

## **An Empirical Study of Machine Learning Algorithms for Cancer Identification**

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### **ABSTRACT:**

A major challenge in the search for a cure for cancer is predicting the illness status of the disease. For instance, distinguishing between benign and malignant tumours helps doctors diagnose cancer more accurately. Although technology advancements produced data on patients with various illness stages, it would be crucial to assess how well machine learning algorithms accomplish predictions. In this article, we suggest employing machine learning algorithms such as a variation of AdaBoost, deepboost, xgboost, and support vector machines. We then analyse them using area under curve and accuracy on actual clinical data linked to thyroid cancer, colon cancer, and liver cancer. Results from experiments demonstrate the SVM's strong performance.

*Keywords:* Boosting, Machine Learning, Cancer Genomics, Medical Diagnosis

### **1.INTRODUCTION:**

By 2020, there may be 15 million more cancer cases [1]. A crucial step in the medical diagnosis of cancer patients would be to give healthcare workers efficient tools to spot cancer situations. Researchers from several fields created computational methods to help with medical diagnostics in order to accomplish this objective. For instance, to enhance the pre-surgical diagnoses of thyroid nodules, Stokowy et al. [2]

suggested a computational strategy based on gene expression data to distinguish malignant from benign thyroid nodules. Zhang *et al.* [3] proposed using extreme learning machine (ELM) applied to several microarray data including lung data, lymphoma data and other prediction performance. As various machine learning researchers have recently proposed computational methods to improve the performance, there is a need to apply these recent machine learning algorithms for clinical data generated via various technologies.

In this paper, we present using Deep Boost, which is a new ensemble learning algorithm [7]; XGBoost, which is a scalable end-to-end tree boosting system [8]; a variant of Adaboost [9]; and support vector machines (SVM) [10, 11]. We apply these machine learning algorithms for cancer identification. Experimental results on real data pertaining to thyroid cancer, colon cancer, and liver cancer show that SVM outperforms the previously mentioned algorithms.

The rest of the paper is organized as follows. Section II reports experimental results of machine learning algorithms on real data pertaining to thyroid cancer, colon cancer and liver cancer. Section III concludes the paper and points out future work.

## 2. PROPOSED SYSTEM:

The proposed system in this paper, we present using Deep Boost, which is a new ensemble learning algorithm XGBoost, which is a scalable end-to-end tree boosting system, a variant of Adaboost and support vector machines (SVM). We apply these machine learning algorithms for cancer identification. Experimental results on real data pertaining to thyroid cancer, colon cancer, and liver cancer show that SVM outperforms the previously mentioned algorithms.

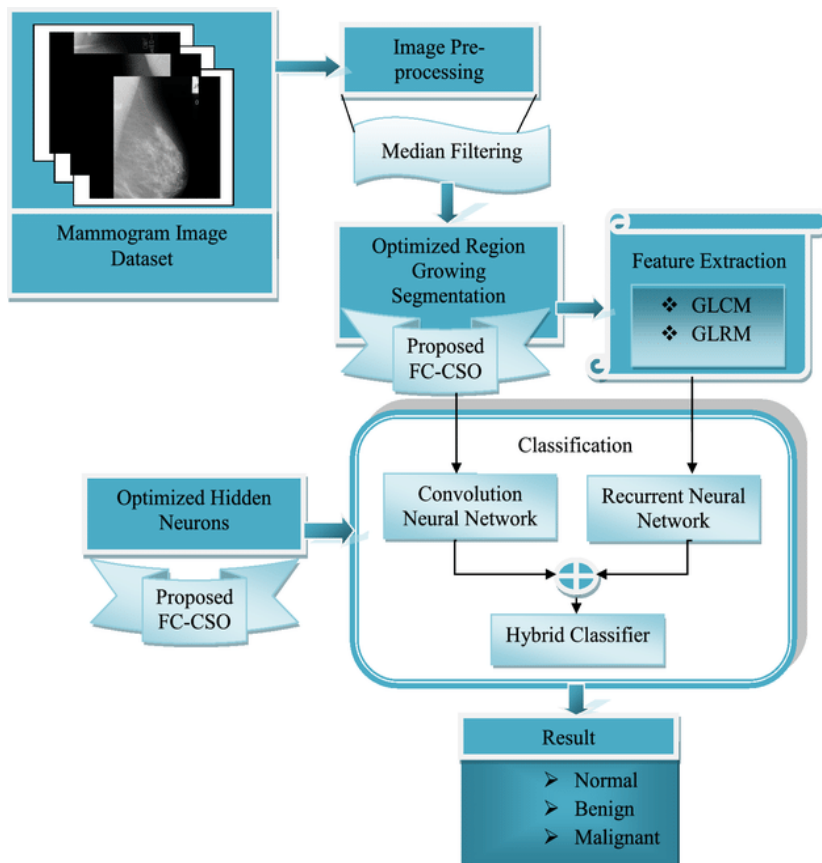
## 3. LITERATURE SURVEY:

T. Chen, and C. Guestrin, "Xgboost: A scalable tree boosting system," *Proceedings of the 22nd ACM SIGKDD International Conference on Knowledge Discovery and Data Mining*. ACM, 2016. pp. 785-794. T. Turki, and J. T. L. Wang, "Reverse Engineering Gene Regulatory Networks Using Sampling and Boosting Techniques," *Machine Learning and Data Mining in Pattern Recognition: 13th International Conference, MLDM 2017, New York, NY, USA, July 15-20, 2017, Proceedings*, P. Perner, ed., pp. 63-77, Cham: Springer International Publishing, 2017. C.-C. Chang, and C.-J. Lin, "LIBSVM: A library for support vector machines," *ACM Transactions on Intelligent Systems and Technology (TIST)*, vol. 2, no. 3, pp. 27, 2011. J. Zhang, S. O. Williams, and H. Wang,

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### Block Diagram:

The below Figure shows the Block diagram of the proposed system which has been



### Hardware requirements:

- Adaboost algorithm
- Deepboost algorithm
- Xg boost algorithm

**4.CONCLUSION:**

For the goal of cancer identification, we suggest employing machine learning algorithms such as DeepBoost, xgboost, a variation of AdaBoost, and SVM. Support vector machines work well, as evidenced by experimental results on actual clinical data relating to thyroid cancer, colon cancer, and liver cancer. Future work will consist of three main objectives: (1) enhancing the performance of machine learning algorithms through the use of transfer learning techniques as in [14], (2) enhancing the performance of the machine learning used in this study, and (3) combining the machine learning used in this study with feature learning techniques.

**FUTURE WORK:**

The EASY FARMING SYSTEM USING IOT FOR EFFICIENT PADDY GROWTH adding components make feasible. By adding automation process the results for the crop field management will give better results.

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