You are currently looking at version 1.0 of this notebook. To download notebooks and datafiles, as well as get help on Jupyter notebooks in the Coursera platform, visit the <u>Jupyter Notebook FAQ</u> course resource. Applied Machine Learning: Module 4 (Supervised Learning, Part Preamble and Datasets In [4]: %matplotlib notebook import numpy as np import pandas as pd import seaborn as sn import matplotlib.pyplot as plt from sklearn.model_selection import train test split from sklearn.datasets import make classification, make blobs from matplotlib.colors import ListedColormap from sklearn.datasets import load breast cancer from adspy shared utilities import load crime dataset cmap bold = ListedColormap(['#FFFF00', '#00FF00', '#0000FF','#000000']) # fruits dataset fruits = pd.read_table('readonly/fruit_data_with_colors.txt') feature names fruits = ['height', 'width', 'mass', 'color score'] X fruits = fruits[feature names fruits] y fruits = fruits['fruit label'] target names fruits = ['apple', 'mandarin', 'orange', 'lemon'] X fruits 2d = fruits[['height', 'width']] y_fruits_2d = fruits['fruit_label'] # synthetic dataset for simple regression from sklearn.datasets import make_regression plt.figure() plt.title('Sample regression problem with one input variable') X_R1, y_R1 = make_regression(n_samples = 100, n features=1, n informative=1, bias = 150.0, noise = 30, random state=0) plt.scatter(X_R1, y_R1, marker= 'o', s=50) plt.show() # synthetic dataset for more complex regression from sklearn.datasets import make_friedman1 plt.figure() plt.title('Complex regression problem with one input variable') X_F1, y_F1 = make_friedman1(n_samples = 100, n_features = 7, random_state=0) plt.scatter(X_F1[:, 2], y_F1, marker= 'o', s=50) plt.show() # synthetic dataset for classification (binary) plt.figure() plt.title('Sample binary classification problem with two informative features') X_C2, y_C2 = make_classification(n_samples = 100, n_features=2, n_redundant=0, n_informative=2, n clusters per class=1, flip y = 0.1, class sep = 0.5, random state=0) plt.scatter(X_C2[:, 0], X_C2[:, 1], marker= 'o', c=y_C2, s=50, cmap=cmap_bold) plt.show() # more difficult synthetic dataset for classification (binary) # with classes that are not linearly separable X_D2, y_D2 = make_blobs(n_samples = 100, n_features = 2, centers = 8, cluster_std = 1.3, $random_state = 4)$ $y_D2 = y_D2 \% 2$ plt.figure() plt.title('Sample binary classification problem with non-linearly separable classes') plt.scatter(X D2[:,0], X D2[:,1], c=y D2, marker= 'o', s=50, cmap=cmap_bold) plt.show() # Breast cancer dataset for classification cancer = load breast cancer() (X_cancer, y_cancer) = load_breast_cancer(return_X_y = **True**) # Communities and Crime dataset (X_crime, y_crime) = load_crime_dataset() FileNotFoundError Traceback (most recent call last) <ipython-input-4-9671647cd5a5> in <module>() 74 # Communities and Crime dataset ---> 75 (X_crime, y_crime) = load_crime_dataset() /home/jovyan/work/adspy_shared_utilities.py in load_crime_dataset() # https://archive.ics.uci.edu/ml/datasets/Communities+and+Crime+Unnormalized 18 **--->** 19 crime = pd.read_table('readonly/CommViolPredUnnormalizedData.txt', sep=',', na_value s='?') 20 # remove features with poor coverage or lower relevance, and keep ViolentCrimesPerPo p target column $columns_{to} = [5, 6] + list(range(11, 26)) + list(range(32, 103)) + [145]$ /opt/conda/lib/python3.6/site-packages/pandas/io/parsers.py in parser_f(filepath_or_buffer, sep, delimiter, header, names, index_col, usecols, squeeze, prefix, mangle_dupe_cols, dtype, engine, converters, true_values, false_values, skipinitialspace, skiprows, nrows, na_values, keep_defau lt_na, na_filter, verbose, skip_blank_lines, parse_dates, infer_datetime_format, keep_date_col, date parser, dayfirst, iterator, chunksize, compression, thousands, decimal, lineterminator, qu otechar, quoting, escapechar, comment, encoding, dialect, tupleize_cols, error_bad_lines, warn_b ad_lines, skipfooter, skip_footer, doublequote, delim_whitespace, as_recarray, compact_ints, use _unsigned, low_memory, buffer_lines, memory_map, float_precision) 644 skip_blank_lines=skip_blank_lines) 645 --> 646 return _read(filepath_or_buffer, kwds) 647 648 parser_f.__name__ = name /opt/conda/lib/python3.6/site-packages/pandas/io/parsers.py in read(filepath or buffer, kwds) 387 388 # Create the parser. --> 389 parser = TextFileReader(filepath_or_buffer, **kwds) 390 391 if (nrows is not None) and (chunksize is not None): /opt/conda/lib/python3.6/site-packages/pandas/io/parsers.py in init (self, f, engine, **kwds) 728 self.options['has_index_names'] = kwds['has_index_names'] 729 -**-**> 730 self. make engine(self.engine) 731 def close(self): 732 /opt/conda/lib/python3.6/site-packages/pandas/io/parsers.py in _make_engine(self, engine) def make engine(self, engine='c'): if engine == 'c': self._engine = CParserWrapper(self.f, **self.options) 924 925 if engine == 'python': /opt/conda/lib/python3.6/site-packages/pandas/io/parsers.py in __init__(self, src, **kwds) kwds['allow_leading_cols'] = self.index_col is not False 1389 -> 1390 self._reader = _parser.TextReader(src, **kwds) 1391 1392 # XXX pandas/parser.pyx in pandas.parser.TextReader.__cinit__ (pandas/parser.c:4184) () pandas/parser.pyx in pandas.parser.TextReader. setup parser source (pandas/parser.c:8449)() FileNotFoundError: File b'CommViolPredUnnormalizedData.txt' does not exist Naive Bayes classifiers In []: from sklearn.naive bayes import GaussianNB from adspy_shared_utilities import plot class regions for classifier X_train, X_test, y_train, y_test = train_test_split(X_C2, y_C2, random_state=0) nbclf = GaussianNB().fit(X train, y train) plot_class_regions_for_classifier(nbclf, X_train, y_train, X_test, y_test, 'Gaussian Naive Bayes classifier: Dataset 1') In []: X_train, X_test, y_train, y_test = train_test_split(X_D2, y_D2, random state=0) nbclf = GaussianNB().fit(X train, y train) plot_class_regions_for_classifier(nbclf, X_train, y_train, X_test, y_test, 'Gaussian Naive Bayes classifier: Dataset 2') Application to a real-world dataset In []: X_train, X_test, y_train, y_test = train_test_split(X_cancer, y_cancer, random_state = 0) nbclf = GaussianNB().fit(X train, y train) print('Breast cancer dataset') print('Accuracy of GaussianNB classifier on training set: {:.2f}' .format(nbclf.score(X train, y train))) print('Accuracy of GaussianNB classifier on test set: {:.2f}' .format(nbclf.score(X_test, y_test))) **Ensembles of Decision Trees** Random forests In []: from sklearn.ensemble import RandomForestClassifier from sklearn.model_selection import train test split from adspy_shared_utilities import plot_class_regions_for_classifier_subplot X_train, X_test, y_train, y_test = train_test_split(X_D2, y_D2, fig, subaxes = plt.subplots(1, 1, figsize=(6, 6)) clf = RandomForestClassifier().fit(X train, y train) title = 'Random Forest Classifier, complex binary dataset, default settings' plot_class_regions_for_classifier_subplot(clf, X_train, y_train, X_test, y_test, title, subaxes) plt.show() Random forest: Fruit dataset In []: from sklearn.ensemble import RandomForestClassifier from sklearn.model_selection import train_test_split from adspy_shared_utilities import plot_class_regions_for_classifier_subplot X_train, X_test, y_train, y_test = train_test_split(X_fruits.as_matrix(), y fruits.as matrix(), random state = 0)fig, subaxes = plt.subplots(6, 1, figsize=(6, 32)) title = 'Random Forest, fruits dataset, default settings' $pair_list = [[0,1], [0,2], [0,3], [1,2], [1,3], [2,3]]$ for pair, axis in zip(pair_list, subaxes): X = X_train[:, pair] y = y_train clf = RandomForestClassifier().fit(X, y) plot_class_regions_for_classifier_subplot(clf, X, y, None, None, title, axis, target names fruits) axis.set_xlabel(feature_names_fruits[pair[0]]) axis.set ylabel(feature names fruits[pair[1]]) plt.tight layout() plt.show() clf = RandomForestClassifier(n estimators = 10, random_state=0).fit(X_train, y_train) print('Random Forest, Fruit dataset, default settings') print('Accuracy of RF classifier on training set: {:.2f}' .format(clf.score(X_train, y_train))) print('Accuracy of RF classifier on test set: {:.2f}' .format(clf.score(X test, y test))) Random Forests on a real-world dataset In []: from sklearn.ensemble import RandomForestClassifier X_train, X_test, y_train, y_test = train_test_split(X_cancer, y cancer, random state = 0) clf = RandomForestClassifier(max features = 8, random state = 0) clf.fit(X_train, y_train) print('Breast cancer dataset') print('Accuracy of RF classifier on training set: {:.2f}' .format(clf.score(X train, y train))) print('Accuracy of RF classifier on test set: {:.2f}' .format(clf.score(X test, y_test))) **Gradient-boosted decision trees** In []: from sklearn.ensemble import GradientBoostingClassifier from sklearn.model_selection import train_test_split from adspy_shared_utilities import plot_class_regions_for_classifier_subplot X_train, X_test, y_train, y_test = train_test_split(X_D2, y_D2, random_state = 0) fig, subaxes = plt.subplots(1, 1, figsize=(6, 6)) clf = GradientBoostingClassifier().fit(X_train, y_train) title = 'GBDT, complex binary dataset, default settings' plot_class_regions_for_classifier_subplot(clf, X_train, y_train, X_test, y_test, title, subaxes) plt.show() Gradient boosted decision trees on the fruit dataset In []: | X_train, X_test, y_train, y_test = train_test_split(X_fruits.as_matrix(), y_fruits.as_matrix(), random_state = 0) fig, subaxes = plt.subplots(6, 1, figsize=(6, 32)) $pair_list = [[0,1], [0,2], [0,3], [1,2], [1,3], [2,3]]$ for pair, axis in zip(pair_list, subaxes): X = X train[:, pair] $y = y_{train}$ clf = GradientBoostingClassifier().fit(X, y) plot class regions for classifier subplot(clf, X, y, None, None, title, axis, target_names_fruits) axis.set_xlabel(feature_names_fruits[pair[0]]) axis.set_ylabel(feature_names_fruits[pair[1]]) plt.tight_layout() plt.show() clf = GradientBoostingClassifier().fit(X train, y train) print('GBDT, Fruit dataset, default settings') print('Accuracy of GBDT classifier on training set: {:.2f}' .format(clf.score(X train, y train))) print('Accuracy of GBDT classifier on test set: {:.2f}' .format(clf.score(X_test, y_test))) Gradient-boosted decision trees on a real-world dataset In []: from sklearn.ensemble import GradientBoostingClassifier X_train, X_test, y_train, y_test = train_test_split(X_cancer, y_cancer, random_state = 0) clf = GradientBoostingClassifier(random state = 0) clf.fit(X_train, y_train) print('Breast cancer dataset (learning rate=0.1, max depth=3)') print('Accuracy of GBDT classifier on training set: {:.2f}' .format(clf.score(X_train, y_train))) print('Accuracy of GBDT classifier on test set: {:.2f}\n' .format(clf.score(X_test, y_test))) clf = GradientBoostingClassifier(learning_rate = 0.01, max_depth = 2, random_state = 0) clf.fit(X_train, y_train) print('Breast cancer dataset (learning rate=0.01, max depth=2)') print('Accuracy of GBDT classifier on training set: {:.2f}' .format(clf.score(X train, y train))) print('Accuracy of GBDT classifier on test set: {:.2f}' .format(clf.score(X_test, y_test))) **Neural networks Activation functions** In []: xrange = np.linspace(-2, 2, 200) plt.figure(figsize=(7,6)) plt.plot(xrange, np.maximum(xrange, 0), label = 'relu') plt.plot(xrange, np.tanh(xrange), label = 'tanh') plt.plot(xrange, 1 / (1 + np.exp(-xrange)), label = 'logistic') plt.legend() plt.title('Neural network activation functions') plt.xlabel('Input value (x)') plt.ylabel('Activation function output') plt.show() **Neural networks: Classification** Synthetic dataset 1: single hidden layer In []: from sklearn.neural network import MLPClassifier from adspy_shared_utilities import plot_class_regions_for_classifier_subplot X_train, X_test, y_train, y_test = train_test_split(X_D2, y_D2, random_state=0) fig, subaxes = plt.subplots(3, 1, figsize=(6,18)) for units, axis in zip([1, 10, 100], subaxes): nnclf = MLPClassifier(hidden layer sizes = [units], solver='lbfgs', random_state = 0).fit(X_train, y_train) title = 'Dataset 1: Neural net classifier, 1 layer, {} units'.format(units) plot class regions for classifier_subplot(nnclf, X_train, y_train, X_test, y_test, title, axis) plt.tight layout() Synthetic dataset 1: two hidden layers In []: from adspy shared utilities import plot class regions for classifier X_train, X_test, y_train, y_test = train_test_split(X_D2, y_D2, random_state=0) nnclf = MLPClassifier(hidden layer sizes = [10, 10], solver='lbfgs', random_state = 0).fit(X_train, y_train) plot class regions for classifier(nnclf, X train, y train, X test, y test, 'Dataset 1: Neural net classifier, 2 layers, 10/10 units') Regularization parameter: alpha In []: X_train, X_test, y_train, y_test = train_test_split(X_D2, y_D2, random_state=0) fig, subaxes = plt.subplots(4, 1, figsize=(6, 23)) for this alpha, axis in zip([0.01, 0.1, 1.0, 5.0], subaxes): nnclf = MLPClassifier(solver='lbfgs', activation = 'tanh', alpha = this alpha, hidden layer_sizes = [100, 100], random_state = 0).fit(X_train, y_train) title = 'Dataset 2: NN classifier, alpha = {:.3f} '.format(this alpha) plot class regions for classifier subplot(nnclf, X train, y train, X test, y test, title, axis) plt.tight_layout()

The effect of different choices of activation function

activation function'.format(this activation)

plt.tight layout()

Neural networks: Regression

In []: from sklearn.neural_network import MLPRegressor

fig, subaxes = plt.subplots(3, 1, figsize=(6,18))

In []: X train, X test, y train, y test = train test split(X D2, y D2, random state=0)

for this_activation, axis in zip(['logistic', 'tanh', 'relu'], subaxes):
 nnclf = MLPClassifier(solver='lbfgs', activation = this activation,

plot class regions for classifier subplot(nnclf, X train, y train,

title = 'Dataset 2: NN classifier, 2 layers 10/10, {} \

fig, subaxes = plt.subplots(2, 3, figsize=(11,8), dpi=70)

for thisaxisrow, thisactivation in zip(subaxes, ['tanh', 'relu']):

y predict output = mlpreg.predict(X predict input)

clf = MLPClassifier(hidden layer sizes = [100, 100], alpha = 5.0,

print('Accuracy of NN classifier on training set: {:.2f}'

for thisalpha, thisaxis in zip([0.0001, 1.0, 100], thisaxisrow):
 mlpreg = MLPRegressor(hidden layer sizes = [100,100],

alpha = thisalpha,

thisaxis.set title('MLP regression\nalpha={}, activation={})'

.format(thisalpha, thisactivation))

X_train, X_test, y_train, y_test = train_test_split(X_cancer, y_cancer, random_state = 0)

random state = 0, solver='lbfgs').fit(X train scaled, y train)

X predict input = np.linspace(-3, 3, 50).reshape(-1,1)

thisaxis.set xlim([-2.5, 0.75])

plt.tight_layout()

scaler = MinMaxScaler()

Application to real-world dataset for classification

from sklearn.preprocessing import MinMaxScaler

X train scaled = scaler.fit transform(X train)

In []: from sklearn.neural network import MLPClassifier

X test scaled = scaler.transform(X test)

print('Breast cancer dataset')

thisaxis.plot(X_train, y_train, 'o')
thisaxis.set_xlabel('Input feature')
thisaxis.set_ylabel('Target value')

alpha = 0.1, hidden_layer_sizes = [10, 10],
random state = 0).fit(X train, y train)

X_train, X_test, y_train, y_test = train_test_split(X_R1[0::5], y_R1[0::5], random_state = 0)

solver = 'lbfgs').fit(X train, y train)

activation = thisactivation,

X test, y test, title, axis)