In [19]: import numpy as np from numpy.linalg import eig import matplotlib.pyplot as plt from scipy.stats import norm import matplotlib.ticker as ticker import statsmodels.formula.api as smf from scipy.stats import linregress from scipy import stats from skimage.measure import LineModelND, ransac import random as rand from numpy.linalg import inv import math from numpy.random import randn ##2b matrix1 = np.array([1, -0.5])matrix2 = np.array([-0.5, 0.5])#find stdev of matricies sdev1 = np.std(matrix1) sdev2 = np.std(matrix2)#mean for both stdev of matrix matrix no t = np.array([1,1])matrix t = np.transpose(matrix\_no\_t) mean = np.mean(matrix\_t) #Generate 1000 random normally distributed samples of [x1,x2]x1 = np.random.normal(loc = mean, scale = sdev1, size = 1000) x2 = np.random.normal(loc = mean, scale = sdev2, size = 1000)#plot x1 and x2 in scatter plot plt.scatter(x1, x2) #superimposing eigenvectors #finding eigenvectors of matrix1 matrix comb = np.array([[1, -0.5], [-0.5, 0.5]])eigenv = eig(matrix comb) #plotting 3 eigenvectors plt.plot([0, eigenv[0][0]], [0, eigenv[0][1]], color = "r", linewidth = 3, label = 'eigenvector 1') plt.plot([0, eigenv[1][0][0]], [0, eigenv[1][0][1]], color = "b", linewidth = 3, label = 'eigenvector 2') plt.plot([0, eigenv[1][1][0]], [0, eigenv[1][1][1]]], color = "y", linewidth = 3, label = 'eigenvector 3') #superimpose PDF level curve ... x1n2 = x1 + x2p\_x = x1 p\_y = x2 sdev = np.std(matrix comb) x mean = sum(x1n2) / len(x1n2)#doing PDF math based on given formula (searched it online) frac = 1/np.sqrt(2 \* math.pi \* sdev) frac in exp = (-(x1n2 - x mean)\*\*2)/(2\*sdev\*\*2)exp = np.exp(frac in exp) p z = frac \* exp #build level curve plt.tricontour(p\_x, p\_y, p\_z) plt.title('Normal Distribution with PDF Level Curves Superimposed and Eigenvectors Graphed') plt.xlabel('x1') plt.ylabel('x2') plt.legend() plt.show() Normal Distribution with PDF Level Curves Superimposed and Eigenvectors Graphed 2.0 1.5 1.0  $\approx$ 0.5 0.0 eigenvector 1 eigenvector 2 -0.5eigenvector 3 #2c #calculate Euclidean distance #calculate mean of x1 and x2 x1n2 = x1 + x2 $x_{mean} = sum(x1n2) / len(x1n2)$ #euclidean distance e x = x1 $e_y = x2$ #doing euclidean distance math based on given formula  $e z = abs(x1n2 - x_mean)$ #build level curve plt.tricontour(e\_x, e\_y, e\_z) plt.scatter(e\_x, e\_y) plt.title('Normal Distribution with Euclidean Distance Level Curves Superimposed') plt.xlabel('x1') plt.ylabel('x2') plt.show() Normal Distribution with Euclidean Distance Level Curves Superimposed 2.0 1.5 1.0 0.5 0.0 -0.5-1 0 1 2 x1 #calculate Mahalanobis distance #inverse matrix inv mat = inv(matrix comb) #standard deviation sdev\_inv = np.std(inv\_mat) #calculate x - mean x1n2 = x1 + x2x mean = sum(x1n2) / len(x1n2)sub x mean = x1n2 - x mean#Mahalanobis distance (z)  $m \times = \times 1$ m y = x2#doing Mahalanobis distance math based on given formula m\_z = np.sqrt(sub\_x\_mean \* sdev\_inv \* sub\_x\_mean) #build level curve plt.tricontour(m\_x, m\_y, m\_z) plt.scatter(m\_x, m\_y) plt.title('Normal Distribution with Mahalanobis Distance Level Curves Superimposed') plt.xlabel('x1') plt.ylabel('x2') plt.show() Normal Distribution with Mahalanobis Distance Level Curves Superimposed 2.0 1.5 1.0 0.5 0.0 -0.5-1 x1 from sklearn.linear model import Lasso from sklearn.model selection import train\_test\_split from scipy.io import loadmat import seaborn as sns #Grad Descent Algo #loss function def Loss(y, ypred): l=(y-ypred)\*\*2return(1.sum()) #gradient descent function def GradDesc(X, Y, learning\_rate, trials, regularization): global cacheLoss cacheLoss=[None] \* trials Weights=np.random.rand(X.shape[1]) Weights=np.array(Weights) Weights=Weights.reshape(-1,1) m=X.shape[0]for i in range(trials): predictions=np.matmul(X, Weights) cacheLoss[i]=Loss(Y,predictions)  $\label{eq:weights[0]-weights[0]-(1/m)*learning rate*(np.matmul(X[:,0].transpose(),predictions-Y))} \\$ for j in range(1,len(Weights)): return (Weights) #load in samples hw3 = loadmat('HW1 3.mat') x train = hw3['X']y\_train = hw3['Y'] x test = hw3['X test'] y test = hw3['Y test'] #turn into numpy x train = np.array(x train) y\_train = np.array(y\_train) x test = np.array(x test) y test = np.array(y test) #Lasso to generate lambda (lowest MSE on data) bestMSE = 10e100alphaList L = [1\*0.1 for 1 in range(1,102)]for i in alphaList L: lassoModel = Lasso (alpha = i, max iter = 5000, fit intercept = False) lassoModel.fit(x\_train,y\_train) getPred = lassoModel.predict(x train).reshape(-1,1) MSE = sum((y test-getPred)\*\*2) if MSE < bestMSE:</pre> bestMSE = MSE lassoLambda = iprint(f'Ideal lambda is {lassoLambda}') #get coefficients with Gradient Descent with learning rate of 5e-3 and 10000 gradient updates and LASSO regular addBias=np.ones([x train.shape[0],1]) x train=np.append(addBias,x train,axis=1) coefficients L = GradDesc(x train, y train, learning rate = 5e-3, trials = 10000, regularization = lassoLambda) #plotting coefficients ax = plt.gca() ax.scatter(alphaList L, coefficients L) ax.set xscale('log') plt.title('LASSO/GB Model Coefficients of Linear Regression') plt.xlabel('alpha') plt.ylabel('coefficient') plt.axis('tight') plt.show() Ideal lambda is 0.1 LASSO/GB Model Coefficients of Linear Regression 0.8 0.6 **mefficient** 0.4 0.2 0.0 -0.2 $10^{-1}$ 10<sup>1</sup> 10° alpha from sklearn import linear model #3b #Train linear model  $alphaList_ls = [1*0.1 for 1 in range(1,101)]$ t = linear\_model.LinearRegression() x delete train = np.delete(x train, 0, 1) #reshape/delete 1. column so we can match train and test sets model = t.fit(x delete train, y train) #model coefficients coefficients\_ls = model.coef\_ #plot coefficients ax = plt.gca() ax.scatter(alphaList\_ls, coefficients\_ls) ax.set\_xscale('log') plt.title('Least Squares Model Coefficients of Linear Regression') plt.xlabel('alpha') plt.ylabel('coefficient') plt.axis('tight') plt.show() Least Squares Model Coefficients of Linear Regression 1.0 0.8 0.6 **mefficient** 0.4 0.2 0.0  $10^{-1}$ 10° alpha #3c In [29]: #Lasso #MSE on test data #best y predcitions from LASSO Y\_pred\_Lasso = lassoModel.predict(x\_test).reshape(-1,1) Y\_true\_Lasso = y\_test MSE\_Lasso = np.square(np.subtract(Y\_true\_Lasso, Y\_pred\_Lasso)).mean() **#Least Squares #MSE** on test data #best y predictions from LS intercept = model.intercept\_ slope = model.coef\_ for i in range(len(x\_test)): #y = mx+b Y\_pred\_LS = intercept + slope \* x\_test[i] Y\_true\_LS = y\_test MSE\_LS = np.square(np.subtract(Y\_true\_LS, Y\_pred\_LS)).mean() print(f"The mean square error on test set (true vs predicted output) for LASSO is {MSE\_Lasso} compared to {MSE\_ The mean square error on test set (true vs predicted output) for LASSO is 6.252351768583674 compared to 8.08601 7848092585 for least squares ... hence as LASSO produces a lower MSE, it is a model with better fit In [30]: #4a #using 1000 samples #load in samples hw4 = loadmat('HW1 4.mat') hw4.keys() #check data xdict = hw4['X']ydict = hw4['Y'] alpha = hw4['alpha'] #convert dict to np.array xlist = list(xdict) ylist = list(ydict) X = np.array(xlist)ytrue = np.array(ylist) #change to 1D arrays to plot X 1D = X[:,0]ytrue\_1D = ytrue[:,0] e = np.random.normal(loc = 0, scale = 0.01) #gaussian noise #added gausian noise  $ytrue_1D = ytrue_1D + e$ degree = 3 plt.scatter(X 1D, ytrue 1D, color = 'b', alpha = 0.5) #plot true model slope, intercept, r value, p value, std err = stats.linregress(X[:,0], ytrue[:,0]) #mtrue = linregress(x, ytrue) #to find slope mtrue[0] and intercept mtrue[1] of true plt.plot(X 1D, slope\*X 1D + intercept, '-y', label = 'true model') #plotting y = mx+b#estimated model with poly least squares (polyfit) polyfit = np.polyfit(X 1D, ytrue 1D, degree) yestimate = -polyfit[0]\*X\*\*3 + polyfit[1]\*X\*\*2 + polyfit[2]\*X + polyfit[3]mest = stats.linregress(X[:,0], yestimate[:,0]) #plot estimated model plt.plot(X 1D, mest[0]\*X 1D + mest[1], '-r', label = 'estimated model') plt.title('Polynomial Least Squares True and Estimated Model') plt.xlabel('X') plt.ylabel('y') plt.axis([-1.05, 1.05, 1, 7]) plt.legend() plt.show() Polynomial Least Squares True and Estimated Model estimated model 6 3 -1.00 -0.75 -0.50 -0.25 0.00 0.25 0.50 #4b #combine X and ytrue data = np.column\_stack([X\_1D,ytrue\_1D]) #fit line using all data model = LineModelND() model.estimate(data) #implement RANSAC with fitted line data to get model r model\_r, inliers = ransac(data, LineModelND, min\_samples = 2, residual\_threshold = 1, max\_trials = 1000) outliers = inliers == False #separate out the outliers from inliers #generate coordinates ytrue= model.predict\_y(X\_1D) yestimate = model\_r.predict\_y(X\_1D) #finding consensus set randset = [] con = [] #consensus set inliers\_count = 0 loop = 5for i in range(loop): #tried my best on this consensus set implementation randset.append(data[rand.randint(0, 1000)]) if randset[i][1] - ytrue[i] <= 0.3: #threshold</pre> con.append(randset[i]) if data[inliers][i][0] == randset[i][0] and data[inliers][i][1]== randset[i][1]: inliers count+=1 if len(con) > 0: if inliers\_count/len(con) \* 100 > 40: #inliers exceed 40% break else: continue loop**+=**1 #plotting consensus set for i in range(len(con)): con\_set = plt.plot(con[i][0], con[i][1], '.y', alpha=1, markersize = 30) #consensus set plt.plot(data[inliers, 0], data[inliers, 1], '.r', alpha=0.6, label = 'Inliers') #inliers plt.plot(data[outliers, 0], data[outliers, 1], '.b', alpha=0.6, label = 'Outliers') #outliers plt.plot(X 1D, ytrue, '-y', label='true model') plt.plot(X\_1D, yestimate, '-r', label='estimated model') plt.title('RANSAC True and Estimated Model with Consensus Set (Yellow Circles)') plt.xlabel('X') plt.ylabel('y') plt.axis([-1.05, 1.05, 1, 7]) plt.legend() plt.show() RANSAC True and Estimated Model with Consensus Set (Yellow Circles) Inliers Outliers true model estimated model 5 3 -0.50 -0.25 0.00 0.25 0.50 -0.75