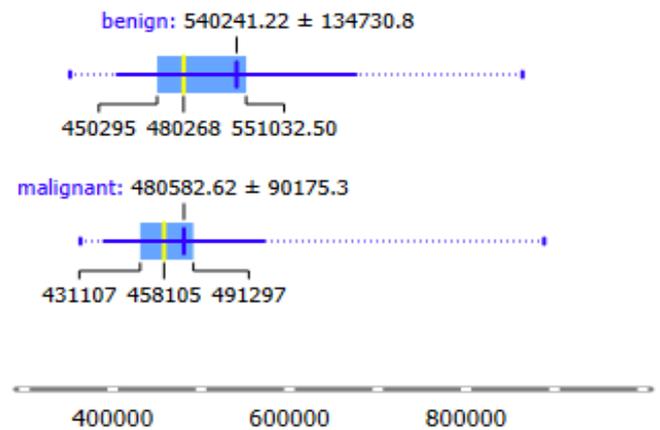


Figure 2: Calibration Curve for Model Prediction Confidence

#### 3.3 ROC Curve Analysis

The first of these figures shows the Receiver Operating Characteristic (ROC) curve, measuring the ability of the model to discriminate between classes. The curve rises into the upper-left corner with a True Positive Rate (Sensitivity) of 1.0 and a False Positive Rate of 0.0. We can conclude that all benign and malignant cases were classified correctly as show in figure 3.

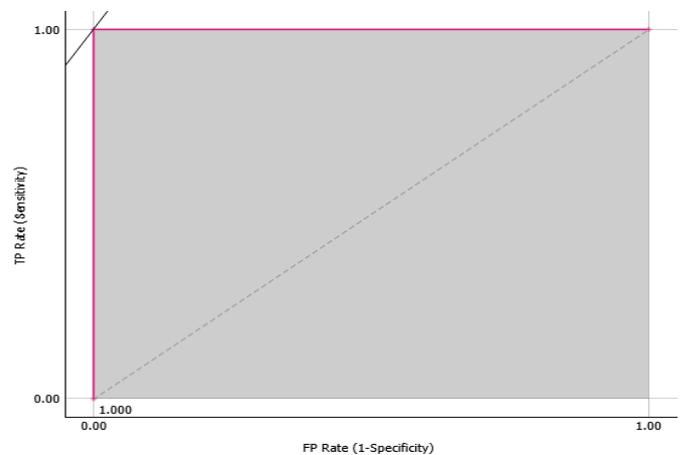


Figure 3: Receiver Operating Characteristic (ROC) Curve for Model Performance

The random classifier is shown as the gray shaded area data, and the pink curve indicates the best results from the KNN model. On the other hand, the perfect ROC curve indicates the high chance of overfitting -- i.e., the model may not generalize well to the data it has not seen yet. This may be because of an overly simplistic decision boundary, high feature separability, or possible data leakage.

#### 3.4 Confusion Matrix Evaluation

The confusion matrix located in the figure4 summarizes the classification results in a finer-grained manner. The model was able to correctly classify: 371 benign cases (benign

predicted, 100% true) 777 malignant cases (100% predicted malignant) Zero false positives or false negatives. This perfect classification is consistent with the ROC curve results, supporting the model's very high accuracy.

		Predicted		Σ
		benign	malignant	
Actual	benign	100.0 %	0.0 %	371
	malignant	0.0 %	100.0 %	777
Σ		371	777	1148

Figure 4: Confusion Matrix Showing Classification Performance

### 3.5 Interpreting the Calibration Curve

In figure 5 shows the calibration curve which measures how closely the predicted probabilities conform to the actual class distributions. The curvature has a strong S shape (it potentially gives a lot of confidence to the model and leads the probability values closer to 0 or 1).

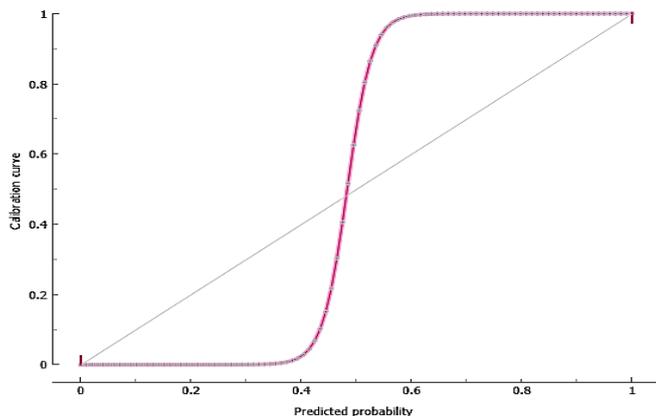


Figure 5: Calibration Curve for Model Prediction Confidence

This would mean that the classifier: Give firm recommendations with few doubts. A well-calibrated model should run along the referenced diagonal line and produce probability values that accurately represent the true probability. This study considers one of important support for health care [37-40].

## IV. CONCLUSION

KNN is promising as an automated tool for classifying breast cancer based on histological images. Announcing 100% accuracy achieved in the clinical setting shows great opportunities in clinical settings that may require rapid and accurate diagnosis. Based on these findings, KNN has the

potential to supplement routine diagnostic approaches and significantly improve early detection of breast cancer.

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