Q1 The data was first extracted from the .dat file and seperated type. the "t" variable holds the discrete time stamps while the "y" variable holds the measured output at said timestamps. Note that the max parameters was increased from 5 to 6 to hindlight the behaviour of $\hat{\sigma}$ later on in part 3 of question 1. In [1]: import pandas as pd import numpy as np param max = 6 # Largest number of parameters data = np.loadtxt('dataHw1.dat') t = data[:, 0].copy()y = data[:,1].copy()A look-up table containing the regressors was generated. The width of this table (# of columns) is determined by the value of the "param_max" variable declaired in the previous cell. The code code cell used for estimating phi will make use of this table by slicing the required section (e.g. only the first 2 columns will be used when estimating 2 parameters). In [2]: phi = np.stack([(t - 1)**n for n in range(param max)], axis=1) # generation of phi look-up table pd.DataFrame(phi, columns=[f't^{i}' for i in range(param max)]) # Prints look-up table bellow Out[2]: t^0 t^1 t^2 t^3 t^4 t^5 1.0 0.0 0.0 0.0 0.0 0.0 1.0 1.0 1.0 1.0 1.0 1.0 32.0 2 1.0 2.0 4.0 8.0 16.0 9.0 27.0 243.0 1.0 3.0 81.0 4.0 16.0 64.0 256.0 1024.0 1.0 625.0 25.0 125.0 3125.0 1.0 5.0 36.0 216.0 1296.0 7776.0 6 1.0 6.0 1.0 7.0 49.0 343.0 2401.0 16807.0 8.0 64.0 512.0 4096.0 32768.0 8 1.0 1.0 9.0 81.0 729.0 6561.0 59049.0 1.0 10.0 100.0 1000.0 10000.0 100000.0 10 1.0 11.0 121.0 1331.0 14641.0 161051.0 11 1.0 12.0 144.0 1728.0 20736.0 248832.0 12 1.0 13.0 169.0 2197.0 28561.0 371293.0 13 1.0 14.0 196.0 2744.0 38416.0 537824.0 In the cell below, the least square estimates for parameters ranging from 1 to 5 (or param_max) are calculated with each iteration of the for loop Q1.1 & Q1.2 In [3]: import pandas as pd from numpy.linalg import inv theta_hat = [] # list for storing theta_hat loss = [] # list for storing the loss functions for i in range(0,param_max): phi_temp = phi[:, 0:i+1] # the "phi" look-up table is sliced as required for each iteration theta_temp = inv(phi_temp.T@phi_temp)@phi_temp.T@y # temporary storage of theta_hat estimate err = (y - phi_temp@theta_temp) # difference between measured output and estimated output theta_hat.append(np.append(theta_temp, [0]*(param_max - i - 1))) # estimated theta_hat for each # iteration are stored here loss.append(err@err/2) # Loss function for each iteration are stored here # Theta_hats for parameter counts ranging from one to 5 (or value of "param_max") are # packaged into a dataframe for presentation in table format df = pd.DataFrame(np.vstack(theta hat), index=[i for i in range(1,param max+1)], columns=[f'Theta{i}' for i in range(1,param max+1)]) df['Loss'] = loss # Loss column is added on far right side of table Out[3]: Theta1 Theta2 Theta3 Theta4 Theta5 Theta6 Loss **1** 51.435013 0.000000 0.000000 24103.092840 0.000000 0.000000 0.000000 0.000000 0.000000 4637.216740 **2** -31.106229 11.791606 0.000000 0.000000 11.150564 -7.711529 1.393081 0.000000 0.000000 0.000000 634.251379 8.137128 -4.576673 0.813574 0.027596 0.000000 0.000000 612.440363 3.497340 -1.992023 -0.011372 563.290183 4.234310 0.346007 0.000000 11.388026 -25.060590 14.203890 -2.876500 0.251265 -0.007504 293.073810 When observing the data in the table above, one can see that the loss function seems to stabilise around 3 parameters with relativley small differences for 4 parameters and up. Although the loss function seems to plateau, the value at which it levels out is still quite high. This is due to the large value of sigma (i.e. sigma = 11). The code in the cell below extracts the row coeresponding to 3 parameters (third row) and is post matrix multiplied by the "phi" look-up table. The row was padded with zeros for parameters 4 and 5 and therfore, can be directly multiplied by the entire look-up table when calculating the estimated y_hat vector. In [4]: import matplotlib.pyplot as mpl mpl.rcParams['figure.figsize'] = [20, 10] num param best = 3 # number of parameters to be used def y_hat_plotter(num_param_best): # extraction of row with 3 parameters excluding the "loss" column theta_hat_true = df.loc[num_param_best, df.columns != 'Loss'] y_hat = phi@theta_hat_true # calculation of estimated y_hat vector mpl.scatter((t - 1), y) # plot measured output y mpl.plot((t - 1), y_hat, 'r') # plot estimated output y_hat mpl.xlabel('t') mpl.ylabel('y_hat') mpl.title(f'y_hat vs. t with {num_param_best} parameters') mpl.legend(labels=['y_hat', 'y'], fontsize='xx-large') y hat_plotter(num_param_best) y_hat vs. t with 3 parameters y_hat У 150 125 100 75 50 25 Clearly, when analysing the graph above (y_hat vs. t), one can see that the curve representing the estimated outputs "y_hat" follows the trend of the measured values very closely. Q1.3 In [5]: sigma = np.sqrt(2*df['Loss']/(np.array([15]*param max) - df.index)) pd.DataFrame(sigma, columns=['Sigma']) Out[5]: Sigma **1** 58.679630 2 26.709885 10.281467 10.552383 10.614049 8.070162 The equation for the estimation of the standard deviation is $\hat{\sigma} = \sqrt{rac{2V(\hat{ heta},t)}{t-n}}$ where $V(\hat{ heta},t) = rac{1}{2} \sum_{i=1}^t (y(i) - \phi^T heta)^2$ Since y(i) is the measured output, it can be further exapnded to $y(i) = y(i)_{actual} + e(i)$ where $y(i)_{actual}$ is the actual output and e(i) is the iid noise. If this is substituted into the loss function, the equation becomes $V(\hat{ heta},t) = rac{1}{2} \sum_{i=1}^t (y(i)_{actual} + e(i) - \phi^T heta)^2$ If we assume that $\phi^T \theta$ ($y(i)_{estimate}$) is so close to $y(i)_{actual}$, that $y(i)_{actual} - \phi^T \theta pprox 0$, the loss function becomes $V(\hat{ heta},t) = rac{1}{2} \sum_{i=1}^t e(i)^2$ Plugging this result back into the equation for estimating $\hat{\sigma}$ we get $\hat{\sigma} = \sqrt{rac{\sum_{i=1}^{t} \overline{e(i)^2}}{t-n}}$ Since the mean of the noise is zero, we can conclude that the estimation for $\hat{\sigma}$ becomes the estimation of the standard variation of the noise. Therfore, if $\hat{\sigma} \approx \sigma$, then $\phi^T \theta$ must be very close to y_{actual} . When observing the table above, one can see that $\hat{\sigma}$ is closes to σ ($\sigma = 11$) when 5 parameters are being estimated and has a sharp drop for 6 parameters. It should be taken into consideration that 15 data points is not very much and the total distribution might not have completly taken the form of a gaussian distribution. For this reason, it would be prudent to use 4 parameters since it is in the middle of the three different parameters closes to 11. y_hat_plotter(4) y hat vs. t with 4 parameters y_hat 150 125 100 / hat 75 50 25 **Q.2** The cell below contains the initialisation of the experiment. All three inputs (i.e. $\delta(t-100)$, u(t-100) and $sin(\frac{2\pi t}{5}) + cos(\frac{4\pi t}{5})$) are created and inserted into a list of three lists. Each individual list will be referenced during their respective iteration of the loops 2 cells below. Additionally, θ and y(t) are also initialised here. In [7]: import numpy as np import pandas as pd from math import cos, sin, pi import scipy as spy from numpy.linalg import inv from numpy import sqrt from scipy.signal import unit_impulse import matplotlib.pyplot as mpl sample_depth = 3000 a1 = 1.3a2 = 0.75b0 = 1.1b1 = -0.35theta0 = np.array([a1, a2, b0, b1])t = [i for i in range(sample_depth)] u t1 = unit impulse(sample depth, 100) # Creating impulse delta(t - 100) u_t2 = np.zeros(sample_depth) # Creating unit step unit(t - 100) $u_t2[np.where(np.arange(0,sample_depth) >= 100)] = 1$ $u_t3 = np.array([sin(2*pi*t[i]/5) + cos(4*pi*t[i]/5))$ for i in t]) # Creating dual freq. sinusoid $u_t = np.stack([u_t1, u_t2, u_t3])$ # $impulse = u_t[0], step = u_t[1], dual freq. sinusoid = u_t3$ sigma = 0.65y0 = np.random.normal(0, sigma) # creating a list of three lists which will be used to store the outputs of all three runs $y = [[y0] \text{ for } i \text{ in } range(len(u_t))]$ theta_hat0 = np.reshape(np.array([0]*4), (-1,1)) # initial theta estimations will be 0 # creating a list of three lists which will be used to store the theta_hats of all three runs theta_hat = [[theta_hat0] for i in range(len(u_t))] The cell below contains two nested loops. The first loop determines which input vector will be used. The second loop, estimates the parameters recursively. During the first iteration, ϕ is set to $-y_{(j)(i-1)}, 0, u_{t(j)(i-1)}, 0$ since $-y_{(j)(i-2)}$ and $u_{t(j)(i-2)}$ do not exist yet. It should also be notted that the original equation given in the assingment was time shifted by 2 time units. This way, the equation relies on past values rather than future ones. Additionally, the final P matrix will be saved for each type of input. This will be used to calculate $\hat{\sigma_{b0}}$ and $\hat{\sigma_{b1}}$ in part 4 of this question. In [8]: p_final = [] # used to derive sigma_hat_b0 and sigma_hat_b1 in part 4 of this question for j in range(len(u t)): p = 100*np.identity(4) # starting P matrix for i in range(1, sample_depth): **if** (i == 1): # accounts for the lack of t-2 data on first iteration $phi = np.array([-y[j][i-1], 0, u_t[j][i-1], 0])$ else: $phi = np.array([-y[j][i-1], -y[j][i-2], u_t[j][i-1], u_t[j][i-2]])$ # changes phi's dimensions from (4,) to [4,1] enabling transpose operations phi = np.asarray(phi).reshape(-1,1) y[j].append(np.reshape(phi.T@theta0 + np.random.normal(0, sigma), ())) p = inv(inv(p) + phi@phi.T)k = p@phi $\label{eq:theta_hat[j]} $$ theta_hat[j].append(theta_hat[j][i-1] + k*(y[j][i] - phi.T@theta_hat[j][i-1])) $$$ # The data collected with this "if" statement will be used in part 4 of Q2 if (i == sample_depth - 1): p final.append(p) df_lst = [pd.DataFrame(np.asarray(theta_hat[i]).reshape(-1,4,), columns=['a1', 'a2', 'b0', 'b1']) for i in range(len(u_t))] In [9]: import matplotlib.pyplot as mpl import seaborn as sns mpl.rcParams['figure.figsize'] = [20, 10] def theta hat ploter(df, title, line width=1.2): graph = sns.lineplot(data=df, dashes=False) graph.axhline(y=a1, color='b', linestyle='--', linewidth=line_width, label='a1') graph.axhline(y=a2, color='y', linestyle='--', linewidth=line_width, label='a2') graph.axhline(y=b0, color='g', linestyle='--', linewidth=line_width, label='b0') graph.axhline(y=b1, color='r', linestyle='--', linewidth=line width, label='b1') mpl.title(title) mpl.ylabel('Magnitude of "Theta hat"') mpl.xlabel('Time Stamps "t"') mpl.legend(bbox_to_anchor=(1.05, 1), loc=2, borderaxespad=0, labels=df lst[2].columns, fontsize='xx-large') mpl.show() theta_hat_ploter(df_lst[0], 'Theta_hat Estimates for Impulse Input') theta_hat_ploter(df_lst[1], 'Theta_hat Estimates for Step Input') Theta_hat Estimates for Impulse Input a1 1.5 a2 b0 b1 1.0 0.5 Magnitude of "Theta hat 0.0 -0.5-1.0Time Stamps "t" Theta_hat Estimates for Step Input a1 1.50 a2 b0 1.25 1.00 0.75 Magnitude of "Theta hat" 0.25 0.00 -0.25 -0.50500 1500 Time Stamps "t" 2000 2500 3000 **Q2.1** In the first graph ('Theta_hat Estimates for Impulse Input'), it can be seen that a1 and a2 converge to their true values where as b0 and b1 do not. This is expected since a1 and a2 are the coefficients of the output regressors (i.e. y(t-1) and y(t-2)) while b0 and b1 are the coefficients of the inputs. Since the output is always stimulated by noise, the recursive least square estimates algorithum can pick-up and track changes in the y(t-1) and y(t-2) regressors and converge a1 and a2 to their true values. b0 and b1 on the other hand do not get stimulated until t=100, the effects of which can be seen in the figure above where b0 and b1 are estimated to be zero (initial condition) until t = 100. The impulse dies immediatley after (t = 101). For this reason, b0 and b1 move but never converge onto their true values. In the second graph ('Theta_hat Estimates for Step Input'), we can see a similar result where a1 and a2 converge to their true values while b0 and b1 do not. a1 and a2 converge for the same reasons they converged for an impulse input, however, b0 and b1 do not converge becasue a unit step is only sufficiently stimulating for one parameter. This can be explained qualitativley by thinking about the effects the input has on the output. At t=100, the output will see a sudden jump, just like in the case for the impulse, however, unlike the impulse, the step continues to contribute to the output. Although the output feels the effects of the input for the rest of the experiment, there is no way of determining by how much a parameter regressor pair is contributing to the output. In other words, the solution is not unique. **Q2.2** Eq. 2.46 from the textbook states that parceval's theorem can be used to determine teh order of excitation of an input signal. The equation is $rac{1}{2\pi}\int_{-\pi}^{\pi}\left|A(e^{j\omega})
ight|^2\Phi(\omega)d\omega>0.$ In this equation, the amplitude $A(e^{j\omega})$ is squared and will therfore always be positive. The frequency content of the signal given by $\Phi(\omega)$ is the component that will determine at what frequency the solution is non-zero. To determine the frequency content of the input signal, the fourier transform of the cosine and sin function will be derived. $F(cos(\omega t)) = \pi(\delta(\omega - \omega_0) + \delta(\omega + \omega_0))$ $F(sin(\omega t)) = -j\pi(\delta(\omega-\omega_0) - \delta(\omega+\omega_0))$ Either of these functions contribute 2 degrees of excitation. Since $\omega_0=rac{4\pi}{5}$ for the cosine term and $\omega_0=rac{2\pi}{5}$ for the sine function, both components contribute to the order of excitation, that is, 4. **Q2.3** In [10]: theta_hat_ploter(df_lst[2], 'Theta_hat Estimates for Dual Frequency Sinuziodal Input') Theta_hat Estimates for Dual Frequency Sinuziodal Input a1 a2 1.25 b0 b1 1.00 Magnitude of "Theta_hat" 0.50 0.25 0.00 500 2000 3000 In [12]: df lst[2] b0 b1 Out[12]: a1 a2 **0** 0.000000 0.000000 0.000000 0.000000 0.491600 0.000000 -0.234074 0.000000 **2** 0.460459 0.224409 -0.298980 -0.106852 0.952607 0.840574 0.734571 0.412290 **4** 1.040661 0.747066 0.917413 0.078551 2995 1.301388 0.749588 1.110715 -0.332253 2996 1.301409 0.749589 1.110638 -0.332022 1.110581 -0.332016 2997 1.301439 0.749607 2998 1.301419 0.749614 1.110515 -0.332121 2999 1.301299 0.749490 1.110709 -0.332487 3000 rows × 4 columns **Q2.4** In [11]: sigma hat b0 = [sigma*sqrt(p final[i][2,2]) for i in range(len(u t))] sigma_hat_b1 = [sigma*sqrt(p_final[i][3,3]) for i in range(len(u_t))] df = pd.DataFrame(np.stack([sigma_hat_b0, sigma_hat_b1]), columns=['Impulse', 'Step', 'Dual Freq.'], index=['sigma_hat_b0', 'sigma_hat_b1']) Out[11]: Impulse Step Dual Freq. **sigma_hat_b0** 0.646929 0.643737 0.014384 0.017412 **sigma_hat_b1** 0.646846 0.643978