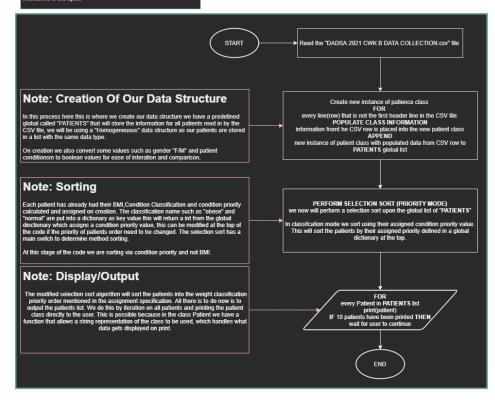
FLOWCHART DESIGN DIAGRAM FOR TASK 1

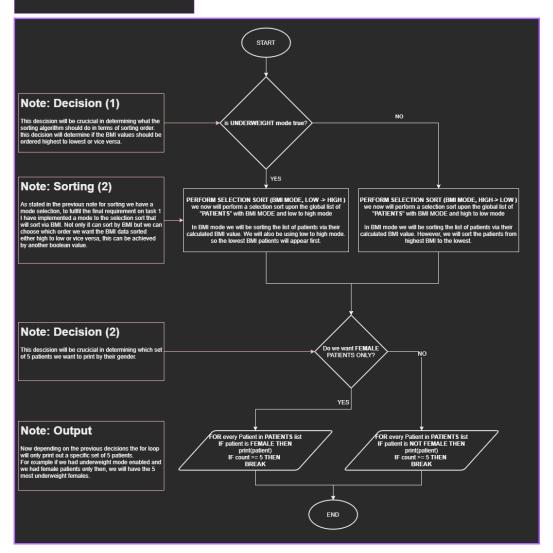
Flowchart for Task 1

The flowchart below describes the process of the solution.py for task 1. It will only cover the read/construction of data and output process for printing patients by their weight classification order as mentioned in the spec.



Flowchart for Task 1

This flow chart covers the process for the last requirement for task 1, I seperated from the other main flow chart for simplicity and readability.



In the flowchart diagram above you can see my design for task 1. In this example I have described and outlined the processes of how my python script will handle and sort the patient information data and from the CSV file.

PSEUDO CODE

Read Datafile

FUNCTION READ_DATAFILE(): called on application start when we want to read in the csv data

global variable reference to patients

global PATIENTS

open the datafile and access it via variable named csvfile

WITH open(datafile name()) AS csvfile:

line_count = 0 tracks the lines

csv reader will read the csv file and separate it with the given delimiter character of a comma spamreader = csv.reader(csvfile, delimiter=',')

FOR row IN spamreader: for every row in the CSV
IF line_count != 0: IGNORE FIRST LINE OF CSV

Create new patient and argument pass the row data into the patient object.

row[0] will be patient name, row[1] will be patient date of birth.

Then append that patient to the GLOBAL patients list referenced at the top of this function

line_count += 1 increment line count, used to track which line to ignore

close the CSV file once we done reading it csvfile.close()

Perform selection sort with no additional arguments, by default the patients will be sorted by condition priority which is determined and calculated on creation of new patient within the constructor

selection_sort(PATIENTS)

In the provided text above you can see my pseudo code for task 1. In this example I am showing the read datafile method. This segment of code is responsible for reading the CSV file and converting it into a homogenous data structure. Each row of the CSV file will have its column values passed into a new instance of patient class and its data simplified to the use of Ternary operators: "Y" == True etc.

Patient Class

CLASS PATIENT:

FUNCTION CONSTRUCTOR(self, name, dob, isfemale, height, weight, bodybuild, smoker, asthmatic, NJT, hypertension, renal rt, ileostomy, parenteral nutrition):

On class construction store passed argument values on this class self

```
self.identity_name = id_name
self.dob = CONVERT STRING DOB TO DATETIME DATA TYPE
self.is_female = is_female
self.height = height
self.weight = weight
self.body_build = body_build
self.smoker = smoker
self.asthmatic = asthmatic
self.nasogastric_tube = nasogastric_tube
self.hypertension = hypertension
self.renal_rt = renal_rt
self.ileostomy = ileostomy
self.parenteral_nutrition = parenteral_nutrition
```

Calculate BMI and BMI Classification and priority on patient creation, as well as their age and assign it to self in the class

```
self.bmi = CALCULATE_BODYMASS_INDEX(self)
self.classification = DETERMINE_CLASSIFICATION(self)
self.conditionPriority = CONVERT BMI CLASSIFICATION TO PRIORITY
self.age = CALCULATE_DOB_TO_AGE(self)
```

In the provided text above you can see my pseudo code for task 1. In this example I am showing the patient class. This class is responsible for holding all data about our patient.

Sorting Algorithm: Selection

```
# selection_sort :: takes in python list of patient objects sorts them by condition priority if sort_via_bmi is false
# if sort via bmi is true then low to high determines the sorting order of the BMI values
FUNCTION SELECTION_SORT(array, sort via bmi = False, low to high = True):
  FOR i IN range(0, len(array)): # for every element in the list named array
    Finding the smallest number in the subarray
    min = i
    for every element in the array from i + 1 we compare the condition values if sort via bmi Boolean is false
    FOR j IN range(i+1,len(array)):
      IF sort_via_bmi: sorts via BMI if argument is provided, default is false
         sorts the BMI low to high by default if set to false the ternary operators will switch the sort order to high to low
         IF ((array[min].bmi > array[j].bmi) IF low to high ELSE (array[min].bmi < array[j].bmi) ):
           min = j # set min from j index found
       ELSE:
         if array[min].conditionPriority > array[j].conditionPriority: # conditionPriority is a int value set on creation of patient
           set min from j index found
           min = j
    IF (min != i):
       # perform swap
      temp = array[i]
      array[i] = array[min]
      array[min] = temp
```

In the provided text above you can see my pseudo code for task 1. In this example I am showing the sorting algorithm used to sort the patients. The sorting algorithm will sort the patients via lowest to highest by default unless the Boolean is given to sort it highest to lowest. There is also another Boolean that determines if we should sort via bmi values or their assigned given condition priority.

Calculate Body Mass Index Function

calculate_bodymass_index :: takes in object type Patient, gets weight and height from patient object # returns float body mass index

FUNCTION CALCULATE_BODYMASS_INDEX(patient):

RETURN patient.weight / patient.height**2

In the provided text above you can see my pseudo code for task 1. In this example I am showing the calculate body mass index function. This function is responsible for calculating the BMI of a given patient object, the patient object will store their weight and height as a public variable. We access these variables to calculate the BMI with the given equation in the spec. the result is then assigned upon the patient object as a new value called BMI.

Determine patient classification Function:

determine_classification :: takes in object type Patient::returns string classification FUNCTION DETERMINE_CLASSIFICATION(patient):

```
depending on the body build we change the overweight to obese threshold :: seems to be only factor changing
IF patient.body_build.lower() == "slim":
  overweight to obese threshold = 28
ELIF patient.body build.lower() == "regular":
 overweight_to_obese_threshold = 29
ELIF patient.body build.lower() == "athletic":
  overweight to obese threshold = 30
depending on the patient BMI and the overweight_to_obese_threshold
return the classification of this patient which is assigned in patient class
IF(patient.bmi < 18.5):
  RETURN "underweight"
ELIF(patient.bmi > 18.5 and patient.bmi < 25):
  RETURN "normal"
ELIF(patient.bmi > 25 and patient.bmi < overweight to obese threshold):
  RETURN "overweight"
ELIF(patient.bmi > overweight to obese threshold):
  RETURN "obese"
```

In the provided text above you can see my pseudo code for task 1. In this example I am showing the determine patient classification. This function is responsible for calculating and assigning a patient classification name. The patient object will have variables stored upon it that contain their body build and bmi. We access these variables to calculate the patient classification to the requirements of the spec. the classification name is then returned and assigned on the patient object.

Calculate Age

```
calculate_dob_to_age :: takes in object type Patient :: returns int age of patient
FUNCTION CALCULATE_DOB_TO_AGE(patient):
    get the current date now
    get the patient born date of birth

calculate the difference from datenow and the born date from the patient :: return int year
```

In the provided text above you can see my pseudo code for task 1. In this example I am showing the calculate dob to age function. This function is responsible for calculating and assigning a patient age. The patient object will have a variable stored upon it that contain date of birth. We access this variable to calculate the patient age by finding the difference with date time now. The returned value is assigned and stored on the patient.