02 Exercise Notebook 2

March 29, 2023

0.1 Exercise 2

In this exercise, we will plot COVID hospital admissions in the UK from March 2020 to February 2023, and you will fit an exponential curve to understand the rise in hospital admissions in a three week period from late December 2021 to early January 2022.

The data is in a file called hospital_cases_2023-02-16.csv (comma-separated-values format). It was downloaded from the official website for UK COVID-19 data.

```
[1]: # If you are running this on Google Colab, uncomment and run the following ⊔

ilines; otherwise ignore this cell

# from google.colab import drive

# drive.mount('/content/drive')
```

```
[2]: import math import numpy as np import matplotlib.pyplot as plt import pandas as pd
```

We will use pandas, a library for data analysis in Python to load and view the data. Pandas uses a structure called a *data frame* to represent and manipulate data tables. All the required commands are included here, so you won't need to learn Pandas for this exercise. But if you are interested in learning more, this is a good place to start.

```
[3]: date hospitalCases
0 27/03/2020 7267
1 28/03/2020 8278
2 29/03/2020 9525
```

The command pd.read_csv loads the data onto a data frame. We have used the .head() command to display the top 3 rows of the data frame.

We can also display a random sample of rows from the data frame using .sample(), or the last few rows using .tail().

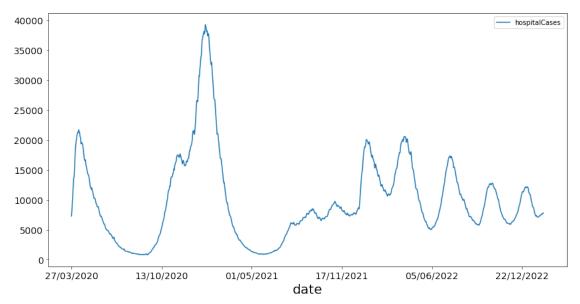
```
[5]: df_hosp.sample(5)
```

```
[5]:
                       hospitalCases
                 date
     273
          25/12/2020
                                20921
     988
          10/12/2022
                                 7533
     621
          08/12/2021
                                 7470
     526
          04/09/2021
                                 7570
     649
          05/01/2022
                                18221
```

You can plot one column against another by just using their column names. Let us plot the hospitalCases column versus date.

```
[6]: plt.rcParams['figure.figsize'] = [14, 7]
  plt.rcParams['axes.titlesize'] = 20
  plt.rcParams['axes.labelsize'] = 20
  plt.rcParams['xtick.labelsize'] = 14
  plt.rcParams['ytick.labelsize'] = 14

  df_hosp.plot(x='date', y='hospitalCases')
  plt.show()
```



Observe the sharp increase in hospital admissions corresponding to each wave; also notice that the peaks after mid-2021 are smaller (due to the vaccines). We now extract the rows spanning a three week period starting 22 December 2021 (when Omicron first spread in the UK) into a data frame called df_part

```
[7]: df_part = df_hosp[635:656]
df_part.head(2)
```

```
[7]: date hospitalCases
635 22/12/2021 8400
636 23/12/2021 8436
```

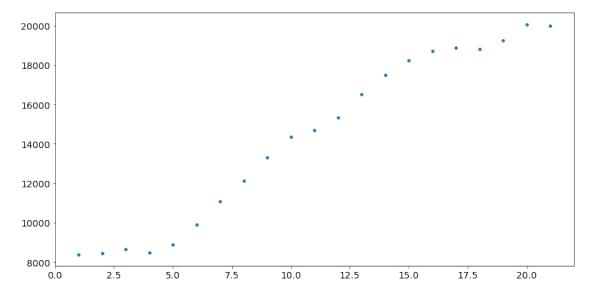
```
[8]: df_part.tail(2)
```

```
[8]: date hospitalCases 654 10/01/2022 20065 655 11/01/2022 19967
```

We now convert the hospitalCases column to a numpy array yvals

0.2 2a) Plot the data yvals vs xvals in a scatterplot

```
[12]: # Your code for scatterplot here
plt.scatter(xvals,yvals, s=15)
plt.show()
```



0.3 2b) Fit an exponential model to the data

From our knowledge of how the virus spreads, we know that the number of infections, hospital admissions etc. should (roughly) follow an exponential curve. We would therefore like to fit a model of the form $y = c_1 e^{c_2 x}$, where y is the number of admissions on day x.

Note that this is a linear model on a log-scale for y. That is, $\log y = \log c_1 + c_2 x$.

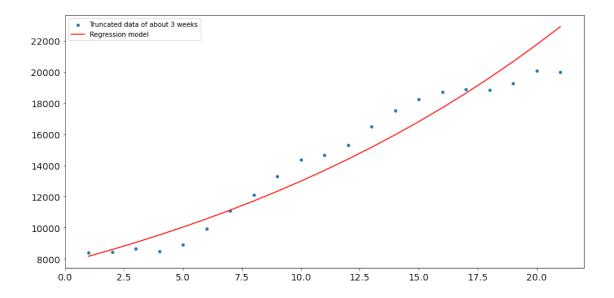
- Fit a linear model for log(yvals) vs xvals, and print the values of c_1 and c_2

• Plot the fit $y = c_1 e^{c_2 x}$ along with the scatterplot of the data

```
[15]: # your code here
      def polyreg(data_matrix: np.array, k: int)->np.array:
          The function returns the the coefficient vector beta, the fit X*beta, and
       \hookrightarrow the vector of residuals y-X*beta
          N, _ = data_matrix.shape
          t, y = [data_matrix[:,i] for i in range(2)]
          X = np.ones((N,k+1))
          for i in range(1,k+1):
              X[:,i]= t**i
          P = np.linalg.inv((X.T).dot(X))
          P = P.dot(X.T)
          beta = P.dot(y)
          fit = np.dot(X, beta)
          residual = y - fit
          return (beta,fit,residual)
      xy_data = np.column_stack([xvals, np.log(yvals)])
      beta, fit, residual = polyreg(xy_data, 1)
      c1, c2 = np.exp(beta[0]), beta[1]
      print(f'Value of C1 coefficient: {round(c1 ,3)}')
      print(f'Value of C2 coefficient: {round(c2 ,3)}')
      plt.scatter(xvals, yvals, s=15, label='Truncated data of about 3 weeks')
      plt.plot(xvals, c1 * np.exp( c2* xvals), 'r', label='Regression model')
      plt.legend()
      plt.plot()
```

Value of C1 coefficient: 7754.298 Value of C2 coefficient: 0.052

[15]: []



0.4 $\,$ 2c) Estimate the weekly growth rate in hospital admissions (in %) over this period

Hint: According to the model, admissions increase every 7 days by a factor of $(c_1e^{c_2(x+7)})/(c_1e^{c_2x})=e^{7c_2}$.

```
[17]: # compute and print weekly growth rate (in %)
growth_rate = 100 * (np.exp( 7.0 * c2) - 1.0)
print(f'Weekly growth rate in hospital admissions over this period:

→{round(growth_rate,2)}%')
```

Weekly growth rate in hospital admissions over this period: 43.47%