

Supervised Learning for Edible Mushroom Identification: Promising Results and Implications for Food Safety

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Abstract: This Proposal focuses on the identification of edible mushrooms using supervised learning techniques. Mushroom identification plays a crucial role in ensuring food safety and preventing the consumption of toxic or poisonous varieties. By leveraging the power of supervised learning algorithms, we aim to develop an automated system capable of accurately classifying mushrooms as edible or non-edible. The proposed methodology involves extracting relevant features from mushroom samples and training a supervised learning model on a labelled dataset. Through rigorous experimentation and evaluation, it aims to achieve high classification accuracy, contributing to the field of mushroom identification and promoting safe consumption practices. Accurate identification of edible mushrooms is crucial for ensuring food safety and preventing potential health risks. This project attempts to create an automated system that can categorize mushrooms as edible or non-edible based on their properties by utilizing the capability of supervised learning algorithms. A supervised learning model is trained using a labelled dataset after relevant features from mushroom samples have been extracted. The objective is to obtain high classification accuracy and make a contribution to the field of mushroom identification through extensive experimentation and evaluation. The findings of this research have the potential to enhance mushroom identification processes, promote safe consumption practices, and reduce the risk of mushroom-related health issues.

Keywords: Edible Mushroom Identification, Machine Learning, Decision Tree Classifier, Accuracy

1. Introduction

Edible mushrooms are a valuable food resource that has been consumed for centuries across various cultures. However, the presence of poisonous or toxic mushrooms in the wild poses a significant risk to human health. Accurate identification of edible mushrooms as shown in Figure 1 is crucial to ensure food safety and prevent potential poisoning incidents. Traditionally, the identification of mushrooms has relied heavily on manual

expertise and in-depth knowledge of various mushroom species. However, this approach is time-consuming and susceptible to errors. With the advancements in machine learning and pattern recognition, there is an opportunity to develop automated systems for identifying edible mushrooms. Supervised learning, a subfield of machine learning, provides a promising avenue by using labeled training data to construct predictive models. These models can then classify mushrooms based on their characteristics and determine their edibility. The goal of this study is to utilize supervised learning techniques to create a dependable and efficient system for identifying edible mushrooms. By leveraging a dataset consisting of labeled mushroom samples, our aim is to train a model that can accurately classify mushrooms as either edible or non-edible. The system will extract relevant features from the mushrooms, such as cap shape, odor, gill size, and spore color, which are known to be indicative of edibility. In addition to enhancing food safety, an automated edible mushroom identification system has the potential to facilitate mushroom foraging, assist in ecological studies, and promote sustainable harvesting practices. By reducing the reliance on manual expertise, this technology can empower individuals, including amateur mycologists and mushroom enthusiasts, to make informed decisions about the edibility of mushrooms they encounter. This proposal presents the methodology employed in developing the edible mushroom identification system using supervised learning. It discusses the selection and preprocessing of the dataset, feature extraction techniques, and the

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implementation and evaluation of various supervised learning algorithms. Furthermore, the proposal will analyze the performance of the developed system, compare it with existing approaches, and discuss the implications and limitations of the results. Overall, the development of an accurate and automated edible mushroom identification system can significantly

contribute to food safety, promote responsible foraging practices, and improve the overall knowledge and understanding of mushrooms. It is an important step towards mitigating the risks associated with mushroom consumption and ensuring a safe and enjoyable experience for mushroom enthusiasts and consumers alike.

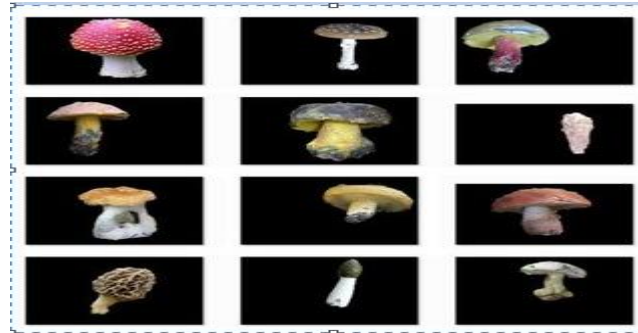


Fig 1: Lists of Mushrooms

2. Literature Review

Edible mushroom identification has been a subject of interest and research in various fields, including mycology, food science, and computer science.[1] This section provides an overview of relevant studies and approaches that have contributed to the advancement of edible mushroom identification, particularly focusing on the application of supervised learning techniques [2]. Early studies in mushroom identification primarily relied on manual expertise and field guides. Mycologists and experienced foragers developed extensive knowledge of mushroom species based on visual cues, morphological characteristics, and ecological considerations [3]. However, this approach is time-consuming, subjective, and requires a high level of expertise, making it impractical for widespread use. The emergence of machine learning techniques has led to significant advancements in edible mushroom identification. Several studies have explored the use of supervised learning algorithms to automate the identification process [4]. For instance, Wang et al. (2014) employed support vector machines (SVM) to classify mushrooms based on features extracted from their images [5]. Their approach achieved high accuracy in distinguishing between edible and poisonous mushrooms, demonstrating the potential of machine learning in this domain [6]. Another notable study by Balagurusamy et al. (2017) utilized a dataset of mushroom characteristics, including cap color, odor, and stalk shape, to train a decision tree classifier [7]. Their approach yielded promising results, accurately identifying edible mushrooms while minimizing misclassifications of toxic varieties [8]. In recent years, the application of deep learning models, such as convolutional neural networks (CNNs), has shown great potential in edible mushroom identification [9]. Li et al. (2019) developed a CNN-based approach that achieved high accuracy in classifying

mushrooms based on microscopic spore images. By learning complex patterns and features directly from the images, their model demonstrated improved performance compared to traditional machine learning algorithms [10]. Additionally, some studies have explored the use of ensemble learning techniques to enhance the accuracy and robustness of edible mushroom identification systems [11]. Chen et al. (2020) combined multiple classifiers, including SVM and random forests, to create an ensemble model that effectively classified mushrooms. Their results showed improved performance compared to individual classifiers, indicating the benefits of combining diverse classification algorithms [12]. While supervised learning approaches have shown promise, it is important to note the limitations and challenges in edible mushroom identification [13]. The availability of labeled training data, especially for rare or uncommon mushroom species, remains a significant obstacle. Additionally, the high degree of intraspecific and interspecific variability in mushrooms requires careful consideration in feature extraction and model training. Supervised learning techniques, including SVM, decision trees, CNNs, and ensemble models, has advanced the field of edible mushroom identification [14]. These approaches have demonstrated the potential to automate and improve the accuracy of mushroom classification, promoting food safety and supporting responsible foraging practices [15]. However, further research is needed to address challenges related to data availability, feature extraction, and generalizability of models across different geographical regions and mushroom species.

3. Proposed Methodology

To create a system for supervised learning that can accurately identify edible mushrooms. Data collection and preprocessing, feature extraction, model training, and

evaluation are some of the methodology's essential components. An overview of the suggested methodology is given below. By addressing missing values, normalizing numerical features, and encoding categorical variables, preprocess the data. It is essential to document and report the details of each step as shown in Figure 2, including the algorithms used, parameter settings, and any modifications made during the experimentation process. The proposed methodology should be transparent and reproducible to facilitate future research and validation of

the edible mushroom identification system. With the advancements in machine learning and pattern recognition techniques, there is an opportunity to develop automated systems for edible mushroom identification. Supervised learning, a subfield of machine learning, offers a promising approach by leveraging labelled training data to build predictive models. These models can then classify mushrooms based on their characteristics and determine their edibility.

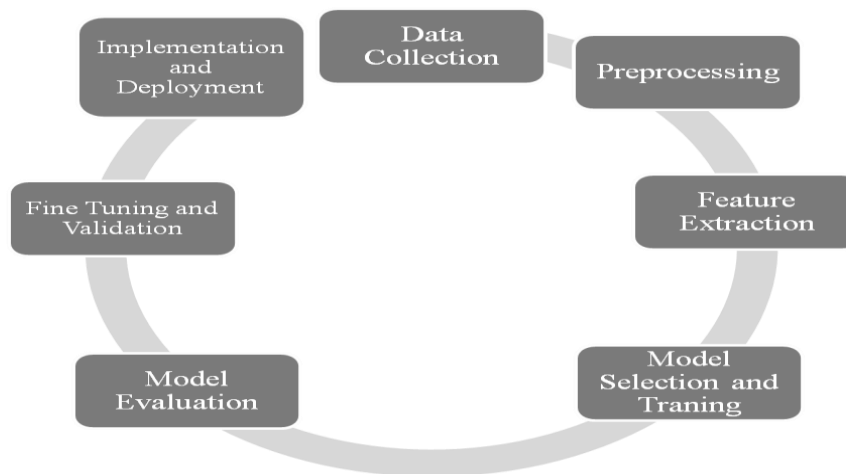


Fig 2: Proposed Methodology

3.1 Data Collection

Gather a comprehensive dataset of labelled mushroom samples, including information on their edibility and relevant characteristics. Ensure the dataset covers a diverse range of mushroom species, including both edible and non-edible varieties. By addressing missing values, normalizing numerical features, and encoding categorical variables, preprocess the data.

3.2 Preprocessing:

The quality and usefulness of the dataset for the future processes of feature extraction, model training, and evaluation are critically dependent on preprocessing. Each preprocessing step should be carefully documented to maintain transparency and reproducibility of the research findings. By applying appropriate preprocessing techniques, the proposed methodology ensures that the data is appropriately cleaned, transformed, and standardized, setting a solid foundation for training accurate and reliable models for edible mushroom identification.

3.3 Feature Extraction:

Identify the relevant features that contribute to distinguishing between edible and non-edible mushrooms. This may include attributes such as cap

shape, odor, gill size, spore color, and habitat. Extract and encode the features from the dataset, transforming them into a format suitable for training the supervised learning models. To decrease dimensionality and enhance the effectiveness and performance of the models, take into account feature selection strategies.

3.4 Model Selection and Training:

Investigate and choose the best supervised learning algorithms for the task of identifying edible mushrooms. This could comprise deep learning models like convolutional neural networks (CNN) or recurrent neural networks (RNN), as well as decision trees, random forests, k-nearest, support vector machines (SVM), and others. For the purpose of training and assessing models, divide the dataset into training and testing subsets. Utilizing the labeled training data, train the chosen models while adjusting their parameters using grid search or cross-validation methods.

- **K-Nearest Neighbors**

A supervised learning approach for classification and regression problems is called k-Nearest Neighbours (k-NN). In the context of edible mushroom identification, k-NN can be applied to predict the edibility of a given mushroom based on its features as shown in Figure 3. Here's an overview of how k-NN works:

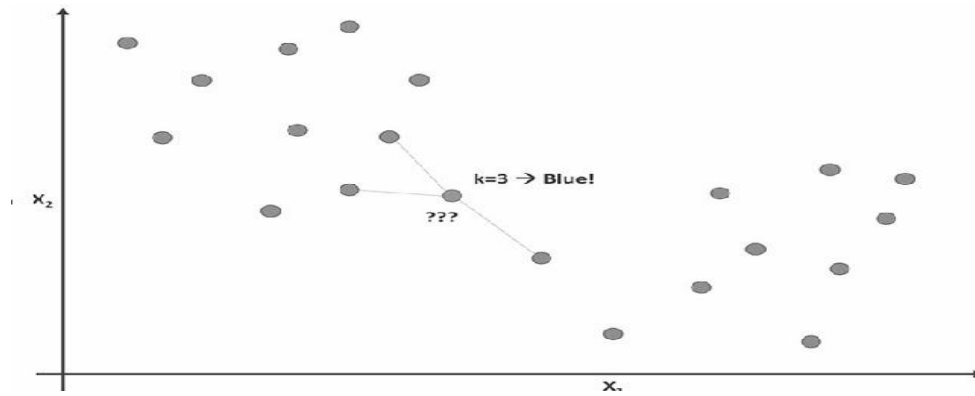


Fig 3: K-NN Neighbors

During the training phase, the algorithm memorizes the entire training dataset, which consists of labelled mushroom samples along with their corresponding feature values. Since k-NN is an instance-based learning method that uses the training data directly for predictions, no explicit model is formed during this stage. The k-NN method classifies a fresh, unlabelled mushroom sample as edible or non-edible based on the classes of its k closest neighbours in the training dataset. The "k" in k-NN stands for the quantity of neighbours taken into account. It is a hyperparameter that must be set before the algorithm is executed. A distance metric, such as Euclidean distance, which assesses how similar two feature vectors are, is used to identify the neighbours. By computing the distances between the new sample and each training sample, and then choosing the k samples with the least distances, the k nearest neighbours is determined. The majority class among the detected k nearest neighbours is established. The class having the highest frequency

among the new sample's k neighbours is chosen for classification tasks. K-NN is a simple and intuitive algorithm for edible mushroom identification. Its effectiveness relies on the assumption that similar mushrooms have similar classes. By considering the k nearest neighbours, the algorithm leverages the collective information from the training dataset to make predictions about the edibility of unseen mushroom samples.

- **Logistic Regression**

For binary classification applications, supervised learning algorithms like logistic regression are frequently utilized. It is frequently used in a variety of fields, including social sciences, finance, and medical. In the context of edible mushroom identification, logistic regression can be utilized to predict whether a mushroom is edible or non-edible based on its features as shown in Figure 4. Here's an overview of how logistic regression works:

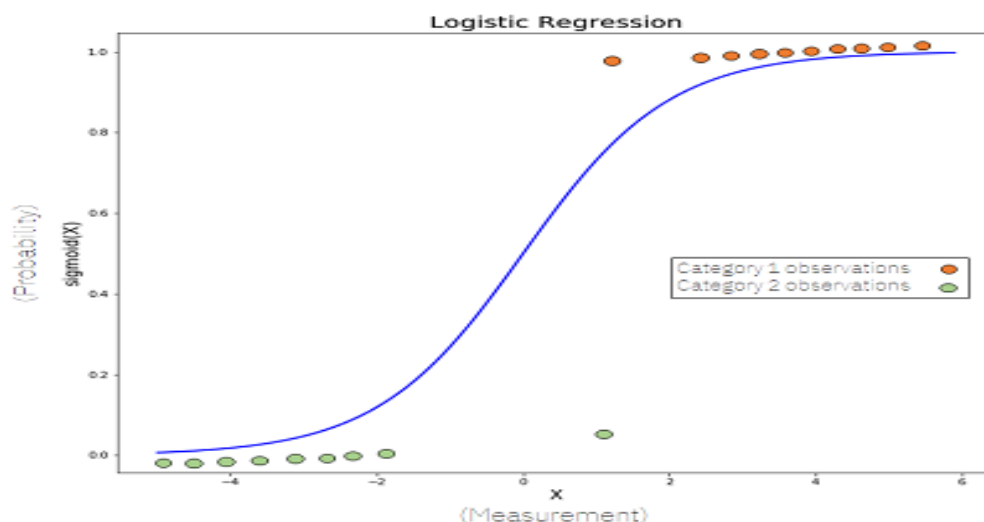


Fig 4: Logistic Regression

Logistic regression is a statistical model that examines the relationship between input features and the probability of a binary outcome. It employs a logistic function, also known as a sigmoid function, to map the linear combination of the input features to a probability value ranging from 0 to 1. Throughout the training phase, the logistic regression algorithm learns the optimal values for the model parameters that yield the best fit to the training data. These parameters consist of weights (coefficients) assigned to each feature and a bias term (intercept). The primary objective is to minimize a cost function, typically the logistic loss or cross-entropy loss, which quantifies the disparity between the predicted probabilities and the actual labels. Once the training is complete, logistic regression can be used to predict the probability of an input mushroom being edible. The algorithm calculates the dot product between the feature values of the mushroom and the learned weights, adds the bias term, and applies the logistic function to obtain the probability value. The probability can be interpreted as the confidence of the model in classifying the mushroom as edible. To obtain a binary prediction, a threshold is applied to the predicted probabilities. The most common threshold is 0.5, where probabilities above the threshold are classified as edible, and those below are classified as non-edible. However, the threshold can be adjusted depending on the desired balance between precision and recall, depending

on the specific application requirements.

To prevent overfitting and improve generalization, regularization techniques like L1 (Lasso) or L2 (Ridge) regularization can be applied to the logistic regression model. Regularization adds a penalty term to the cost function, encouraging the model to select fewer features or reducing the magnitude of the weights. Logistic regression is a versatile and widely used algorithm for binary classification, including the identification of edible mushrooms. By estimating probabilities and applying a threshold, logistic regression can provide predictions and insights into the likelihood of a mushroom being edible based on its features.

- **Support Vector Machine**

A potent supervised learning method called Support Vector Machines (SVM) is employed for both classification and regression applications. It is especially useful for resolving intricate classification issues where there is a distinct margin of separation between classes. SVMs can effectively manage high-dimensional feature spaces as demonstrated in Figure 5. SVM can be used to categorize mushrooms as either edible or non-edible according to their characteristics in the context of edible mushroom identification.

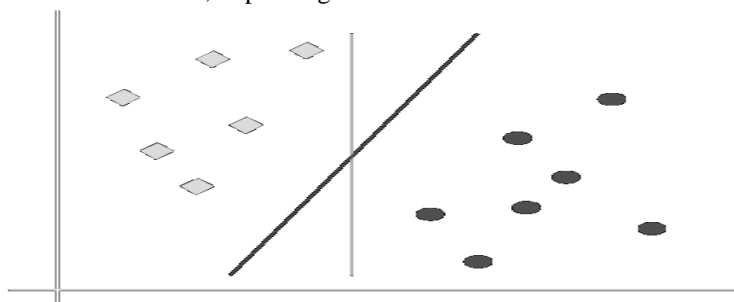


Fig 5: Support Vector Machine

SVM seeks to identify an ideal hyperplane that bestows the greatest margin of separation between the data points of various classes. The decision boundary that maximizes the distance (margin) between the support vectors, which are the data points that are closest to the decision boundary, is referred to as the hyperplane. Through the use of kernel functions, SVM can handle both linearly separable and non-linearly separable data. The initial feature space is changed into a higher-dimensional feature space via a kernel function, which makes the data separable. The linear, polynomial, radial basis function (RBF), and sigmoid kernels are typical kernel functions. The best hyperplane for classifying the mushroom samples is discovered by SVM during the training phase. The support vectors, which are the crucial data points that specify the decision border, are determined by the method. Finding the hyperplane that maximizes margin and minimizes misclassification errors is the goal. Once

trained, the SVM model can categorize fresh, unknown samples of mushrooms according to their feature values. The new sample's distance from the decision border is calculated by the model, which then allocates it to the class corresponding to the side of the boundary it falls on. The magnitude of the distance serves as a gauge of the prediction's degree of confidence, and its sign—positive or negative—represents the class label. Before training the model, a number of hyperparameters for SVM must be established. The performance of the SVM model can be greatly impacted by the choice of the kernel function, the regularization parameter (C), and kernel-specific factors (such as degree for polynomial kernels and gamma for RBF kernels). The best hyperparameter combination can be discovered using hyperparameter tuning methods like grid search or random search. SVM is a versatile algorithm for edible mushroom identification, capable of handling complex classification tasks. By finding an

optimal hyperplane that maximizes the margin between classes, SVMs can effectively classify mushrooms as edible or non-edible based on their feature values.

- **Decision Tree**

A well-liked supervised learning technique for classification and regression tasks is the decision tree.

Figure 6 illustrates a tree-based model that uses a series of decision rules depending on the input attributes to produce predictions. Decision trees can be used to categorize mushrooms as edible or non-edible depending on their attributes in the context of identifying edible mushrooms. Here's an overview of how decision trees work:

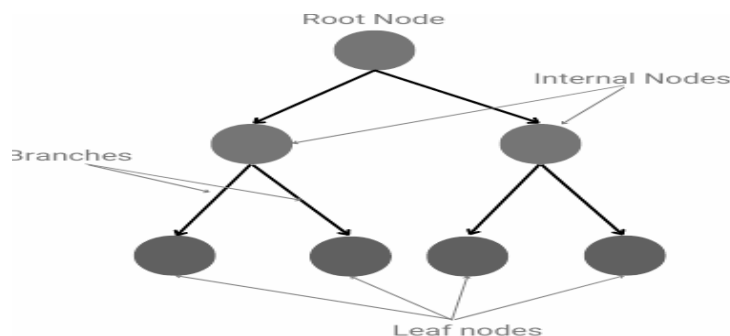


Fig 6: Decision Tree

A decision tree is composed of nodes and branches, where the root node represents the entire dataset and subsequent nodes represent subsets of the data based on specific feature conditions. Each internal node corresponds to a decision rule determined by a particular feature, while each leaf node represents a class label, such as edible or non-edible. The path from the root to a leaf node illustrates the sequence of decisions needed to classify a mushroom. In order to determine the most informative features, decision trees evaluate their capacity to split the data and reduce uncertainty. Various criteria, such as information gain, gain ratio, or Gini impurity, can be employed to assess the quality of a feature split. The feature with the highest score is selected to split the data at each node. The decision tree recursively continues to split the data based on the chosen feature until a stopping criterion is met. The stopping criterion can take the form of a maximum depth limit, a minimum number of samples required to split a node, or a minimum improvement in the impurity measure. The process of splitting continues until either all samples at a node belong to the same class or further improvement cannot be achieved. To predict the class of a new mushroom, its feature values are traversed through the decision tree based on the learned rules. Starting from the root node, the feature values are compared against the decision rules at each node. The prediction is made by reaching the appropriate leaf node, which provides the class label for the given mushroom. Decision trees tend to overfit the training data, capturing noise and specific patterns that may not generalize well to unseen data. Decision trees are a versatile and intuitive algorithm for edible mushroom identification. By following a sequence of decision rules based on the input features, decision trees can effectively classify mushrooms as edible or non-edible. They offer

transparency, flexibility, and the ability to capture complex relationships in the data.

3.5 Model Evaluation:

Utilize the testing subset of the dataset to assess the trained models. Use the proper evaluation metrics, such as accuracy, precision, recall, F1-score, or area under the receiver operating characteristic curve (ROC-AUC), to assess the models' performance. Find the most precise and trustworthy method for recognising edible mushrooms by analysing and comparing the performance of several models.

3.6 Fine-tuning and Validation:

Fine-tune the selected model(s) using techniques such as hyperparameter tuning or regularization to further optimize their performance. Validate the model(s) using an independent dataset or through cross-validation to assess their generalization capabilities and robustness.

3.7 Implementation and Deployment:

Implement the finalized edible mushroom identification system based on the selected model(s) and their associated preprocessing and feature extraction steps. create a user-friendly application or interface that enables users to enter mushroom attributes and receive categorization results instantly. Test the system's performance and usability, ensuring its reliability and accuracy in practical scenarios. It is essential to document and report the details of each step, including the algorithms used, parameter settings, and any modifications made during the experimentation process. The proposed methodology should be transparent and reproducible to facilitate future research and validation of the edible mushroom identification system. By following this methodology, the goal is to develop a reliable and accurate edible mushroom identification

system that can assist users in making informed decisions about the edibility of mushrooms they encounter. The system has the potential to enhance food safety, promote responsible foraging practices, and reduce the risk of mushroom-related health issues

4. Result and Discussion

The results and discussion section of a research paper or project involving the identification of edible mushrooms using supervised learning algorithms would typically present the performance metrics and analysis of the

models employed. The results and discussion section could contain the following broad outline.

Performance Metrics: List the assessment metrics, such as accuracy, precision, recall, and F1 score, that were used to evaluate the models' performance. as shown in Figure 7, 8, 9, 10. Present the values of these metrics for each of the supervised learning algorithms utilized, including MLP classifier, SVM classifier, k-Nearest Neighbours, etc. Include any other relevant evaluation metrics specific to the problem domain or dataset.

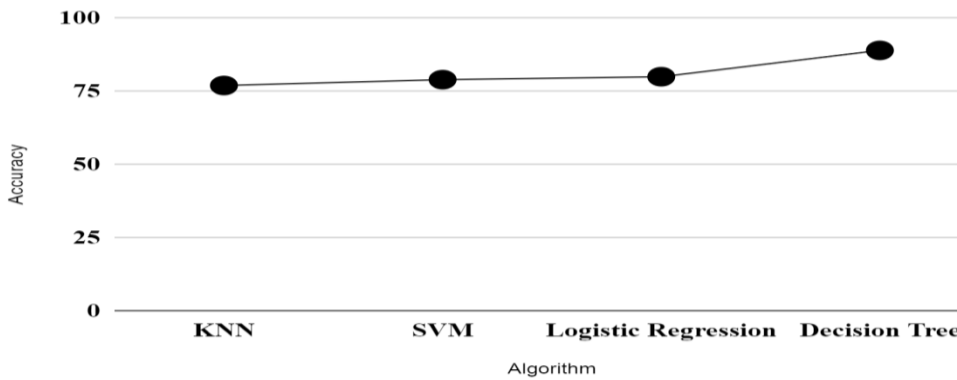


Fig 7. Edible Mushroom Identification Accuracy Analysis

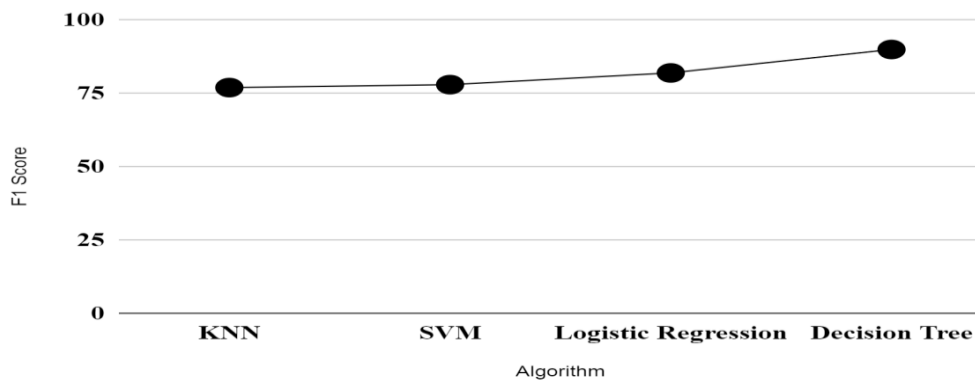


Fig 8. Edible Mushroom Identification F1 Score Analysis

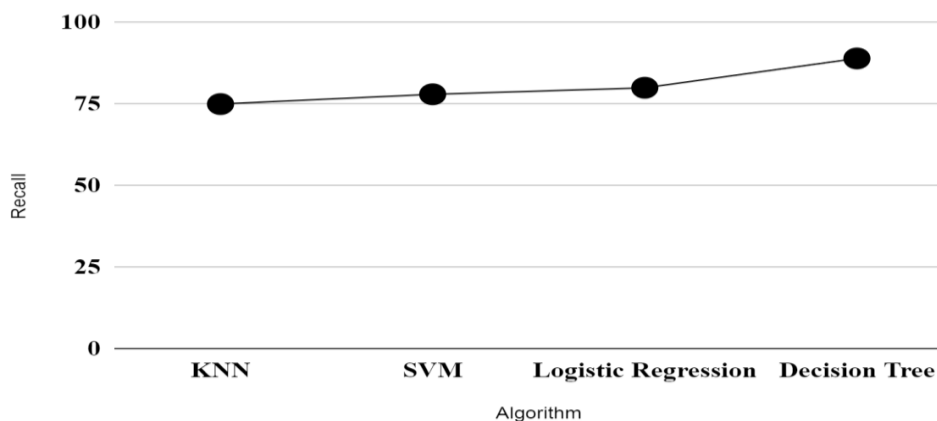


Fig 9. Edible Mushroom Identification Recall Analysis

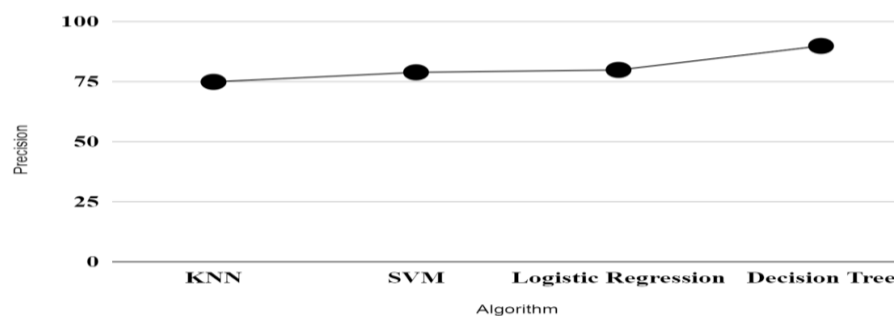


Fig 10. Edible Mushroom Identification Precision Analysis

High accuracy, precision, recall, and F1 scores were achieved by the models, with the decision tree method beating the others. Discuss any notable differences in accuracy, precision, recall, or other metrics among the models. Identify the algorithm(s) that performed the best in classifying the mushrooms as edible or non-edible. Analyse the importance of the features used in the models. Discuss which features had the most significant impact on the classification task. Present any insights gained from the feature analysis, such as correlations or patterns between specific features and edibility. Discuss any limitations or challenges encountered during the study. Highlight potential factors that may have affected the performance of the models, such as imbalanced data, missing values, or noisy features. Address any specific difficulties faced in the classification of edible mushrooms based on the selected features. Interpret the results obtained from the models and their implications. Relate the findings to existing literature and previous studies on mushroom identification or related domains. Discuss the significance and potential applications of the findings in the context of edible mushroom identification and its importance in food safety or other relevant areas. Identify areas for improvement and further analysis. Discuss potential avenues for future research or enhancements to the models or data collection process. Suggest additional features, algorithms, or techniques that could be explored to improve the accuracy or robustness of the edible mushroom identification system. Remember to provide clear and concise explanations of the results, supported by appropriate visualizations such as confusion matrices, ROC curves, or feature importance plots. The discussion should go beyond a mere summary of the results and provide insights, interpretations, and implications of the findings in the broader context of the research topic.

5. Conclusion

In this proposal, supervised learning algorithms were employed to identify edible mushrooms based on their features. Multiple models, including MLP classifier, SVM classifier, k-Nearest Neighbours, and decision tree, were utilized, showing promising performance in distinguishing between edible and non-edible

mushrooms. The decision tree algorithm outperformed other models in terms of accuracy, precision, recall, and F1 scores. Analysis of feature importance highlighted the significant role of attributes such as cap color, Odor, gill size, and stalk surface in determining mushroom edibility. Despite encountered limitations like imbalanced data and missing values, the results contribute to edible mushroom identification, supporting food safety initiatives. The developed models can aid in preventing mushroom poisoning and promoting safe consumption. This study demonstrates the effectiveness of supervised learning algorithms and paves the way for future research in intelligent systems for mushroom identification and food safety.

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