

Exploratory Data Analysis (EDA) & SVM Modeling Report

Dataset Overview

- **Dataset Name:** dataset.csv
 - **Target Variable:** Disease
 - **Features:** Symptom-based (e.g., Symptom_1 to Symptom_N)
 - **Preprocessing Applied:** Label encoding, binary encoding, standard scaling
 - **Objective:** Classify diseases based on reported symptoms
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Target Variable

The Disease column is a **multi-class categorical** variable, encoded numerically using LabelEncoder.

A bar plot (not shown here) should be used to visualize class distribution and detect imbalance.

Feature Insights

- The feature columns represent symptoms with categorical string values.
 - Many symptom values are missing (NaN or blank).
 - All symptom values were unified into a unique indexed list before being **binary encoded** for dimensionality reduction.
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⚙️ Data Preprocessing Pipeline

1. **Symptom Encoding:** Mapped all unique symptoms to integers.
 2. **Binary Encoding:** Converted symptom indexes into binary to reduce dimensionality.
 3. **Feature Scaling:** Used StandardScaler to normalize values.
 4. **Data Split:** 80/20 train-test split using train_test_split.
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Feature Selection

Used a **Random Forest Classifier** to identify and retain only the most important features via SelectFromModel.

Helped improve SVM performance by removing noise.

SVM Modeling

- **Classifier Used:** Support Vector Machine (SVC)
- **Kernel:** RBF (Radial Basis Function)
- **Hyperparameter Tuning:** GridSearchCV
- **Scoring Metric:** Accuracy

Results: - Best Cross-validation Accuracy: 0.9929 - Tuned Model Accuracy (Test Set): 0.9919

Evaluation

Used `classification_report()` for detailed evaluation. Output was visualized using a **heatmap** for precision, recall, and F1-score per class.

- **Number of Classes:** 41
- **Visualization Enhancements:**
 - Small font for class names
 - Padding and larger figure size for clarity
 - Colored summary row for average scores

(Include this visualization in your report or slide deck.)

Explainability (Optional)

- shap was imported, though not applied in the notebook.
 - SHAP can be used for model interpretation and feature contribution explanation if needed.
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Summary

- Binary encoding handled high-cardinality categorical features effectively.
 - Random Forest feature selection improved model focus.
 - SVM with hyperparameter tuning achieved nearly perfect classification.
 - Class-wise metrics were visualized to improve interpretability.
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Recommendations

- Add `df.info()` and `df.describe()` summaries to show structure and distribution.
- Visualize class distribution and missing data using seaborn/matplotlib.
- Export SHAP visualizations if interpretability is needed.