

# **Explainable Classification Models**



classification models that can be easily explainable



If you're looking for classification models that prioritize explainability, there are several options:

- 1. **Logistic Regression**: One of the simplest and most widely used classification algorithms. It models the probability of the default class using the logistic function.
- 2. **Decision Trees**: These models partition the feature space into a set of rectangular regions, and each region is assigned a class label. Decision trees are easily interpretable and can be visualized.
- 3. **Random Forests**: An ensemble learning method that builds multiple decision trees and merges their predictions. While random forests are more complex than individual decision trees, they can still provide insights into feature importance.
- 4. Gradient Boosting Machines (GBM): Similar to random forests, GBM builds an ensemble of decision trees, but it does so sequentially, with each tree correcting errors made by the previous one. Despite being more complex, GBM can still offer some level of interpretability, especially in terms of feature importance.
- 5. **Linear Support Vector Machines (SVM)**: SVM finds the hyperplane that best separates classes in high-dimensional space. When the data is linearly separable, SVM can provide clear decision boundaries and feature weights.
- 6. **K-Nearest Neighbors (KNN)**: KNN classifies new data points based on the majority class of their nearest neighbors. While KNN itself is straightforward, explaining predictions might require visualizing the nearest neighbors.
- 7. **Naive Bayes**: This algorithm is based on Bayes' theorem with the assumption of independence between features. It's simple and easy to implement, making it interpretable, though its assumption of feature independence might not hold in many real-world scenarios.
- 8. **Rule-Based Models**: These models directly represent decision rules inferred from the data. Examples include OneR (One Rule), which selects one rule based on a single feature, and RIPPER (Repeated Incremental Pruning to Produce Error Reduction), which builds a set of rules.

Each of these models has its strengths and weaknesses in terms of accuracy, complexity, and interpretability, so the choice depends on your specific requirements and constraints.





in python, write a code that uses linear sym to classification tasks



Sure, here's a simple example of how you can use Linear SVM for classification tasks in Python using the popular Scikit-learn library:

```
python
 from sklearn import datasets
 from sklearn.model_selection import train_test_split
 from sklearn.svm import LinearSVC
 from sklearn.metrics import accuracy_score
 # Load dataset (example using the Iris dataset)
 iris = datasets.load iris()
 X = iris.data
 y = iris.target
 # Split data into train and test sets
 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
 # Initialize and train the Linear SVM model
 svm model = LinearSVC()
 svm_model.fit(X_train, y_train)
 # Predict on the test set
 y_pred = svm_model.predict(X_test)
 # Calculate accuracy
 accuracy = accuracy_score(y_test, y_pred)
 print("Accuracy:", accuracy)
```

## In this code:

- We load the Iris dataset from Scikit-learn.
- Split the data into training and testing sets using `train test split`.
- Initialize the Linear SVM model using `LinearSVC`.
- Train the model using the training data.
- Make predictions on the test data.
- Calculate the accuracy of the model using `accuracy\_score` from Scikit-learn.

You can replace the Iris dataset with your own dataset by loading it into `x` and `y` variables. This code gives you a basic framework to start with Linear SVM classification tasks in Python.



give me a python code to explain a prediction on a trained LinearSVC





Certainly! To explain a prediction made by a trained LinearSVC model, you can use techniques such as examining the coefficients of the model or visualizing decision boundaries. Here's an example code that demonstrates how to explain a prediction by inspecting the coefficients of the LinearSVC model:

```
python
 from sklearn import datasets
 from sklearn.model selection import train test split
 from sklearn.svm import LinearSVC
 from sklearn.metrics import accuracy_score
 import numpy as np
 # Load dataset (example using the Iris dataset)
 iris = datasets.load_iris()
 X = iris.data
 y = iris.target
 # Split data into train and test sets
 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
 # Initialize and train the Linear SVM model
 svm_model = LinearSVC()
 svm_model.fit(X_train, y_train)
 # Select a sample from the test set to explain
 sample_index = 0
 sample = X_test[sample_index]
 true_label = y_test[sample_index]
 # Predict on the sample
 predicted_label = svm_model.predict([sample])[0]
 # Explain the prediction by examining coefficients
 coefficients = svm model.coef
 intercept = svm_model.intercept_
 print("Coefficients:", coefficients)
 print("Intercept:", intercept)
 # Optionally, you can also calculate the decision score
 decision_score = np.dot(coefficients, sample) + intercept
 print("Decision score:", decision_score)
 # Print the true and predicted labels
 print("True label:", true_label)
 print("Predicted label:", predicted_label)
```

#### In this code:

- We train a LinearSVC model on the Iris dataset.
- We select a sample from the test set to explain.
- We predict the label of the sample using the trained model.
- We examine the coefficients and intercept of the model to understand its decision boundary.
- Optionally, we calculate the decision score, which is the result of the dot product of the coefficients and the sample plus the intercept.
- Finally, we print the true and predicted labels to compare them.



You can further extend this code to visualize the decision boundaries or to use other techniques for explaining model predictions, such as LIME (Local Interpretable Model-agnostic Explanations) or SHAP (SHapley Additive exPlanations).



there is a more clear way to explain the prediction of a LinearSVC using the coefficients and intercepts?



Certainly! When explaining a prediction from a LinearSVC model, you can provide a more detailed breakdown of how each feature contributes to the final decision. Here's how you can do it:

- 1. Display the feature names (if available) along with their corresponding coefficients.
- 2. Show the contribution of each feature to the decision score.
- 3. Sum these contributions along with the intercept to get the final decision score.

Here's a more detailed and clear way to explain the prediction:

```
python
 from sklearn import datasets
 from sklearn.model_selection import train_test_split
 from sklearn.svm import LinearSVC
 import numpy as np
 # Load dataset (example using the Iris dataset)
 iris = datasets.load iris()
 X = iris.data
 y = iris.target
 # Feature names for clarity
 feature names = iris.feature names
 # Split data into train and test sets
 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
 # Initialize and train the Linear SVM model
 svm model = LinearSVC()
 svm_model.fit(X_train, y_train)
 # Select a sample from the test set to explain
 sample index = 0
 sample = X_test[sample_index]
 true_label = y_test[sample_index]
 # Predict on the sample
 predicted label = svm model.predict([sample])[0]
 # Get coefficients and intercept
 coefficients = svm_model.coef_
 intercept = svm_model.intercept_
 # Explain the prediction
 print(f"Explaining prediction for sample index {sample_index}:")
 print(f"True label: {true_label}")
 print(f"Predicted label: {predicted_label}\n")
 decision_scores = np.dot(coefficients, sample) + intercept
```



```
# Detailed breakdown
for i, class_label in enumerate(iris.target_names):
    print(f"Class {class_label} decision score breakdown:")
    for feature, coef in zip(feature_names, coefficients[i]):
        contribution = coef * sample[feature_names.index(feature)]
        print(f" Feature: {feature}, Coefficient: {coef:.4f}, Value:
{sample[feature_names.index(feature)]:.4f}, Contribution: {contribution:.4f}")
    print(f" Intercept: {intercept[i]:.4f}")
    print(f" Decision score: {decision_scores[i]:.4f}\n")
```

#### In this code:

- 1. Feature Names: We use `feature names` from the dataset for clarity.
- 2. **Decision Scores**: For each class, we calculate and display the contribution of each feature to the decision score.
- 3. **Explanation**: For each class, the detailed breakdown shows how each feature, along with its coefficient and value, contributes to the decision score.

This approach gives a clear and detailed explanation of how the LinearSVC model made its prediction for a specific sample, making it easier to understand the influence of each feature.



can you wrap the code that explain the prediction in a single python function?



Sure! Here is the code wrapped into a single Python function that explains the prediction of a `LinearSVC` model:

```
python
 from sklearn import datasets
 from sklearn.model_selection import train_test_split
 from sklearn.svm import LinearSVC
 import numpy as np
 def explain_prediction(svm_model, sample, feature_names, class_names):
     Explains the prediction of a LinearSVC model for a given sample.
     Parameters:
     svm model (LinearSVC): The trained LinearSVC model.
     sample (array): The sample to explain.
     feature_names (list): List of feature names.
     class_names (list): List of class names.
     # Predict on the sample
     predicted_label = svm_model.predict([sample])[0]
     # Get coefficients and intercept
     coefficients = svm model.coef
     intercept = svm_model.intercept_
     decision scores = np.dot(coefficients, sample) + intercept
     print(f"Explaining prediction for the given sample:")
     print(f"Predicted label: {class_names[predicted_label]}\n")
```



```
# Detailed breakdown
    for i, class label in enumerate(class names):
        print(f"Class {class_label} decision score breakdown:")
        for feature, coef in zip(feature_names, coefficients[i]):
             contribution = coef * sample[feature_names.index(feature)]
print(f" Feature: {feature}, Coefficient: {coef:.4f}, Value:
{sample[feature_names.index(feature)]:.4f}, Contribution: {contribution:.4f}")
        print(f"
                  Intercept: {intercept[i]:.4f}")
        print(f" Decision score: {decision_scores[i]:.4f}\n")
# Load dataset (example using the Iris dataset)
iris = datasets.load iris()
X = iris.data
y = iris.target
# Feature names and class names for clarity
feature names = iris.feature names
class_names = iris.target_names
# Split data into train and test sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
# Initialize and train the Linear SVM model
svm model = LinearSVC()
svm_model.fit(X_train, y_train)
# Select a sample from the test set to explain
sample index = 0
sample = X test[sample index]
# Explain the prediction
explain_prediction(svm_model, sample, feature_names, class_names)
```

In this function `explain\_prediction`:

## 1. Parameters:

- `svm\_model`: The trained `LinearSVC` model.
- `sample`: The sample to explain.
- `feature\_names`: List of feature names.
- `class\_names`: List of class names.

#### 2. Explanation:

- Predict the label for the given sample.
- Calculate the decision scores for each class by computing the dot product of coefficients and sample values plus the intercept.
- For each class, print out the contribution of each feature to the decision score, including the feature's name, coefficient, value, and contribution.

You can call this function with any sample from your test set to get a detailed explanation of the prediction.



rewrite the code but the sample is a single instance





Certainly! Here's the code with a modified `explain\_prediction` function that takes a single instance (sample) and provides a detailed explanation of the prediction:

# python from sklearn import datasets from sklearn.model\_selection import train\_test\_split from sklearn.svm import LinearSVC import numpy as np def explain\_prediction(svm\_model, sample, feature\_names, class\_names): Explains the prediction of a LinearSVC model for a given sample. Parameters: svm\_model (LinearSVC): The trained LinearSVC model. sample (array): The sample to explain. feature\_names (list): List of feature names. class\_names (list): List of class names. # Predict on the sample predicted label = svm model.predict([sample])[0] # Get coefficients and intercept coefficients = svm model.coef intercept = svm\_model.intercept decision\_scores = np.dot(coefficients, sample) + intercept print(f"Explaining prediction for the given sample:") print(f"Predicted label: {class\_names[predicted\_label]}\n") # Detailed breakdown for i, class label in enumerate(class names): print(f"Class {class\_label} decision score breakdown:") print(f" Decision score: {decision\_scores[i]:.4f}\n") # Load dataset (example using the Iris dataset) iris = datasets.load iris() X = iris.datay = iris.target # Feature names and class names for clarity feature\_names = iris.feature\_names class\_names = iris.target\_names # Split data into train and test sets $X_{\text{train}}$ , $X_{\text{test}}$ , $y_{\text{train}}$ , $y_{\text{test}}$ = $train_{\text{test}}$ split(X, Y, $test_{\text{size}} = 0.3$ , $test_{\text{size}} = 0.3$ , $test_{\text{test}}$ size = 0.3, $test_{\text{te$ # Initialize and train the Linear SVM model svm model = LinearSVC() svm\_model.fit(X\_train, y\_train) # Select a sample from the test set to explain $sample_index = 0$ sample = X\_test[sample\_index] # Explain the prediction explain\_prediction(svm\_model, sample, feature\_names, class\_names)



## **Explanation:**

### 1. Function `explain\_prediction`:

- Takes a trained `LinearSVC` model, a single sample, feature names, and class names as inputs.
- Predicts the label for the given sample.
- Calculates the decision scores for each class by computing the dot product of coefficients and sample values plus the intercept.
- Prints out the detailed contribution of each feature to the decision score for each class, including the feature's name, coefficient, value, and contribution.

## 2. Usage:

- Load the Iris dataset and split it into training and test sets.
- o Train a `LinearSVC` model.
- Select a single instance from the test set.
- Call the `explain\_prediction` function with the trained model, the selected sample, feature names, and class names to get a detailed explanation of the prediction.



the "decision\_scores = np.dot(coefficients, sample) + intercept" part of the code is not working when sample is a single instance



You're right. The issue arises because the `sample` should be a 1D array, and `np.dot` needs the shapes to be compatible. In this case, you should reshape the `sample` to ensure that it works correctly with `np.dot`. Here's the corrected code:

```
python
 from sklearn import datasets
 from sklearn.model selection import train test split
 from sklearn.svm import LinearSVC
 import numpy as np
 def explain_prediction(svm_model, sample, feature_names, class_names):
     Explains the prediction of a LinearSVC model for a given sample.
     Parameters:
     svm_model (LinearSVC): The trained LinearSVC model.
     sample (array): The sample to explain.
     feature_names (list): List of feature names.
     class_names (list): List of class names.
     # Predict on the sample
     predicted label = svm model.predict([sample])[0]
     # Get coefficients and intercept
     coefficients = svm model.coef
     intercept = svm_model.intercept_
     # Calculate decision scores
```



```
decision scores = np.dot(coefficients, sample) + intercept
    print(f"Explaining prediction for the given sample:")
    print(f"Predicted label: {class_names[predicted_label]}\n")
    # Detailed breakdown
    for i, class_label in enumerate(class_names):
        print(f"Class {class_label} decision score breakdown:")
        for j, feature in enumerate(feature_names):
            contribution = coefficients[i][\overline{j}] * sample[j]
            print(f" Feature: {feature}, Coefficient: {coefficients[i][j]:.4f}, Value:
print(f" Decision score: {decision_scores[i]:.4f}\n")
# Load dataset (example using the Iris dataset)
iris = datasets.load_iris()
X = iris.data
y = iris.target
# Feature names and class names for clarity
feature_names = iris.feature_names
class names = iris.target names
# Split data into train and test sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
# Initialize and train the Linear SVM model
svm model = LinearSVC()
svm_model.fit(X_train, y_train)
# Select a sample from the test set to explain
sample index = 0
sample = X test[sample index]
# Explain the prediction
explain_prediction(svm_model, sample, feature_names, class_names)
```

## **Explanation of Changes:**

## 1. `sample` Handling:

- Ensure `sample` is passed as a 1D array directly.
- `np.dot(coefficients, sample)` works correctly with `sample` as a 1D array, because `coefficients` is a 2D array (classes × features).

#### 2. Explanation Breakdown:

• The function `explain\_prediction` iterates over the classes and features to explain the contribution of each feature to the decision score.

This version ensures that the `np.dot` operation works correctly with the given sample and provides a clear explanation of the prediction.

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