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from sklearn.datasets import load_iris
from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
import pandas as pd

iris = load_iris()

df = pd.DataFrame(data=iris.data, columns=iris.feature_names)

df.head()

{"summary":{"\n  \"name\": \"df\",\n  \"rows\": 150,\n  \"fields\": [\n    {\n      \"column\": \"sepal length (cm)\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.8280661279778629,\n        \"min\": 4.3,\n        \"max\": 7.9,\n        \"num_unique_values\": 35,\n        \"samples\": [\n          6.2,\n          4.5,\n          5.6\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      },\n      \"column\": \"sepal width (cm)\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.435866284936698,\n        \"min\": 2.0,\n        \"max\": 4.4,\n        \"num_unique_values\": 23,\n        \"samples\": [\n          2.3,\n          4.0,\n          3.5\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      },\n      \"column\": \"petal length (cm)\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 1.7652982332594667,\n        \"min\": 1.0,\n        \"max\": 6.9,\n        \"num_unique_values\": 43,\n        \"samples\": [\n          6.7,\n          3.8,\n          3.7\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      },\n      \"column\": \"petal width (cm)\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.7622376689603465,\n        \"min\": 0.1,\n        \"max\": 2.5,\n        \"num_unique_values\": 22,\n        \"samples\": [\n          0.2,\n          1.2,\n          1.3\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      }\n    ]\n  ],\n  \"type\": \"dataframe\", \"variable_name\": \"df\"}

df['target'] = iris.target

df.head()

{"summary":{"\n  \"name\": \"df\",\n  \"rows\": 150,\n  \"fields\": [\n    {\n      \"column\": \"sepal length (cm)\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.8280661279778629,\n        \"min\": 4.3,\n        \"max\": 7.9,\n        \"num_unique_values\": 35,\n        \"samples\": [\n          6.2,\n          4.5,\n          5.6\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      },\n      \"column\": \"sepal width (cm)\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.435866284936698,\n        \"min\": 2.0,\n        \"max\": 4.4,\n        \"num_unique_values\": 23,\n        \"samples\": [\n          2.3,\n          4.0,\n          3.5\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      },\n      \"column\": \"petal length (cm)\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 1.7652982332594667,\n        \"min\": 1.0,\n        \"max\": 6.9,\n        \"num_unique_values\": 43,\n        \"samples\": [\n          6.7,\n          3.8,\n          3.7\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      },\n      \"column\": \"petal width (cm)\",\n      \"properties\": {\n        \"dtype\": \"number\",\n        \"std\": 0.7622376689603465,\n        \"min\": 0.1,\n        \"max\": 2.5,\n        \"num_unique_values\": 22,\n        \"samples\": [\n          0.2,\n          1.2,\n          1.3\n        ],\n        \"semantic_type\": \"\",\n        \"description\": \"\"\n      }\n    ]\n  ],\n  \"type\": \"dataframe\", \"variable_name\": \"df\"}

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df.tail()
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df['species'] = df['target'].map({0: 'setosa', 1: 'versicolor', 2: 'virginica'})

df.head()

{"summary": {"name": "df", "rows": 150, "fields": [{"column": "sepal length (cm)", "properties": {"dtype": "number", "std": 0.8280661279778629, "min": 4.3, "max": 7.9, "num_unique_values": 35, "samples": [4.5, 5.6], "semantic_type": "number"}, "description": "sepal length (cm)"}, {"column": "sepal width (cm)", "properties": {"dtype": "number", "std": 0.435866284936698, "min": 2.0, "max": 4.4, "num_unique_values": 23, "samples": [2.3, 4.0, 3.5], "semantic_type": "number"}, "description": "sepal width (cm)"}, {"column": "petal length (cm)", "properties": {"dtype": "number", "std": 1.7652982332594667, "min": 1.0, "max": 6.9, "num_unique_values": 43, "samples": [3.8, 3.7], "semantic_type": "number"}, "description": "petal length (cm)"}, {"column": "petal width (cm)", "properties": {"dtype": "number", "std": 0.7622376689603465, "min": 0.1, "max": 2.5, "num_unique_values": 22, "samples": [0.2, 1.2, 1.3], "semantic_type": "number"}, "description": "petal width (cm)"}, {"column": "target", "properties": {"dtype": "number", "std": 0, "min": 0, "max": 2, "num_unique_values": 3, "samples": [0, 1, 2], "semantic_type": "number"}, "description": "target"}, {"column": "species", "properties": {"dtype": "category", "num_unique_values": 3, "samples": ["setosa", "versicolor", "virginica"], "semantic_type": "category"}, "description": "species"}]}, "type": "dataframe", "variable_name": "df"}

df.tail()

{"summary": {"name": "df", "rows": 5, "fields": [{"column": "sepal length (cm)", "properties": {"dtype": "number", "std": 0.30331501776206193, "min": 5.9, "max": 6.7, "num_unique_values": 5, "samples": [6.3, 5.9], "semantic_type": "number"}, "description": "sepal length (cm)"}, {"column": "sepal width (cm)", "properties": {"dtype": "number", "std": 0.17051384725156017, "min": 2.4, "max": 4.4, "num_unique_values": 19, "samples": [2.5, 4.4, 3.7], "semantic_type": "number"}, "description": "sepal width (cm)"}, {"column": "petal length (cm)", "properties": {"dtype": "number", "std": 1.4639146609994677, "min": 1.0, "max": 6.9, "num_unique_values": 43, "samples": [3.8, 3.7, 5.0], "semantic_type": "number"}, "description": "petal length (cm)"}, {"column": "petal width (cm)", "properties": {"dtype": "number", "std": 0.7622376689603465, "min": 0.1, "max": 2.5, "num_unique_values": 22, "samples": [0.2, 1.2, 1.3], "semantic_type": "number"}, "description": "petal width (cm)"}, {"column": "target", "properties": {"dtype": "number", "std": 0, "min": 0, "max": 2, "num_unique_values": 3, "samples": [0, 1, 2], "semantic_type": "number"}, "description": "target"}, {"column": "species", "properties": {"dtype": "category", "num_unique_values": 3, "samples": ["setosa", "versicolor", "virginica"], "semantic_type": "category"}, "description": "species"}]}, "type": "dataframe", "variable_name": "df"}

```

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```

```

X = df.drop(['target', 'species'], axis=1)
y = df['target']

```

```

X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.3, random_state=42)

```

```

clf = DecisionTreeClassifier()

```

```

clf.fit(X_train, y_train)

```

```

DecisionTreeClassifier()

```

```

predictions = clf.predict(X_test)
accuracy = accuracy_score(y_test, predictions)
print("Accuracy:", accuracy)

```

```

Accuracy: 1.0

```

```

print("\nEnter flower details to predict species:")
sepal_length = float(input("Sepal length (cm): "))
sepal_width = float(input("Sepal width (cm): "))
petal_length = float(input("Petal length (cm): "))

```

```

petal_width = float(input("Petal width (cm): "))
user_input = [[sepal_length, sepal_width, petal_length, petal_width]]
prediction = clf.predict(user_input)[0]
predicted_species = iris.target_names[prediction]
print("\n The predicted species is:", predicted_species)

```

Enter flower details to predict species:

Sepal length (cm): 3

Sepal width (cm): 2

Petal length (cm): 1

Petal width (cm): 4

NameError Traceback (most recent call last)

/tmp/ipython-input-4-433288988.py in <cell line: 0>()

5 petal_width = float(input("Petal width (cm): "))

6 user_input = [[sepal_length, sepal_width, petal_length, petal_width]]

----> 7 prediction = clf.predict(user_input)[0]

8 predicted_species = iris.target_names[prediction]

9 print("\n The predicted species is:", predicted_species)

NameError: name 'clf' is not defined

```

from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split
from sklearn.datasets import load_iris
import pandas as pd

```

Load data and create DataFrame

iris = load_iris()

df = pd.DataFrame(data=iris.data, columns=iris.feature_names)

df['target'] = iris.target

df['species'] = df['target'].map({0: 'setosa', 1: 'versicolor', 2: 'virginica'})

print("\nEnter flower details to predict species:")

sepal_length=float(input("sepal length(cm):"))

sepal_width=float(input("sepal width(cm):"))

petal_length=float(input("petal length(cm):"))

petal_width=float(input("petal width(cm):"))

user_input = [[sepal_length, sepal_width, petal_length, petal_width]]

Define clf here

clf = DecisionTreeClassifier()

Fit the classifier

X = df.drop(['target', 'species'], axis=1)

```
y = df['target']
X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.3, random_state=42)
clf.fit(X_train, y_train)
```

```
prediction = clf.predict(user_input)[0]
predicted_species = iris.target_names[prediction]
print("\n The predicted species is:", predicted_species)
```

Enter flower details to predict species:

sepal length(cm):3

sepal width(cm):2

petal length(cm):1

petal width(cm):4

The predicted species is: setosa

```
/usr/local/lib/python3.11/dist-packages/sklearn/utils/
validation.py:2739: UserWarning: X does not have valid feature names,
but DecisionTreeClassifier was fitted with feature names
  warnings.warn(
```

```
print(iris.target_names)
```

```
['setosa' 'versicolor' 'virginica']
```