

EE3801 Data Engineering

Laboratory Exercise (LAB-I)

Assignment release date: Sept 02, 2020; Briefing: Sept 03, 2020

Date submission due: Sept 09, 2020

<u>Grading:</u> Your ASSIGNMENT will be graded out of 100 marks and the final weight of this assignment is <u>10%</u>. The guidelines to be followed, as explained in the template will also carry marks. So please adhere to the guidelines.

NOTE: This is an individual lab exercise and NO DISCUSSIONS AND EXCHANGE OF SOLUTION IDEAS BETWEEN ANY STUDENTS ARE ALLOWED. If we come to know of this <u>in any form</u>, we will be taking relevant disciplinary actions. Please follow this guideline strictly.

<u>Concepts used</u>: Data frames, data extraction from a given dataset, performing simple computations, use of pandas dataframe methods for data wrangling - extraction, handling missing data, transforming a DF to a target DF, and interpretation of results

Data required for this assignment: bodyfat2.csv, bodyfat3.csv

Consider **bodyfat2.csv** dataset given to you. This data is captured for individuals (each individual representing a row) as an outcome of a uniform medical testing.

(1a) **Compute** the mean, median, and sum, for each individual for columns starting from neck onwards till wrist (in your dataset) and print the top 3 and bottom 3 values; store all values in a dataframe [6 rows by 3 columns]. Display your results using meaningful messages. Give your dataframe a meaningful name. [10 Marks]

- (1b) **Compute** the mean, median, and sum, for each feature using the respective methods and print your results clearly <u>with meaningful messages</u>. **NOTE:** You can use ".describe()" <u>only to compare</u> your results, if you wish. Give your dataframe a meaningful name. [5 Marks]
- (2) In bodyfat2.csv dataset, <u>for every feature (other than age, weight and height features)</u>, <u>identify the individuals</u> that have maximum and minimum fat. Store your results (<u>max value, corresponding feature, min value, corresponding feature</u>) in a dataframe and display with a meaningful message. So you may store as a 12 x 4 size dataframe. The individuals are to be captured as their respective row indices. [10 Marks] (Sample DF is shown below)

| Feature | Max value | Individual ID | Min value | Individual ID |
|---------|-----------|---------------|-----------|---------------|
| density | | | | |
| bodyfat | | | | |
| neck | | | | |
| | | | | |

Note: You can use ".describe()" <u>only to compare</u> your results, if you wish, on max and min values.

- (3) Find **number of entries** (individuals) **in each feature** (column) that fall within 10% of standard deviation from its respective mean and median metrics. Store your results as a dataframe and display with meaningful messages. [15 Marks]
- (4) In **bodyfat3.csv** data given to you, <u>count the number of missing values in every feature</u> and print your results as a DF clearly with a meaningful message. [5 Marks]
- (5a) Copy your **bodyfat3.csv** as **bodyfat3b.csv**. In **bodyfat3b.csv** dataset, for each feature, write a python code to replace the missing values with MEAN of that feature. Compute the **difference in mean values** for each feature by comparing it with the <u>original mean from **bodyfat2.csv** dataset</u>. Display your results using meaningful messages always. [15 Marks]
- (5b) Copy your *bodyfat3.csv* as *bodyfat3c.csv*. Using *bodyfat3c.csv*, repeat (5a) using **MEDIAN** metric and report your findings. [15 Marks]

- (5c) Use the results of 5(a) and 5(b) to compare the accuracies and state your inference on the results. [10 Marks]
- (6) Consider the results of Problem 1(b) [bodyfat2 dataset]. For every feature, normalize the values using the expression:

$$x'_{i,f} = \frac{x_{i,f} - \mu_f}{\sigma_f}$$

Where μ_f denotes the feature mean and σ_f denotes the feature standard deviation.

- (i) Store all the results in a **separate dataframe**. Print the top 3 and bottom 3 rows from this new dataframe. [2 Marks]
- (ii) For each feature (all 15 features) in this new dataframe, compute the number of individuals that are greater than the respective feature's mean and store your results as a Series. Print the series with a meaningful message. [8 marks]
- Overall presentation of your results with clarity, meaningful names to DFs, meaningful messages in your output, and comments: 5 Marks