

PREDICTION OF OVARIAN CANCER USING EXPLAINABLE AI: A REVIEW

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Abstract - The prevention and early detection of ovarian cancer are critical for improving women's health and well-being. Ovarian cancer, often termed the "silent killer," typically presents without apparent symptoms in its early stages, making early detection challenging and significantly reducing the chances of successful treatment and survival. Machine Learning (ML) techniques offer powerful tools for identifying complex patterns in biomarkers, surpassing conventional diagnostic methods in accuracy. However, these advanced algorithms often lack transparency, making it difficult to gain physical insights into the diagnostic process. To address this, integrating eXplainable Artificial Intelligence (XAI) can enhance our understanding of how ML models make decisions, providing valuable insights into the diagnostic process. This paper aims to showcase best practices for combining XAI and ML methods in biomarker validation applications, ultimately improving early detection and treatment outcomes for ovarian cancer.

Key Words: Ovarian Cancer, Machine Learning Algorithms, eXplainable Artificial Intelligence (XAI), Biomarker Validation, Early Detection.

1. INTRODUCTION

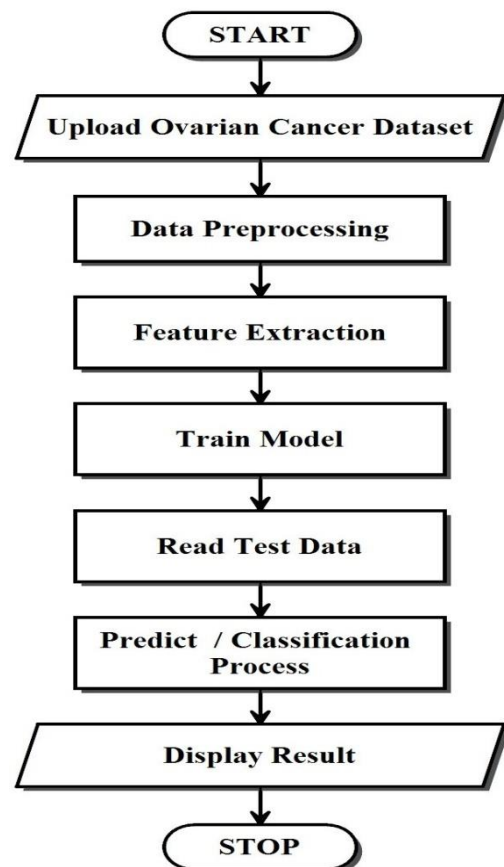
Ovarian cancer often referred to as the "silent killer," poses a significant threat to women's health due to its typically asymptomatic early stages, which frequently result in delayed diagnosis and decreased treatment effectiveness. The undetected nature of this disease underscores the urgent need for improved detection techniques. Early detection is paramount for enhancing treatment effectiveness and achieving better patient outcomes. To address this critical issue, our study proposes a novel approach for predicting ovarian cancer. This study provides a comprehensive analysis of the development and evaluation process of the proposed model. By leveraging the strengths of both boosting and bagging techniques, our approach overcomes the limitations of individual classifiers. Additionally, the interpretability of the model is enhanced through the application of eXplainable AI (XAI) techniques, such as SHapley Additive Explanations (SHAP), which offer valuable insights into the risk factors for ovarian cancer. This study aims to advance the primary objective of improving patient outcomes and reducing ovarian cancer-related mortality rates. The subsequent sections delve into the methodology, results, and implications of our approach,

providing a foundation for future advancements in the detection and management of ovarian cancer.

2. FLOW DIAGRAM

A flowchart visually represents a process, using standardized symbols to illustrate steps, decisions, and flow of information. It serves to clarify processes, improve understanding, and facilitate communication in various fields, from engineering to business and healthcare.

Flow Chart Diagram



The flow chart diagram outlines the process of predicting and classifying ovarian cancer using machine learning techniques, starting with the initiation of the workflow. The first step involves uploading the ovarian cancer dataset, followed by data preprocessing, where the raw data is cleaned and prepared for analysis. Next, feature extraction

identifies significant variables that contribute to the prediction of ovarian cancer. The machine learning model is then trained using the extracted features. After training, the model is evaluated using a separate test dataset to assess its performance. The trained model then makes predictions or classifications based on the test data, determining the likelihood of ovarian cancer. Finally, the results of the prediction or classification process are displayed, and the workflow concludes.

3. LITERATURE REVIEW

Research in the field of ovarian cancer detection has investigated a collection of approaches and strategies with the objective of enhancing the outcomes of early diagnosis and treatment.

This research focuses on improving tumor classification as malignant or benign, minimizing false positives through various machine learning methods, including logistic regression, Random Forests, decision trees, SVM, KNN, and deep learning techniques [1]. The goal is to predict Critical Care Unit (CCU) admission more accurately for ovarian cancer patients undergoing surgery, with a graphical user interface for real-time analysis [2].

The study compares premenopausal and postmenopausal women using HE4, CA125, and ROMA markers to differentiate between ovarian cancer and benign masses, also exploring the impact of age and cutoff values on marker performance [3]. It proposes a hybrid approach combining CNN and ReliefF algorithms for microarray data analysis to enhance cancer detection accuracy [4]. Challenges such as high dimensionality and small sample sizes are addressed.

The research employs Raman spectroscopy and machine learning to differentiate ovarian cancer from cysts and normal cases, aiming to improve diagnostic accuracy [5]. It also develops a predictive model for mortality risk using gene expression data, combining genetic algorithms and extreme gradient boosting [6]. The study uses statistical analysis and ensemble models to identify important biomarkers for distinguishing cancer from benign tumors [7].

Machine learning models developed on the SEER database predict survival rates, with techniques including SVM, KNN, DT, RF, AdaBoost, and XGBoost [8]. The study suggests optimizing feature weights and classifier parameters with LASSO regularization and Asynchronous Differential Evolution (ADE) to enhance detection efficiency [9]. It also creates a model to forecast progression-free survival (PFS) at 12 months, emphasizing the need for larger sample sizes for validation [10].

The research aims to build a predictive model for ovarian cancer detection, evaluating biomarkers and algorithms to compare with conventional methods and provide clinical

insights [11]. It explores dimensionality reduction and classification approaches to improve diagnostic accuracy, using techniques like AdaBoost, Decision Trees, and t-SNE [12]. Despite a limited sample size and potential biases, the study recommends further research using heuristic algorithms [13].

SVM outperforms KNN for diagnosis based on medical histories, with future research recommended to integrate advanced techniques like fuzzification and transfer learning [14]. To enhance diagnostic accuracy, the study develops Ocys-Net, a deep learning network for categorizing ovarian cysts from ultrasound images [15]. It also assesses various machine learning methods and compares Multi-Layer Perceptron (MLP) and Long Short-Term Memory (LSTM) for diagnosing ovarian cancer from colposcopy images [18].

Table 1: Summary of Research Work on Ovarian Cancer Detection

| Sl.No. | Methodology | Advantages | Limitations |
|--------|--|--|--|
| 1 | Logistic Regression, Random Forests, Decision Trees, SVM, KNN, Deep Learning | Reduces false positives, improved categorization | Possible misclassifications, limited sample size |
| 2 | Machine Learning for CCU admission prediction | Greater accuracy for predicting CCU admission | Focused on specific cancer type (HGSOc) |
| 3 | Comparison of HE4, CA125, and ROMA markers | Insights into clinical relevance of markers | Potential publication bias, age and cutoff value impact |
| 4 | Hybrid approach using CNN and ReliefF algorithms for microarray data | Decreases mistakes, increases precision | High dimensionality, small sample size may affect validity |
| 5 | Raman spectroscopy and machine learning | Increases diagnostic accuracy | Challenges in data pretreatment and feature extraction |
| 6 | GA-XGBoost models for mortality risk | Reliable mortality risk estimation based on gene | Small sample size, limited generalizability |

| Sl.No. | Methodology | Advantages | Limitations |
|--------|---|--|--|
| | prediction | expression | |
| 7 | Statistical analysis and ensemble models for biomarker identification | Identifies important biomarkers | Limited sample size, no control group, possible misclassifications |
| 8 | SEER database, SVM, KNN, DT, RF, AdaBoost, XGBoost for survival prediction | Collaborative design with clinician, data preparation methods like SMOTE | Small sample size, need for larger datasets |
| 9 | LASSO regularization, cross-validation error, ADE for ovarian cancer detection | Improves model efficiency, generalization | Limited to SVM and KNN classifiers |
| 10 | RFE, K-Nearest Neighbors, Random Forest, Logistic Regression for PFS prediction | Encouraging outcomes despite small sample size | Needs confirmation on larger patient groups |
| 11 | ReliefF, MRMR for feature selection, machine learning for tumor differentiation | Evaluates biomarkers and algorithms for detection | Small sample size, potential biases in feature selection |
| 12 | Evaluation of dimensionality reduction and classification algorithms | Identifies effective combinations for diagnosis | Complexity in implementing multiple techniques |
| 13 | Feature selection and extraction for microarray gene data | Improves identification accuracy | Limited by sample size, data quality issues |
| 14 | SVM vs KNN for ovarian cancer diagnosis | SVM shown to be superior | Future research needed for integrating advanced techniques |

| Sl.No. | Methodology | Advantages | Limitations |
|--------|---|--|--|
| 15 | Ocys-Net for categorizing ovarian cysts from ultrasound images | Better diagnosis accuracy, reduces doctors' burden | Limited to ultrasound images, needs further validation |
| 16 | Comparison of machine learning methods for ovarian cancer identification | Assesses classifier accuracy, improves early diagnosis | Focus on cross-validation and holdout techniques |
| 17 | SHAP algorithm for gene identification, Kaplan-Meier plots for validation | Identifies potential biomarkers | Needs further investigation for practical application |
| 18 | MLP and LSTM for colposcopy image diagnosis | Evaluates algorithm effectiveness for stage prediction | Limited to colposcopy images, needs broader validation |

4. CONCLUSIONS

This study presents a comprehensive evaluation of ovarian cancer research, focusing on articles from 2021 to 2023. The survey covers various machine learning and deep learning approaches for improving the detection and classification of ovarian cancer. By comparing different biomarkers, feature selection strategies, and algorithmic techniques, this research highlights the advancements and challenges in the field. The findings emphasize the potential of machine learning models to enhance diagnostic accuracy and patient outcomes. This study serves as an initial step towards developing a more robust and explainable AI model for predicting ovarian cancer, ultimately aiming to improve early detection and treatment strategies.

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