import matplotlib.pyplot as plt mpl.rc('axes', labelsize=14) mpl.rc('xtick', labelsize=12) mpl.rc('ytick', labelsize=12) # Where to save the figures PROJECT_ROOT_DIR = "." CHAPTER ID = "ensembles" IMAGES_PATH = os.path.join(PROJECT_ROOT_DIR, "images", CHAPTER_ID) os.makedirs(IMAGES_PATH, exist_ok=**True**) def save_fig(fig_id, tight_layout=True, fig_extension="png", resolution=300): path = os.path.join(IMAGES_PATH, fig_id + "." + fig_extension) print("Saving figure", fig_id) if tight_layout: plt.tight_layout() plt.savefig(path, format=fig_extension, dpi=resolution) options(jupyter.plot_mimetypes = c("text/plain", "image/png")) Prepare the data. Load breast cancer data In [50]: from sklearn.datasets import load_breast_cancer cancer = load_breast_cancer() print(cancer.target[0:20]) print(list(cancer.target names)) print(cancer.data[0:5]) print(list(cancer.feature_names)) ['malignant', 'benign'] [[1.799e+01 1.038e+01 1.228e+02 1.001e+03 1.184e-01 2.776e-01 3.001e-01 1.471e-01 2.419e-01 7.871e-02 1.095e+00 9.053e-01 8.589e+00 1.534e+02 6.399e-03 4.904e-02 5.373e-02 1.587e-02 3.003e-02 6.193e-03 2.538e+01 1.733e+01 1.846e+02 2.019e+03 1.622e-01 6.656e-01 7.119e-01 2.654e-01 4.601e-01 1.189e-01] [2.057e+01 1.777e+01 1.329e+02 1.326e+03 8.474e-02 7.864e-02 8.690e-02 7.017e-02 1.812e-01 5.667e-02 5.435e-01 7.339e-01 3.398e+00 7.408e+01 5.225e-03 1.308e-02 1.860e-02 1.340e-02 1.389e-02 3.532e-03 2.499e+01 2.341e+01 1.588e+02 1.956e+03 1.238e-01 1.866e-01 2.416e-01 1.860e-01 2.750e-01 8.902e-02] [1.969e+01 2.125e+01 1.300e+02 1.203e+03 1.096e-01 1.599e-01 1.974e-01 1.279e-01 2.069e-01 5.999e-02 7.456e-01 7.869e-01 4.585e+00 9.403e+01 6.150e-03 4.006e-02 3.832e-02 2.058e-02 2.250e-02 4.571e-03 2.357e+01 2.553e+01 1.525e+02 1.709e+03 1.444e-01 4.245e-01 4.504e-01 2.430e-01 3.613e-01 8.758e-02] [1.142e+01 2.038e+01 7.758e+01 3.861e+02 1.425e-01 2.839e-01 2.414e-01 1.052e-01 2.597e-01 9.744e-02 4.956e-01 1.156e+00 3.445e+00 2.723e+01 9.110e-03 7.458e-02 5.661e-02 1.867e-02 5.963e-02 9.208e-03 1.491e+01 2.650e+01 9.887e+01 5.677e+02 2.098e-01 8.663e-01 6.869e-01 2.575e-01 6.638e-01 1.730e-01] [2.029e+01 1.434e+01 1.351e+02 1.297e+03 1.003e-01 1.328e-01 1.980e-01 1.043e-01 1.809e-01 5.883e-02 7.572e-01 7.813e-01 5.438e+00 9.444e+01 1.149e-02 2.461e-02 5.688e-02 1.885e-02 1.756e-02 5.115e-03 2.254e+01 1.667e+01 1.522e+02 1.575e+03 1.374e-01 2.050e-01 4.000e-01 1.625e-01 2.364e-01 7.678e-02]] ['mean radius', 'mean texture', 'mean perimeter', 'mean area', 'mean smoothness', 'mean compactness', 'mean concavity , 'mean concave points', 'mean symmetry', 'mean fractal dimension', 'radius error', 'texture error', 'perimeter erro r', 'area error', 'smoothness error', 'compactness error', 'concavity error', 'concave points error', 'symmetry error ', 'fractal dimension error', 'worst radius', 'worst texture', 'worst perimeter', 'worst area', 'worst smoothness', ' worst compactness', 'worst concavity', 'worst concave points', 'worst symmetry', 'worst fractal dimension'] Problem 1. Break the data into training (80%)/testing data(20%). Estimate a tree classification model with maximum depth of 2. Plot the tree and calculate the accuracy rate. Predict target using all features, don't forget to set random numbers to 42. In [58]: # Starting point import random from sklearn.tree import DecisionTreeClassifier from sklearn import tree import pydotplus X = cancer.data y = cancer.target random.seed(42) import os In [59]: from sklearn.model selection import train test split X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42) tree clf = DecisionTreeClassifier(max depth=2, random state=42) tree_clf.fit(X_train, y_train) Out[59]: DecisionTreeClassifier(ccp alpha=0.0, class weight=None, criterion='gini', max depth=2, max features=None, max leaf nodes=None, min_impurity_decrease=0.0, min_impurity_split=None, min_samples_leaf=1, min_samples_split=2, min_weight_fraction_leaf=0.0, presort='deprecated', random state=42, splitter='best') In [60]: from graphviz import Source from sklearn.tree import export_graphviz export_graphviz(tree clf, out file=os.path.join(IMAGES PATH, "cancer tree.dot"), feature names=cancer.feature names, class_names=cancer.target_names, rounded=True, filled=True Source.from_file(os.path.join(IMAGES_PATH, "cancer_tree.dot")) Out[60]: mean concave points <= 0.051 gini = 0.467samples = 455value = [169, 286]class = benign True False worst radius <= 16.83 worst concave points <= 0.147 gini = 0.107gini = 0.204samples = 282samples = 173value = [16, 266]value = [153, 20]class = benign class = malignant gini = 0.037gini = 0.488gini = 0.5gini = 0.029samples = 263samples = 19samples = 35samples = 138value = [5, 258]value = [11, 8]value = [136, 2]value = [17, 18]class = malignant class = malignant class = benign class = benign In [60]: In [61]: from sklearn.metrics import accuracy score y_pred = tree_clf.predict(X_test) acc = accuracy_score(y_test, y_pred) print("Testing accuracy score for a tree with max depth 2 is", acc) Testing accuracy score for a tree with max depth 2 is 0.9298245614035088 Problem 2: Estimate an unrestricted tree on training data and test it on testing data. Find two most important features and create a scatter plot of malignant and benign tumors along the two axes of two most important feature. Hint: For example of a graph look at: https://stackoverflow.com/questions/12487060/matplotlib-color-according-to-class-labels Do you think the data need rotation? In [62]: tree full = DecisionTreeClassifier(random state=42) tree_full.fit(X_train, y_train) sorted feat = tree full.feature importances for name, score in zip(cancer["feature names"], tree full.feature importances): print(name, score) mean radius 0.0 mean texture 0.05847766231107586 mean perimeter 0.0 mean area 0.0 mean smoothness 0.0 mean compactness 0.0 mean concavity 0.0 mean concave points 0.6914195549049809 mean symmetry 0.0 mean fractal dimension 0.0 radius error 0.0 texture error 0.0 perimeter error 0.0 area error 0.011982573676838769 smoothness error 0.0012367800829339453 compactness error 0.0 concavity error 0.0062757755065447375 concave points error 0.015930814747382796 symmetry error 0.0 fractal dimension error 0.018554466715001834 worst radius 0.05229926933685694 worst texture 0.017445161675930944 worst perimeter 0.051493960584869665 worst area 0.0 worst smoothness 0.009233190446208121 worst compactness 0.0 worst concavity 0.0 worst concave points 0.06565079001137543 worst symmetry 0.0 worst fractal dimension 0.0 In [70]: worst_rad = X_train[:,7] worst_conc_per = X_train[:,27] malignancy = y train In [72]: from matplotlib.colors import ListedColormap, LinearSegmentedColormap cmap = ListedColormap(['red','green']) fig = plt.figure(figsize=(8,8)) plt.scatter(worst_rad, worst_conc_per, c=y_train, cmap=cmap) plt.xlabel("mean concave points") plt.ylabel("Worst Concave Points") plt.show() 0.30 0.25 Worst Concave Points 0.05 0.00 0.000 0.025 0.050 0.075 0.100 0.125 0.150 0.175 0.200 mean concave points **Problem 3** Report accuracy using top 2 features on the full data from problem 2. Loop over 10000 random number between -1 and 1 (from -3 to 3 radians) to find an optimal rotation angle. Report accuracy improvelemt over unrotated data. In [77]: np.random.seed(42) # Unrotated accuracy Imp tr = np.column_stack((X_train[:,7],X_train[:,27])) Imp_test = np.column_stack((X_test[:,7],X_test[:,27])) tree clf.fit(Imp tr, y train) y_pred = tree_clf.predict(Imp_test) acc_best = accuracy_score(y_test, y_pred) print(f'Unrotated accuracy is {acc_best}') angle list = np.random.uniform(-3,3,10000)#Imp = np.column_stack((worst_rad, worst_conc_per)) **for** a **in** range(10000): angle = angle list[a] rotation_matrix = np.array([[np.cos(angle), -np.sin(angle)], [np.sin(angle), np.cos(angle)]]) Xr = Imp_tr.dot(rotation_matrix) Xr_test = Imp_test.dot(rotation_matrix) tree_clf.fit(Xr, y_train) y_pred = tree_clf.predict(Xr_test) acc_new = accuracy_score(y_test, y_pred) if acc new > acc best: acc_best = acc_new angle best = angle print(f'best rotated accuracy is {acc} , best angle is {angle_best}') # Create rotaton matrix 2x2 for each point on a 2-dimensional plane rotation_matrix = np.array([[np.cos(angle), -np.sin(angle)], [np.sin(angle), np.cos(angle)]]) Xr = Imp.dot(rotation matrix) colors = ['red','green'] fig = plt.figure(figsize=(8,8)) plt.scatter(Xr[:,0], Xr[:,1], c=y, cmap=ListedColormap(colors)) plt.xlabel("Worst Radius") plt.ylabel("Worst Concave Points") plt.show() Unrotated accuracy is 0.9122807017543859 best rotated accuracy is 0.9298245614035088 , best angle is -1.0919791501688167 0.175 0.150 0.125 Points 0.100 0.025 0.000 -0.30-0.25-0.20-0.150.00 -0.05Worst Radius **Problem 4** In the main data drop variables used in the problem 3. Add instead the rotated variables (substitution). Estimate accuracy score using with a tree classifier with max depth = 2 (Same as in problem 1). How much did we gain from rotation? In [81]: rotation_matrix = np.array([[np.cos(angle_best), -np.sin(angle_best)], [np.sin(angle_best), np.cos(angle_best)]) Xr = Imp tr.dot(rotation matrix) Xr_test = Imp_test.dot(rotation_matrix) X_new_tr = X_train X new test = X test $X_new_tr[:,7] = Xr[:,0]$ $X \text{ new_tr[:,27]} = Xr[:,1]$ $X_{\text{new_test}[:,7]} = Xr_{\text{test}[:,0]}$ $X_{\text{new_test}}[:,27] = Xr_{\text{test}}[:,1]$ In [84]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42) tree clf = DecisionTreeClassifier(max depth=2, random state=42) tree_clf.fit(X_train, y_train) y_pred = tree_clf.predict(X_test) acc_not_rot = accuracy_score(y_test, y_pred) X_new_tr, X_new_test, y_train, y_test = train_test_split(X_new, y, test_size=0.2, random_state=42) tree clf = DecisionTreeClassifier(max depth=2, random state=42) tree_clf.fit(X_new_tr, y_train) y_pred = tree_clf.predict(X_new_test) acc_r = accuracy_score(y_test, y_pred) print("Testing accuracy score for a tree with max depth 2 is", acc_r) gain_acc = acc_r - acc_not_rot print("The accuracy gain from rotation is", gain_acc) Testing accuracy score for a tree with max depth 2 is 0.956140350877193 The accuracy gain from rotation is 0.02631578947368418 **Problem 5** Generate samples of 100, 10,000 and 100,000, moons using the code below. Set random seed at 42. Split data in training and testing sets. Estimate separately Logistic, Random Forest, SVC and the hard voting classifier. What happens to the accuracy score as you increase the number of observations? Measyre and report the time it takes for each estimation. In [85]: from sklearn.model selection import train test split from sklearn.datasets import make moons X, y = make moons(n samples=500, noise=0.30, random state=42)X train, X test, y train, y test = train test split(X, y, random state=42) In [86]: from sklearn.ensemble import RandomForestClassifier from sklearn.ensemble import VotingClassifier from sklearn.linear_model import LogisticRegression from sklearn.svm import SVC log_clf = LogisticRegression(random_state=42) rnd clf = RandomForestClassifier(random state=42) svm clf = SVC(random state=42) # voting classifier has syntaxis akin to pipeline voting clf = VotingClassifier(estimators=[('lr', log_clf), ('rf', rnd_clf), ('svc', svm_clf)], voting='hard') In [89]: from sklearn.metrics import accuracy_score import time for N in [100,10000,100000]: X, y = make_moons(n_samples=N, noise=0.30, random_state=42) X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=42) print(N, "Observations") for clf in (log clf, rnd clf, svm clf, voting clf): start = time.time() clf.fit(X train, y train) y_pred = clf.predict(X_test) print(clf.__class__.__name__, accuracy_score(y_test, y_pred), f'Time: {time.time() - start} ') 100 Observations LogisticRegression 0.96 Time: 0.0045928955078125 RandomForestClassifier 0.92 Time: 0.14564275741577148 SVC 0.92 Time: 0.0022542476654052734 VotingClassifier 0.96 Time: 0.1466691493988037 10000 Observations LogisticRegression 0.8588 Time: 0.017252445220947266 RandomForestClassifier 0.9092 Time: 0.8374078273773193 SVC 0.9184 Time: 0.6378459930419922 VotingClassifier 0.9176 Time: 1.4783236980438232 100000 Observations LogisticRegression 0.85392 Time: 0.12099838256835938 RandomForestClassifier 0.90508 Time: 11.833928108215332 SVC 0.91332 Time: 66.16663932800293 VotingClassifier 0.90976 Time: 78.46461272239685 Answer: As we increase the number of observations the quality of prediction using Logistic goes down, which SVC takes the lead. SVC though is very slow. Though generally Voting classifier performs better it is inferior to SVC with the large number of data points. **Problem 6** Generate data using the code provided below. Using testing accuracy as metric, estimate bagging random trees estimator with 200 estimators. Try different numbers of samples: 10, 30, 50, and 200. What is optimal number of samples to be used? BaggingClassifier(DecisionTreeClassifier(random_state=42), n_estimators=5, max_samples=?, bootstrap=True, n_jobs=-1, random_state=42) In [110]: X, y = make moons(n samples=500, noise=0.40, random state=42) X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=42) In [111]: from sklearn.ensemble import BaggingClassifier from sklearn.tree import DecisionTreeClassifier from sklearn.metrics import accuracy score import numpy as np def bagtree(S): bag clf = BaggingClassifier(DecisionTreeClassifier(random_state=42), n_estimators=200, max samples=S, bootstrap=True, n jobs=-1, random state=42) bag_clf.fit(X_train, y_train) y_pred = bag_clf.predict(X_test) return accuracy_score(y_test, y_pred) $bag_acc = []$ for i in [10, 30, 50,200]: acc = bagtree(i) vec2 = np.column stack((i,acc)) bag_acc.append(vec2) print(acc) print(bag acc) 0.76 0.832 0.848 0.832 [array([[10. , 0.76]]), array([[30. , 0.832]]), array([[50. , 0.848]]), array([[200. , 0.832]])] Answer 10 samples is clearly not enough, but there is little difference between 100, 300 and 1000 samples. **Problem 7** Find optimal learning rate, number of estimators and maximum depth using GradientBoostingClassifier, and Randomize grid search. Set a grid: learning rate from 0.01 to 3, number of estimators from 1 to 20, and maximum depth from 1 to 10. Try 300 iterations. Example for randomizeSearch: rnd search = RandomizedSearchCV(forest reg, param distributions=param distribs, n iter=?, cv=5, scoring='neg mean squared error', random state=42) gbrt = GradientBoostingClassifier(max_depth=?, n_estimators=?, learning_rate = ?, random_state=42) Which estimator was the best? What was the accuracy of the best estimator? In [114]: X, y = make_moons(n samples=2000, noise=0.40, random state=42) In [120]: import random from sklearn.ensemble import GradientBoostingClassifier from sklearn.model selection import RandomizedSearchCV from scipy.stats import randint from scipy.stats import uniform grad = GradientBoostingClassifier(random state=42) param distribs = { 'n estimators': randint(low=1, high=20), 'learning rate': uniform(0.01,3), 'max depth': randint(low=1, high=10), rnd search = RandomizedSearchCV(grad, param distributions=param distribs, n iter=300, cv=5, scoring='neg mean squared error', random_state=42) rnd search.fit(X, y) from sklearn.model selection import cross val score bestest = rnd_search.best_estimator_ scores = np.mean(cross_val_score(bestest, X, y, cv=5)) print("Best estimator", bestest) print("Produced accuracy",scores) Best estimator GradientBoostingClassifier(ccp_alpha=0.0, criterion='friedman_mse', init=None, learning rate=1.0452137440800489, loss='deviance', max depth=1, max features=None, max leaf nodes=None, min impurity decrease=0.0, min impurity split=None, min samples leaf=1, min samples split=2, min_weight_fraction_leaf=0.0, n_estimators=19, n_iter_no_change=None, presort='deprecated', random state=42, subsample=1.0, tol=0.0001, validation_fraction=0.1, verbose=0, warm_start=False)

Produced accuracy 0.864000000000001

In []:

In [49]: # Python ≥3.5 is required

import sklearn

Common imports
import numpy as np

np.random.seed(42)

%matplotlib inline

To plot pretty figures

import matplotlib as mpl

assert sys.version_info >= (3, 5)

Scikit-Learn ≥0.20 is required

assert sklearn.__version__ >= "0.20"

to make this notebook's output stable across runs

import sys

import os