• Linear Regression • Multiple Linear Regression • Polynomial Linear Regression • Support Vector Regression Decision Tree Regression · Random Forest Regression Clustering • K-Means Clustering · Hierarchical Clustering process · first we import all the libraries · than we read our data • than we do some data exploration before the analysis just to understand the data best . so if we fuond some null values or if we want to chane them than we can do that • we can change the data types of te data • so there are some data exploration commands • info, head, tail, isnull, notnull, unique, nunqiue, dtypes, inhdex, columns, counts, value_counts and mamy more • now we wil check the corelation b/w the dqata and present that data on the heatmap by sns function · noow we can drop thos ecolumns that aere not usefull for our data • and we check corelation so if theer are some features that are highly sorelated than we can drop one of thattwo nfeaturse now its good two select the data and splity that data into input and the out put like x and y · now if trhe data is categorical tan we can do lablel encoding otherwise dont • before the lable encoding we select thge xtreain and y train variables and select the random state and test size than we do lable encoding • now after lable encoding we do normalization • and if the data is not in 2d state than we reeshaoe the data and then do normalization • now after fit and transform the normalization to our xtrain and xtest data we will go for the model selection EDA(exploratry data analysis) import pandas as pd import numpy as np import matplotlib.pyplot as plt import seaborn as sns df.info() df.describe() df.isnull() df.corr() df.dtypes df.shape df.columns df.index sns.countplot(df['c_name']) plt.show() sns.countplot(df.diagnosis) plt.show() sns.heatmap(df.c_name plt.plot(df['c_name1'], df['c_name2']) plt.figure(figsize=(15,9)) plt.scatter(df.c_name1, df.c_name2) plt.show() plt.figure(figsize=(15,9)) sns.jointplot(df.c_name1, df.c_name2) plt.show() sns.pairplot(df[['c_name1', 'c_name2', 'c_name3', 'c_name4']]) plt.show() df[['c_name1', 'c_name2', 'c_name3', 'c_name4']].plot() prepocessing In []: 1 = [] for i in df['diagnosis']: **if** i **==**'M': 1.append(1) else: 1.append(0) col= ['diagnosis'] for i in col: df[i] = le.fit_transform(df[i]) from sklearn.preprocessing import LabelEncoder le = LabelEncoder() # select x and y from sklearn.model_selection import train_test_split x_train, x_test, y_train, y_test = train_test_split(x,y, test_size= 0.3, random_state=41) from sklearn.preprocessing import MinMaxScaler scaler = MinMaxScaler() # SO HERE WE GOT ONE D ARRAY but for the normalization wneed the 2d arrayso we reshace our data x = data.c_name1.values.reshape(-1,1) # or we can do this also y = data.c_name2.values.reshape(-1,1) $x_{train} = np.array(x_{train}).reshape(-1,1)$ $x_{test} = np.array(x_{test}).reshape(-1,1)$ scaler.fit(x_train) x_train = scaler.transform(x_train) x_test = scaler.transform(x_test) use in every model # y_test y_pred for all from sklearn.metrics import confusion_matrix from sklearn.metrics import accuracy_score from sklearn.metrics import precision_score from sklearn.metrics import recall_score from sklearn.metrics import f1_score from sklearn.metrics import classification_report 1. linear regresion (only two features) • multyple regression is same in which we use more then two features as input # now we will import th modle to fit our data from sklearn.linear_model import LinearRegression lr = LinearRegression() lr.fit(x_train, y_train) lr.coef_ # it will give eus the value of m which is slope lr.intercept_ # it will give us the value of c which is intercept # now we have x value m and c now we have to do is # y = mx + c so all we have now we can pridict the y values y_pred = lr.predict(x_test) plt.scatter(x_test, y_test, label = 'baby', c ='r') plt.plot(x_test, y_pred, label= 'hubby', c= 'g') plt.title('graph final') plt.legend() plt.show() # now we can also check the various errors so we can that too from sklearn.metrics import mean_squared_error np.sqrt(mean_squared_error(y_test, y_pred)) # we hav e a command r2score to check the accuracy from sklearn.metrics import r2_score r2_score(y_test, y_pred) print('R sq: ', linear_reg.score(x, y)) print('Correlation: ', math.sqrt(linear_reg.score(x, y))) # we can use tis also **Polynomial Regression** • same as linear regresion from sklearn.preprocessing import PolynomialFeatures from sklearn.linear_model import LinearRegression polynominal_regression = PolynomialFeatures(degree=4) x_polynomial = polynominal_regression.fit_transform(x,y) # %% fit linear_regression = LinearRegression() linear_regression.fit(x_polynomial,y) y_head2 = linear_regression.predict(x_polynomial) from sklearn.metrics import r2_score print("r_square score: ", r2_score(y,y_head2)) Logistic regression • knn and logi we can use both for same prob from sklearn.linear_model import LogisticRegression lgr = LogisticRegression() lgr.fit(x_train, y_train) y_pred = lgr.predict(x_test) # y_test y_pred for all from sklearn.metrics import confusion_matrix from sklearn.metrics import accuracy_score from sklearn.metrics import precision_score from sklearn.metrics import recall_score from sklearn.metrics import f1_score from sklearn.metrics import classification_report knn from sklearn.neighbors import KNeighborsClassifier knn = KNeighborsClassifier(n_neighbors=3) knn.fit(x_train, y_train) y_pred = knn.predict(x_test) # y_test y_pred for all from sklearn.metrics import confusion_matrix from sklearn.metrics import accuracy_score from sklearn.metrics import precision_score from sklearn.metrics import recall_score from sklearn.metrics import f1_score from sklearn.metrics import classification_report confusion_matrix(y_test, prediction) Naive Bayes Classification from sklearn.model_selection import train_test_split x_train, x_test, y_train, y_test = train_test_split(x,y,test_size = 0.3,random_state=1) # %% Naive bayes from sklearn.naive_bayes import GaussianNB nb = GaussianNB()nb.fit(x_train,y_train) nb.score(x_test,y_test) #%% confusion matrix $y_pred = nb.predict(x_test)$ y_true = y_test from sklearn.metrics import confusion_matrix cm = confusion_matrix(y_true,y_pred) # %% cm visualization import seaborn as sns f, ax = plt.subplots(figsize =(5,5)) sns.heatmap(cm, annot = True, linewidths=0.5, linecolor="red", fmt = ".0f", ax=ax) plt.xlabel("y_pred") plt.ylabel("y_true") plt.show() decission tree · everry thing is same in this model to used for categorical data from sklearn.tree import DecisionTreeClassifier dtree = DecisionTreeClassifier() dtree.fit(x_train,y_train) dtree.predict(x_test) Random forest from sklearn.ensemble import RndomForestClassifier rf = RndomForestClassifier(n_estimators = 51) rf.fit(x_train,y_train) rf.predict(x_test) **SVM** two types 1.linear from sklearn.svm import SVC svm = SVC(random_state = 1) svm.fit(x_train,y_train) from sklearn.svm import SVC svm = SVC(kernel = 'linear') svm.fit(x_train,y_train) 2.kerneled svm from sklearn.preprocessing import StandardScaler sc1 = StandardScaler() x_olcekli = sc1.fit_transform(x) sc2 = StandardScaler() y_olcekli = sc2.fit_transform(y) from sklearn.svm import SVR

svr_reg = SVR(kernel = 'rbf')
svr_reg.fit(x_olcekli,y_olcekli)

y_head = svr_reg.predict(x_olcekli)

how to apply models in different conditions

Classification

Regression

Naive Bayes ClassificationDecision Tree ClassificationRandom Forest Classification

K-Nearest Neighbour (KNN) ClassificationSupport Vector Machine (SVM) Classification