```
# Install dtreeviz library !pip -qq install dtreeviz
```

```
- 91.8/91.8 kB 4.1 MB/s eta 0:00:00
import numpy as np
from matplotlib import pyplot as plt
from matplotlib.colors import ListedColormap
from sklearn import datasets
from sklearn import tree
from sklearn.tree import DecisionTreeClassifier, DecisionTreeRegressor
import graphviz
import dtreeviz
#from dtreeviz.trees import dtreeviz
# to import color map
# to import DT classifier and Regressor
# to import graphviz
# import the dtreeviz module directly
# The dtreeviz function is within the dtreeviz module
# Prepare the data into X (predictor) and Y (target), considering sepal length and width
iris = datasets.load_iris()
X = iris.data[:, 2:]
y = iris.target
# Fit the DT classifier with max depth = 2
clf = DecisionTreeClassifier(max_depth = 2, random_state=1234)
model = clf.fit(X, y)
text representation = tree.export text(clf)
# Display result
print(text representation)
# Save in figure
with open("decision_tree.log", "w") as f_out:
  f out.write(text representation)
# Save in figure
with open("decision tree.log", "w") as f out:
  f_out.write(text_representation)
# Save the figure to the .png file
fig.savefig("decision tree.png")
# Predicting probability of a class
clf.predict proba([[5, 1.5]])
# Predicting a class
```

```
clf.predict([[5, 1.5]])
# Load iris dataset and define variables with sepal length and width
iris = datasets.load iris()
X = iris.data[:, 2:]
y = iris.target
# First tree without restrictions
tree clf = DecisionTreeClassifier(random state=42)
tree_clf.fit(X, y)
# Second tree with hyperparameters
                   DecisionTreeClassifier(max depth
                                                        =2,
                                                                 min samples leaf
                                                                                        =1,
tree clf2
            =
min_samples_split = 2, random_state=2)
tree clf2.fit(X, y)
# Define a function for plotting decision boundary
def plot decision boundary(clf, X, y, axes=[0, 7.5, 0, 3], iris = True, legend=False,
plot training=True):
  # define array for x1 and x2 axes
  x1s = np.linspace(axes[0], axes[1], 100)
  x2s = np.linspace(axes[2], axes[3], 100)
  # make N-D coordinate arrays for vectorized evaluations of N-D scalar/vector fields over
N-D grids
  x1, x2 = np.meshgrid(x1s, x2s)
  # the numpy.ravel() functions returns contiguous flattened array(1D array with all the
input-array elements and with the same type a
  X new = np.c [x1.ravel(), x2.ravel()]
  # predict and reshape the y_pred according to x
  y_pred = clf.predict(X_new).reshape(x1.shape)
  # module is used for mapping numbers to colors or color specification conversion in a 1-
D array of colors also known as colormap
  custom_cmap = ListedColormap(['#fafab0','#9898ff','#a0faa0'])
  plt.contourf(x1, x2, y pred, alpha=0.3, cmap=custom cmap)
  if plot training:
    # plot Setosa in yellow
    plt.plot(X[:, 0][y==0], X[:, 1][y==0], "yo", label="Iris setosa")
    # plot Versicolor in blue
    plt.plot(X[:, 0][y==1], X[:, 1][y==1], "bs", label="Iris versicolor")
    # plot Virginica in green
    plt.plot(X[:, 0][y==2], X[:, 1][y==2], "g^", label="Iris virginica")
    plt.axis(axes)
  if iris:
    # define x axes label
    plt.xlabel("Sepal length", fontsize=14)
    # define y_axes label
```

```
plt.ylabel("Sepal width", fontsize=14)
 if legend:
    plt.legend(loc="lower right", fontsize=14)
# Plot both the decision tree
plt.figure(figsize=(8, 4))
# call the plot decision boundary function for tree clf
plot decision boundary(tree clf, X, y)
plt.plot([2.45, 2.45], [0, 3], "k-", linewidth=2)
plt.plot([2.45, 7.5], [1.75, 1.75], "k--", linewidth=2)
plt.plot([4.95, 4.95], [0, 1.75], "k:", linewidth=2)
plt.plot([4.85, 4.85], [1.75, 3], "k:", linewidth=2)
plt.title("No restrictions", fontsize=16)
plt.text(1.40, 1.0, "Depth=0", fontsize=15)
plt.text(3.2, 1.80, "Depth=1", fontsize=13)
plt.text(4.05, 0.5, "(Depth=2)", fontsize=11)
plt.figure(figsize=(8, 4))
# call the plot decision boundary function for tree clf2
plot decision boundary(tree clf2, X, y)
plt.plot([2.45, 2.45], [0, 3], "k-", linewidth=2)
plt.plot([2.45, 7.5], [1.75, 1.75], "k--", linewidth=2)
plt.plot([4.95, 4.95], [0, 1.75], "k:", linewidth=2)
plt.plot([4.85, 4.85], [1.75, 3], "k:", linewidth=2)
plt.title("Regularizing Hyperparameters", fontsize=16)
plt.text(1.40, 1.0, "Depth=0", fontsize=15)
plt.text(3.2, 1.80, "Depth=1", fontsize=13)
plt.text(4.05, 0.5, "(Depth=2)", fontsize=11)
plt.show()
# Prepare the data
# Instead of using load boston, we will use fetch california housing
from sklearn.datasets import fetch california housing
housing = fetch california housing()
X = housing.data
y = housing.target
housing.feature names
# Fit the regressor, set max depth = 3
dt = DecisionTreeRegressor(max_depth=3, random_state=1234)
model = dt.fit(X, y)
text representation = tree.export text(dt)
print(text representation)
# Visualize tree using plot tree
import matplotlib.pyplot as plt
from sklearn import tree
```

```
fig = plt.figure(figsize=(15, 10))
# Instead of using a subset, provide all feature names if the tree was trained with all
features
_ = tree.plot_tree(dt, feature_names=housing.feature_names, filled=True)
# Visualize tree using graphviz
dot_data = tree.export_graphviz(dt, out_file=None,
                  feature names=housing.feature names,
                  filled=True)
graphviz.Source(dot_data, format="png")
# Define plot regression function
def plot_regression_predictions(tree_reg, X, y, axes=[0.3, 1, 0, 60], ylabel="$y$"):
  # creating the x-axes grid in array
  x1 = np.linspace(axes[0], axes[1], 500).reshape(-1, 1)
  # define y
  y pred = tree reg.predict(x1)
  plt.axis(axes)
  plt.xlabel("$x 1$", fontsize=18)
  if ylabel:
    plt.ylabel(ylabel, fontsize=18, rotation=0)
  plt.plot(X, y, "b.")
  # plot y hat (predicted values) in red line in both graphs
  plt.plot(x1, y_pred, "r.-", linewidth=2, label=r"$\hat{y}$")
# Take index value 4 i.e, NOX
X = housing.data[:,4:5]
y = housing.target
# Define model with no hyperparameter
dt = DecisionTreeRegressor(random_state=1234)
dt.fit(X, y)
# Define model with maximum depth = 8
dt2 = DecisionTreeRegressor(max depth=8, random state=1234)
dt2.fit(X, y)
fig, axes = plt.subplots(ncols=2, figsize=(10, 4), sharey=True)
plt.sca(axes[0])
# Plot decision boundary
plot_regression_predictions(dt2, X, y)
for split, style in ((0.1973, "k-"), (0.0917, "k--"), (0.7718, "k--")):
# Plot the fit regression line
plt.plot([split, split], [-0.2, 1], style, linewidth=2)
plt.legend(loc="upper center", fontsize=18)
plt.title("max depth=8", fontsize=14)
plt.sca(axes[1])
# Plot dcision boundary
```

```
plot_regression_predictions(dt, X, y, ylabel=None)
for split, style in ((0.1973, "k-"), (0.0917, "k--"), (0.7718, "k--")):
# Plot the fit regression line
plt.plot([split, split], [-0.2, 1], style, linewidth=2)
plt.title("max_depth=No restriction", fontsize=14)
plt.text(0.3,1, "X1 = NOX feature", fontsize=10)
plt.show()
```