import numpy as np import matplotlib.pyplot as plt from sklearn import svm, datasets from sklearn.model_selection import train_test_split from sklearn.metrics import confusion matrix, ConfusionMatrixDisplay **Exercice I** 1- Lecture des données In [2]: # Chargement des données iris = datasets.load_iris() 2- Partition des données In [3]: # On considère 50% pour l'entrainement et 50%=0.5 pour le test X, y = iris.data[:, :2], iris.target X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.5) class names = iris.target names 3- Affichage de la dispersion des données In [4]: # Sur l'axe des x on aura Sepal length et sur l'axe de y Sepal width # On a trois couleurs pour les trois espèces # et deux marker pour entrainement et test plt.scatter(X_train[:, 0], X_train[:, 1], label="train", edgecolors='k', c=y_train, cmap=plt.cm.coolwarm) plt.scatter(X_test[:, 0], X_test[:, 1], label="test", marker='*', c=y_test, cmap=plt.cm.coolwarm) plt.xlabel('Sepal length') plt.ylabel('Sepal width') plt.title("LinearSVC") plt.show() LinearSVC 4.0 3.5 Sepal width 2.5 2.0 5.0 4.5 7.0 7.5 Sepal length 4- Conception du classificateur In [5]: # On fixe le paramètre de régularisation C=1.0 # On fixe random state à 10 C = 1.0lin_svc = svm.LinearSVC(C=C, random_state =10) lin_svc.fit(X_train, y_train) /Users/nmezghani/opt/anaconda3/lib/python3.8/site-packages/sklearn/svm/_base.py:1206: ConvergenceWarning: Liblinear failed to converge, increa se the number of iterations. warnings.warn(LinearSVC(random_state=10) 5- Taux de classification In [6]: # On teste sur l'ensemble des données de test lin_svc.score(X_test, y_test) Out[6]: 6- Matrice de confusion In [7]: !pip install -U scikit-learn Requirement already satisfied: scikit-learn in /Users/nmezghani/opt/anaconda3/lib/python3.8/site-packages (1.0.2) Requirement already satisfied: numpy>=1.14.6 in /Users/nmezghani/opt/anaconda3/lib/python3.8/site-packages (from scikit-learn) (1.20.3) Requirement already satisfied: scipy>=1.1.0 in /Users/nmezghani/opt/anaconda3/lib/python3.8/site-packages (from scikit-learn) (1.7.3) Requirement already satisfied: threadpoolctl>=2.0.0 in /Users/nmezghani/opt/anaconda3/lib/python3.8/site-packages (from scikit-learn) (2.2.0) Requirement already satisfied: joblib>=0.11 in /Users/nmezghani/opt/anaconda3/lib/python3.8/site-packages (from scikit-learn) (1.1.0) In [8]: # La matrice de confusion peut être non-normalisée # Dans ce cas, on affiche le nombre d'observation # ou normalisée par rapport au nombre d'observation totale # par classe # le paramètre precision permet de fixer le nombre de # chiffre après la virgule np.set printoptions(precision=2) titles options = [("Confusion matrix, without normalization", None), ("Normalized confusion matrix", "true"), for title, normalize in titles options: disp = ConfusionMatrixDisplay.from_estimator(lin_svc, X test, y test, display labels=class names, cmap=plt.cm.Blues, normalize=normalize, disp.ax .set title(title) print(title) print(disp.confusion matrix) plt.show() Confusion matrix, without normalization [[25 0 0] [1 15 10] [0 4 20]] Normalized confusion matrix [[1. 0. 0.] [0.04 0.58 0.38] 0.17 0.83]] .01 Confusion matrix, without normalization 0 0 25 setosa 20 15 Frue label 1 10 versicolor - 10 - 5 20 0 4 virginica setosa versicolor virginica Predicted label Normalized confusion matrix 0 0 setosa 0.8 0.6 0.038 0.58 0.38 versicolor 0.4 - 0.2 0.83 0.17 virginica 0 0.0 setosa versicolor virginica Predicted label 7- Visualisation des frontières de décision In [9]: # Créer la surface de décision discretisée x_{min} , $x_{max} = X[:, 0].min() - 1, X[:, 0].max() + 1$ y_{min} , $y_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1$ # Pour afficher la surface de décision on va discrétiser l'espace avec un pas h $h = max((x_max - x_min) / 100, (y_max - y_min) / 100)$ xx, yy = np.meshgrid(np.arange(x_min, x_max, h), np.arange(y_min, y_max, h)) # Surface de décision linéaire Z = lin svc.predict(np.c_[xx.ravel(), yy.ravel()]) Z = Z.reshape(xx.shape)plt.contourf(xx, yy, Z, cmap=plt.cm.coolwarm, alpha=0.8) # On ajoute les observations utilisées pour d'apprentissage et pour le test également plt.scatter(X_train[:, 0], X_train[:, 1], label="train", edgecolors='k', c=y_train, cmap=plt.cm.coolwarm) plt.scatter(X_test[:, 0], X_test[:, 1], label="test", marker='*', c=y_test, cmap=plt.cm.coolwarm) plt.xlabel('Sepal length') plt.ylabel('Sepal width') plt.title("LinearSVC") plt.show() LinearSVC 5.0 4.5 4.0 Sepal width 3.5 3.0 2.5 2.0 1.5 1.0 6 Sepal length 8- Utilisation des modèles lin_svc et svc In [10]: # Les modèles linéaires LinearSVC() et SVC(kernel='linear') # traitent les problèmes multi-classes de manière différente # linearSVC utilise One-vs-All alors que SVC utilise One-vs-One lin_svc = svm.LinearSVC(C=C).fit(X_train, y_train) svc = svm.SVC(kernel='linear', C=C).fit(X train, y train) titles = ['SVC with linear kernel', 'LinearSVC (linear kernel)'] fig = plt.figure(figsize=(12, 4.5)) for i, clf in enumerate((svc, lin_svc)): plt.subplot(1, 2, i + 1)plt.subplots_adjust(wspace=0.4, hspace=0.4) Z = clf.predict(np.c_[xx.ravel(), yy.ravel()]) # Utiliser une palette de couleurs Z = Z.reshape(xx.shape)plt.contourf(xx, yy, Z, cmap=plt.cm.coolwarm, alpha=0.8) # Afficher aussi les points d'apprentissage plt.scatter(X train[:, 0], X train[:, 1], label="train", edgecolors='k', c=y train, cmap=plt.cm.coolwarm) plt.scatter(X_test[:, 0], X_test[:, 1], label="test", marker='*', c=y_test, cmap=plt.cm.coolwarm) plt.xlabel('Sepal length') plt.ylabel('Sepal width') plt.title(titles[i]) plt.show() /Users/nmezghani/opt/anaconda3/lib/python3.8/site-packages/sklearn/svm/_base.py:1206: ConvergenceWarning: Liblinear failed to converge, increa se the number of iterations. warnings.warn(SVC with linear kernel LinearSVC (linear kernel) 5.0 5.0 4.5 4.5 4.0 4.0 Sepal width Sepal width 3.5 3.5 3.0 3.0 2.5 2.5 2.0 2.0 1.5 1.5 1.0 1.0 6 Sepal length Sepal length 9- Comparaison des résultats Dans les deux cas, on obtient des classifieurs linéaires qui produisent des frontières de décision linéaires. Ces frontières sont légèrement différentes. 10- Utilisation des 4 features In [11]: # On utilise les quatres caractéristiques pour l'entrainement du modèle X, y = iris.data, iris.target X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.5) lin svc = svm.LinearSVC(C=C, random state =10) lin_svc.fit(X_train, y_train) lin_svc.score(X_test, y_test) /Users/nmezghani/opt/anaconda3/lib/python3.8/site-packages/sklearn/svm/ base.py:1206: ConvergenceWarning: Liblinear failed to converge, increa se the number of iterations. warnings.warn(0.92 Out[11]: L'utilisation des quatre caractéristiques permet d'améliorer les taux de classifications (80% par rapport à 92%) **Excercice II** In [12]: from sklearn import tree 1- Partition des données In [13]: # On considère 70% pour l'entrainement et 30%=0.3 pour le test X_train, X_test, y_train, y_test = train_test_split(X, y, train_size=0.7, random_state=0) class_names = iris.target_names 2- Conception d'un arbre de décision de profondeur max de 3 In [14]: clf = tree.DecisionTreeClassifier(max depth = 3) clf.fit(X_train, y_train) DecisionTreeClassifier(max depth=3) Out[14]: In [15]: tree.plot_tree(clf, filled=True) $[Text(0.375, 0.875, 'X[2] \le 2.35 \cdot gini = 0.664 \cdot gngles = 105 \cdot gngles = [34, 32, 39]'),$ Out[15]: $Text(0.25, 0.625, 'gini = 0.0 \setminus samples = 34 \setminus value = [34, 0, 0]'),$ $Text(0.5, 0.625, 'X[2] \le 4.95 \cdot = 0.495 \cdot = 71 \cdot = [0, 32, 39]'),$ $Text(0.25, 0.375, 'X[3] \le 1.65 \cdot ngini = 0.161 \cdot nsamples = 34 \cdot nvalue = [0, 31, 3]'),$ $Text(0.125, 0.125, 'gini = 0.0 \setminus samples = 30 \setminus value = [0, 30, 0]'),$ $Text(0.375, 0.125, 'gini = 0.375 \setminus samples = 4 \setminus value = [0, 1, 3]'),$ $Text(0.75, 0.375, 'X[2] \le 5.05 \cdot gini = 0.053 \cdot gini = 37 \cdot gini$ $Text(0.625, 0.125, 'gini = 0.375 \setminus samples = 4 \setminus value = [0, 1, 3]'),$ Text(0.875, 0.125, 'gini = 0.0\nsamples = 33\nvalue = [0, 0, 33]')] X[2] <= 2.35 gini = 0.664 samples = 105 value = [34, 32, 39] X[2] <= 4.95 ginl = 0.495 samples = 71 value = [0, 32, 39] 3- Taux de classification In [16]: clf.predict(X test) array([2, 1, 0, 2, 0, 2, 0, 1, 1, 1, 2, 1, 1, 1, 1, 0, 1, 1, 0, 0, 2, 1, Out[16]: 0, 0, 2, 0, 0, 1, 1, 0, 2, 1, 0, 2, 2, 1, 0, 2, 1, 1, 2, 0, 2, 0, 0]) In [17]: # Taux de classification clf.score(X test, y test) 0.977777777777777 Out[17]: Matrice de confusion In [18]: disp = ConfusionMatrixDisplay.from estimator(clf, X_test, y test, display labels=class names, cmap=plt.cm.Blues, disp.ax_.set_title(title) print(title) print(disp.confusion matrix) plt.show() Normalized confusion matrix [[16 0 0] [0 17 1] [0 0 11]] Normalized confusion matrix - 16 0 setosa - 14 - 12 - 10 True label 0 17 1 versicolor - 8 6 0 11 virginica - 2 setosa versicolor virginica Predicted label 4- Conception d'un arbre de décision avec un nombre minimum d'échantillons requis par feuille égal à 20. In [19]: clf = tree.DecisionTreeClassifier(min samples leaf = 20) clf.fit(X train, y train) DecisionTreeClassifier(min_samples_leaf=20) Out[19]: In [20]: tree.plot tree(clf, filled=True) $[Text(0.4, 0.83333333333333333, 'X[2] \le 2.35 \cdot ngini = 0.664 \cdot nsamples = 105 \cdot nvalue = [34, 32, 39]'),$ Out[20]: $Text(0.2, 0.5, 'gini = 0.0 \setminus samples = 34 \setminus value = [34, 0, 0]'),$ $Text(0.6, 0.5, 'X[2] \le 4.95 \cdot = 0.495 \cdot = 71 \cdot = [0, 32, 39]'),$ $X[2] \le 2.35$ gini = 0.664samples = 105 value = [34, 32, 39] $X[2] \le 4.95$ gini = 0.0gini = 0.495samples = 34samples = 71value = [34, 0, 0] value = [0, 32, 39] gini = 0.161gini = 0.053samples = 34 samples = 37 value = [0, 31, 3]value = [0, 1, 36] 5- Taux de classification In [21]: clf.score(X test, y test) 0.9111111111111111 Out [21]: Matrice de confusion In [22]: disp = ConfusionMatrixDisplay.from estimator(clf, X_test, y_test, display labels=class names, cmap=plt.cm.Blues, disp.ax_.set_title(title) print(title) print(disp.confusion matrix) plt.show() Normalized confusion matrix [[16 0 0] [0 17 1] [0 3 8]] Normalized confusion matrix - 16 16 0 setosa - 14 - 12 - 10 Frue label 17 0 1 versicolor 8 6 - 4 0 3 virginica - 2 setosa versicolor virginica Predicted label Le taux de classification global est 91%. Le taux de classification de la classe Setosa est de 16/16=100%, celui de la classe virginica est de 8/11=72% 6 & 7- Calcul du taux de classification en fonction des paramètres mdepth et msplit In [23]: from sklearn import model_selection X train, X test, y train, y test = model selection.train test split(iris.data, iris.target, test_size=0.95, random_state=0) for mdepth in [1, 2, 3, 4, 5, 6, 7]: clf = tree.DecisionTreeClassifier(max_depth=mdepth) clf = clf.fit(X_train, y_train) print(clf.score(X_test, y_test)) for msplit in [2, 3, 5, 10, 15, 20]: clf = tree.DecisionTreeClassifier(min samples split=msplit) clf = clf.fit(X train, y train) print(clf.score(X_test, y_test)) 0.6573426573426573 0.7972027972027972 0.8881118881118881 0.8881118881118881 0.7062937062937062 0.7972027972027972 0.7972027972027972 0.7062937062937062 0.8881118881118881 0.6573426573426573 0.32167832167832167 0.32167832167832167 0.32167832167832167 8- Optimisation des paramètres mdepth et msplit In [24]: from sklearn import model selection from sklearn.tree import DecisionTreeClassifier from sklearn.model selection import GridSearchCV In [25]: X_train, X_test, y_train, y_test = model_selection.train_test_split(iris.data, iris.target, test size=0.30, random state=0) pgrid = {"max_depth": [1, 2, 3, 4, 5, 6, 7], "min samples split": [2, 3, 5, 10, 15, 20]} grid search = GridSearchCV(DecisionTreeClassifier(), param grid=pgrid, cv=10) grid_search.fit(X_train, y_train) grid search.best estimator .score(X test, y test) print("Best parameters: ", grid_search.best_params_) Best parameters: {'max depth': 3, 'min samples split': 2} In []: In []: