



**Team:**

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Abed

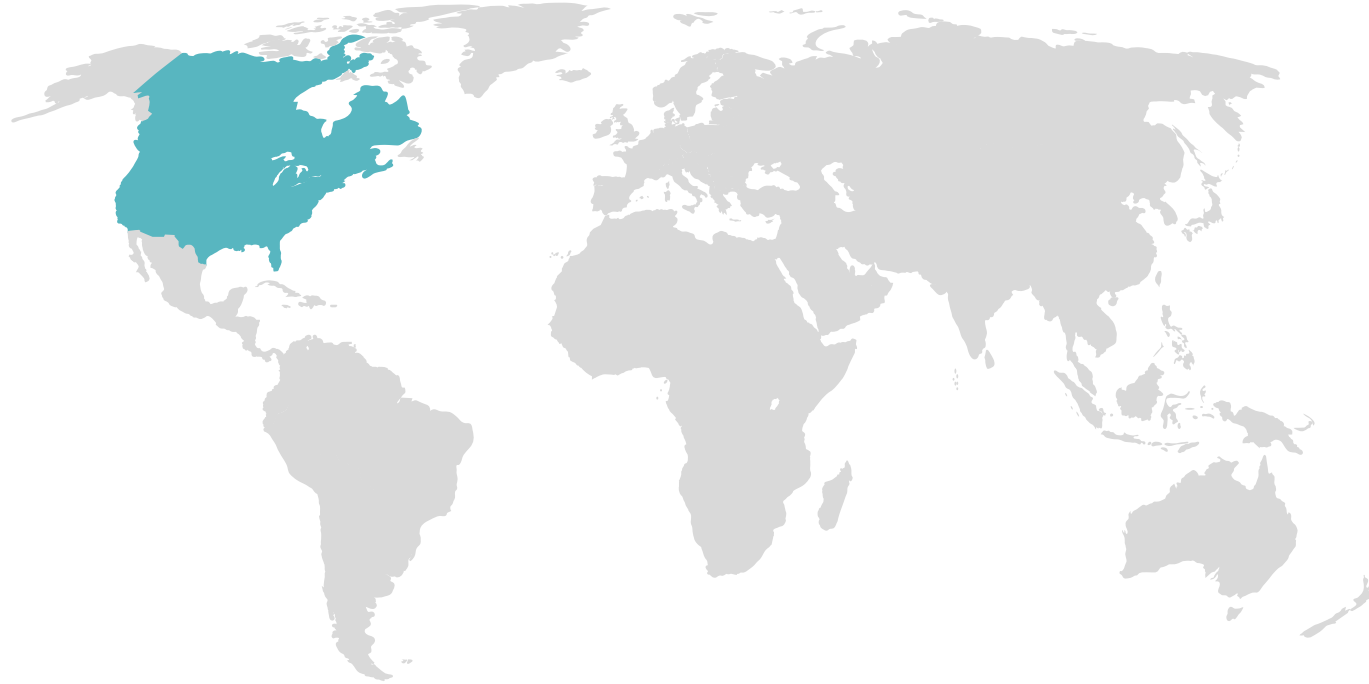
Saleh Abbas

Fatima Twam

Kayan Abukhaizaran

# Medication Non – Adherence

United States



■ **2/3 of Non-Adherent Americans with Prescriptions**

■ **\$300 BILLION Costs to the US Health Care System**

■ **125,000 Premature Deaths**

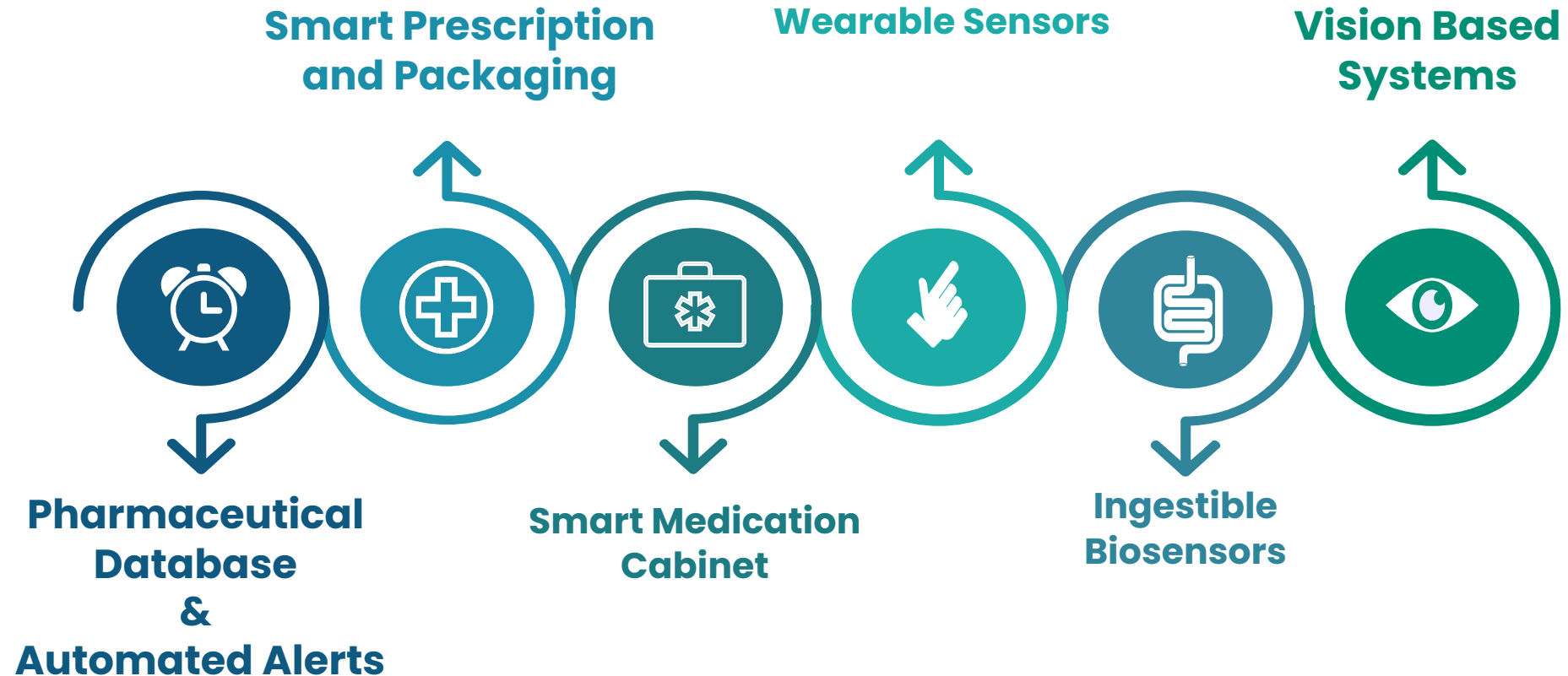
■ **25% of Hospital Admissions**

# Consequences on Healthcare system

Medication non – adherence effects on healthcare system



# SOLUTIONS IN THE MARKET



# TARGET MARKET



## Annual Health Expenditure

Spent on Chronic and Mental Health Diseases



Adults have a chronic disease



Adults have two or more chronic diseases

## **Variable Selection**

Our domain expert team chose the  
needed variables

01

## **Ethical Approval was Obtained**

From IAH Ethical Committee

02

## **EDA Pre-existing Data**

03

## **Data Collection stage preperation**

Feedback was sent from our team to  
hospital to ensure data quality

04

## **Link to EDA Report**

05

[https://drive.google.com/drive/folders/1XsJkP1InuYm\\_A5k5ERISvCM\\_qGGGRtGL?usp=sharing](https://drive.google.com/drive/folders/1XsJkP1InuYm_A5k5ERISvCM_qGGGRtGL?usp=sharing)

# **Our EDA Report**

Diagnosis  
Admission Date and time  
Discharge Date and time  
Gender  
Age  
Height  
Weight  
Past Medical History  
Past Surgical History  
Medication History  
Smoking and Alcohol intake

### **Observations:**

Blood Pressure,  
Heart Rate,  
Respiratory Rate ,  
Temperature

### **Medication Orders**

#### **Lab Tests :**

CBC, Liver Function  
Tests, Lipid Profile

# Variables Selection

The data were obtained from the Health Information System (HIS) of Istishari Arab Hospital (IAH).

# LEGAL AND ETHICAL REQUIREMENTS



المستشفى الإستشاري العربي  
ISTISHARI ARAB HOSPITAL

أكرم من جبر مستشفى

## IAH Research Application Form

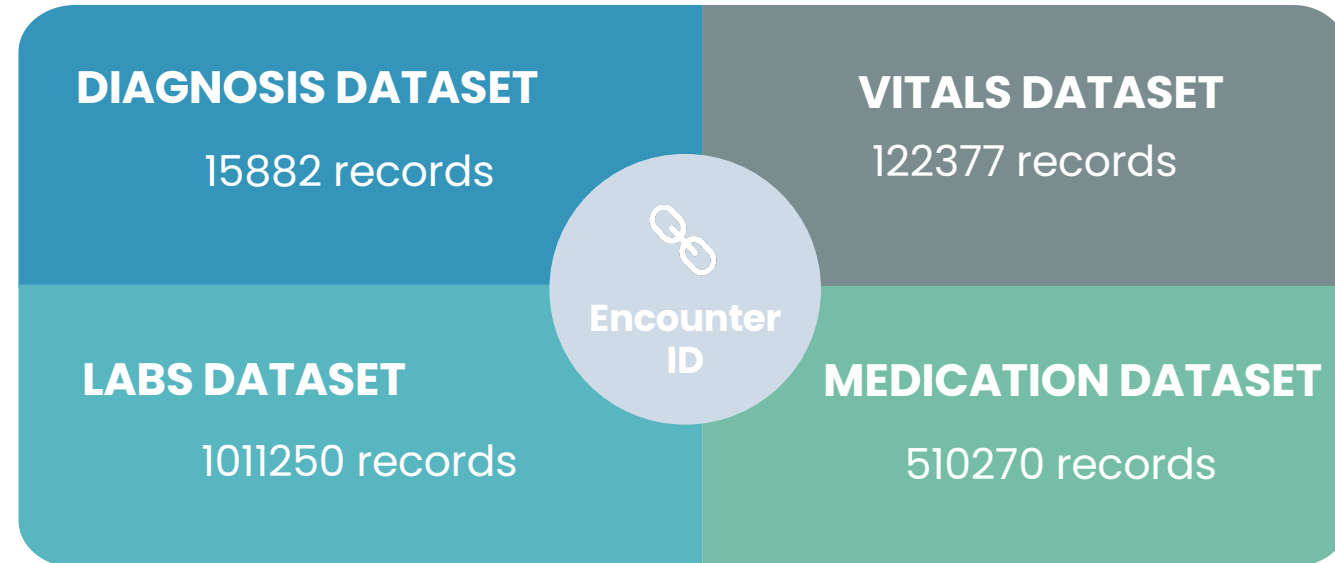
Date	19/01/2022
Name of investigator	Kayan Abukhaizaran
Mobile No.	0599133218
Email	kayan@iah.ps
Expected start date	
Expected completion date	
Name of Company/University	Birzeit University
Attached needed	
Investigator CV	<input type="checkbox"/> Yes <input type="checkbox"/> No
Study Proposal	<input type="checkbox"/> Yes <input type="checkbox"/> No
Consent Form	<input type="checkbox"/> Yes <input type="checkbox"/> No
Data Collection Tools	<input type="checkbox"/> Yes <input type="checkbox"/> No
Informed Consent (Arabic & English)	<input type="checkbox"/> Yes <input type="checkbox"/> No
For COO Office	
Receiving Date	
Application completed	<input type="checkbox"/> Yes <input type="checkbox"/> No
COO Director Note	
Transfer Date	
COO director Sig.	
For Ethical Committee	
Receiving Date	24/1/2022
Ethical Committee Approval	<input checked="" type="checkbox"/> Yes <input type="checkbox"/> No
Ethical Committee Note	Analytical research for non medical purposes Low ethical potential 25/1/2022
Head of Ethical committee Sig.	Dr. [Signature]
CEO Note	OK
CEO Sig.	[Signature]

• For Non Experimental Research only

26/1/2022



# OUR DATASETS



## Data collection phase:

IAH created a query so that the data will be provided to our team on a monthly basis.

# Exploratory Data Analysis

The below mentioned steps were conducted on all the datasets.

**Loading Dataset**

**Checking for  
missing values**

**Checking for  
Data Type**

**Splitting Date  
and Time**

**Saving  
as csv file**

## Checking for Data Type

```
[ ] DAG.info()
```

```
<class 'pandas.core.frame.DataFrame'>  
Int64Index: 15882 entries, 0 to 15881  
Data columns (total 11 columns):  
#   Column                                Non-Null Count  Dtype  
---  ---  
0   #                                     15882 non-null  int64  
1   Encounter ID                         15882 non-null  float64  
2   Patient ID                           15882 non-null  float64  
3   Gender                               15882 non-null  object  
4   Age                                  15882 non-null  float64  
5   Encounter Type                       15882 non-null  object  
6   Current_Department_Name              15882 non-null  object  
7   Diagnosis                             15880 non-null  object  
8   First Physician Note                 15788 non-null  object  
9   Amission Date                       15882 non-null  datetime64[ns]  
10  Discharge Date                      15875 non-null  datetime64[ns]  
dtypes: datetime64[ns](2), float64(3), int64(1), object(5)  
memory usage: 1.5+ MB
```

## Checking for Data Type

```
[ ] Vital.info()
```

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 122377 entries, 0 to 122376
Data columns (total 11 columns):
#   Column                Non-Null Count  Dtype
---  -
0   ENCOUNTER_ID          122377 non-null float64
1   PATIENT_ID            122377 non-null float64
2   READ_DATE             122377 non-null datetime64[ns]
3   HEIGHT                122377 non-null float64
4   WEIGHT                122377 non-null float64
5   TEMP                  78218 non-null  float64
6   PULSE                 86567 non-null  float64
7   RESPIRATORY_RATE      122377 non-null float64
8   RES_RATE              11018 non-null  float64
9   BP_SYSTOLIC           122377 non-null float64
10  BP_DIASTOLIC           122377 non-null float64
dtypes: datetime64[ns](1), float64(10)
memory usage: 11.2 MB
```

## Checking for Data Type

```
[ ] Lab.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1011250 entries, 0 to 1011249
Data columns (total 13 columns):
#   Column                Non-Null Count  Dtype
---  -
0   #                      1011250 non-null  int64
1   Encounter ID           1011250 non-null  int64
2   Order Type             1011250 non-null  object
3   Patient ID             1011250 non-null  int64
4   Order ID               1011250 non-null  int64
5   Result Category        1011250 non-null  object
6   Product Name           1011250 non-null  object
7   Result Name            1011250 non-null  object
8   Result Notes           1011250 non-null  object
9   Normal Range           893704 non-null   object
10  Unit                   960930 non-null   object
11  RESULT_DATETIME        1011250 non-null  datetime64[ns]
12  APPROVE_DATETIME       1011250 non-null  datetime64[ns]
dtypes: datetime64[ns](2), int64(4), object(7)
memory usage: 100.3+ MB
```

## Checking for Data Type

```
[ ] Med.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 510270 entries, 0 to 510269
Data columns (total 6 columns):
#   Column                Non-Null Count  Dtype
---  -
0   EncounterID           510270 non-null  int64
1   Patient ID           510270 non-null  int64
2   Product Name         510270 non-null  object
3   Instructions          502883 non-null  object
4   Order Date Time      510270 non-null  datetime64[ns]
5   Applied Date Time    510270 non-null  datetime64[ns]
dtypes: datetime64[ns](2), int64(2), object(2)
memory usage: 23.4+ MB
```

## Splitting Date and Time

### ▼ Splitting Date and Time

```
[ ] amission_date = []  
    amission_time = []  
    for amission_datetime in CDAG['Amission Date']:  
        amission_date.append(amission_datetime.date())  
        amission_time.append(amission_datetime.time().replace(microsecond=0))
```

```
[ ] discharge_date = []  
    discharge_time = []  
    for discharge_datetime in CDAG['Discharge Date']:  
        if(discharge_datetime is not pd.NaT):  
            discharge_date.append(discharge_datetime.date())  
            discharge_time.append(discharge_datetime.time().replace(microsecond=0))  
        else:  
            discharge_date.append(0)  
            discharge_time.append(0)
```

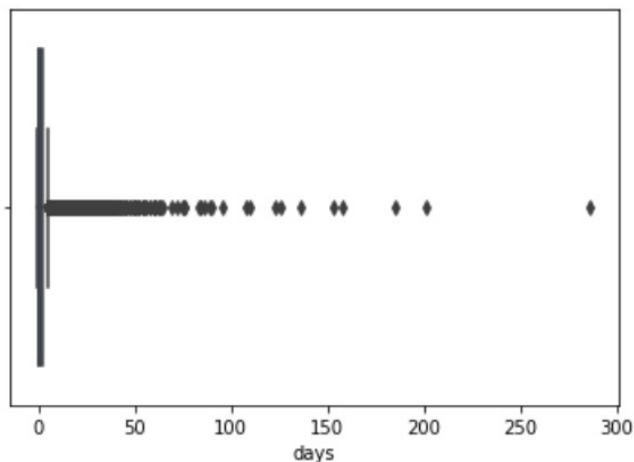
```
[ ] CDAG['AMMISSION_DATE'] = amission_date  
    CDAG['AMMISSION_TIME'] = amission_time  
    CDAG['DISCHARGE_DATE'] = discharge_date  
    CDAG['DISCHARGE_TIME'] = discharge_time
```



# Calculating Lengths of Stay (LOS)

```
✓ [19] CDAG['days']=np.round(CDAG['days'], decimals=2)
```

```
✓ [20] sns.boxplot(CDAG['days'])
```



```
✓ 0s ▶ EDAG.describe()
```



	Encounter ID	Patient ID	Age	days
count	15879.000000	1.587900e+04	15879.000000	15872.000000
mean	258863.571132	1.538143e+08	46.199194	2.453103
std	22254.918077	7.381813e+09	22.137496	6.875492
min	219931.000000	1.930000e+02	1.000000	-0.650000
25%	239986.500000	7.296900e+04	31.000000	0.270000
50%	258958.000000	8.914700e+04	50.000000	0.690000
75%	278023.500000	9.944250e+04	63.000000	2.100000
max	297909.000000	4.201611e+11	98.000000	286.010000





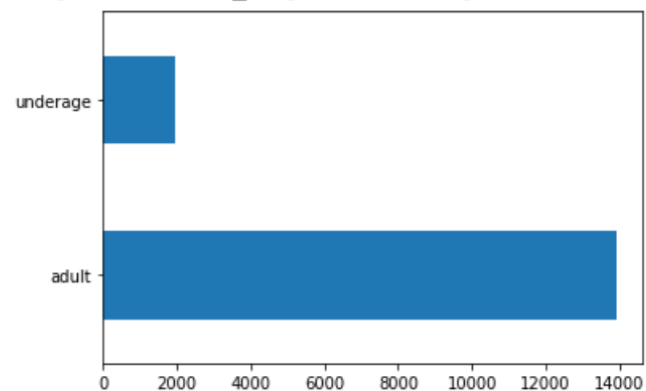


# Categorizing Age Groups

```
[ ] conditions = [  
    (EDAG['Age'] <= 18),  
    (EDAG['Age'] > 18)  
]  
  
values = ['underage', 'adult']  
  
EDAG['Age_Group'] = np.select(conditions, values)
```

```
[ ] EDAG['Age_Group'].value_counts()[0:20].plot(kind='barh')
```

<matplotlib.axes.\_subplots.AxesSubplot at 0x7f3f04d86990>



```
[ ] Adult_EDAG=EDAG.loc[EDAG['Age'] > 18]
```



## Data Extraction from “First Physician Note”

---

```
[55] pmh=0
      for x in DAG['First Physician Note']:
          gg=str(x)
          if ('past medical history' in gg.lower()) or ('past medical' in gg.lower()) or ('medical history' in gg.lower()) or ('pmh' in gg.lower()) or ('pmhx' in gg.lower()):
              pmh=pmh+1
      print(pmh)
```

7284

### **Data collection phase:**

Our requested variables will be obtained from new screens currently in the testing phase

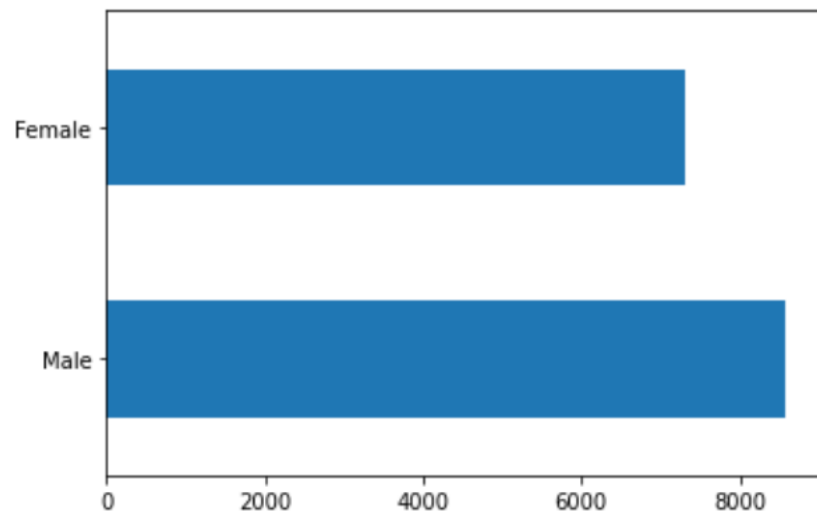


# Checking for Gender values

---

```
[ ] EDAG[ 'Gender' ].value_counts()[ :20 ].plot(kind='barh')
```

<matplotlib.axes.\_subplots.AxesSubplot at 0x7f3f04a90a90>





# Mapping “Diagnosis” with ICD-10

---

```
[ ] icd = pd.read_excel('/content/drive/MyDrive/Data-science-project/mappingData/ICD Data List.xlsx')
```

```
[ ] icd.info()
```

```
[ ] del icd['ICD_CODING']  
del icd['CATEGORY_CODE']
```

```
[ ] icd['ICD_FULL_DESC']=icd['ICD_FULL_DESC'].str.lower()  
icd['ICD_FULL_DESC']=icd['ICD_FULL_DESC'].str.strip()
```

```
[ ] C2DAG['Diagnosis']=C2DAG['Diagnosis'].str.lower()  
C2DAG['Diagnosis']=C2DAG['Diagnosis'].str.strip()
```

```
[ ] dagMapping=C2DAG.merge(icd, left_on='Diagnosis',right_on='ICD_FULL_DESC', how='left')
```

```
[ ] dagMapping.info()
```

Double-click (or enter) to edit

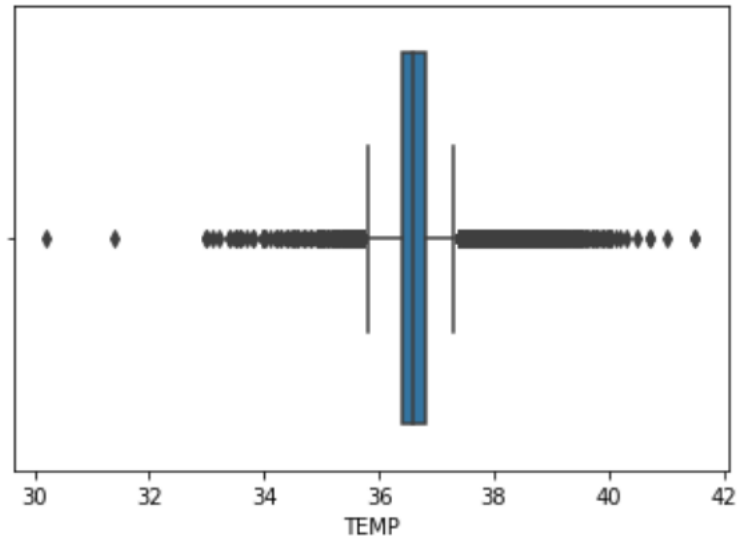
```
[ ] del dagMapping['ICD_FULL_DESC']  
del dagMapping['Diagnosis']
```



# Validating the “Temp”

```
✓ [105] EVitals['TEMP'] = EVitals['TEMP'].replace(np.nan, 0)
```

```
✓ [107] sns.boxplot(EVitals.loc[(EVitals['TEMP'] > 30)&(EVitals['TEMP'] < 45)]['TEMP'])
```



## Data collection phase:

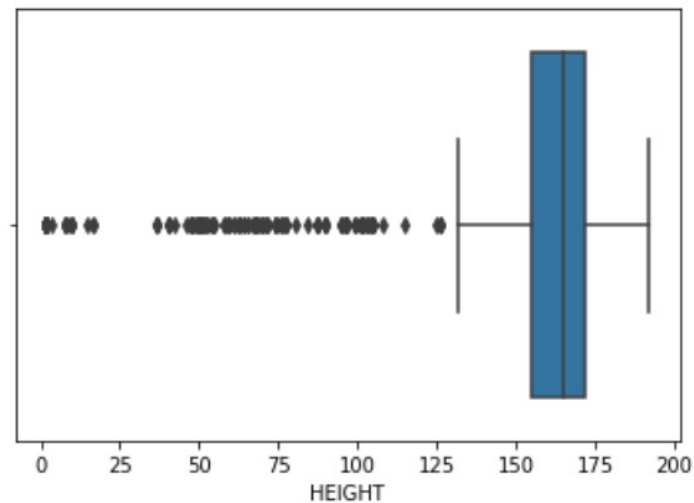
To overcome the outliers and fault values due to human errors. We provided the IAH with validation tools to be applied on the observation fields.



# Validating the “Height”

✓ [91] moreThanZero=EVitals.loc[EVitals['HEIGHT'] > 0]['PATIENT\_ID'].unique()  
0s

✓  sns.boxplot(EVitals.loc[EVitals['HEIGHT'] > 0]['HEIGHT'])  
0s



# Validating the “Weight”



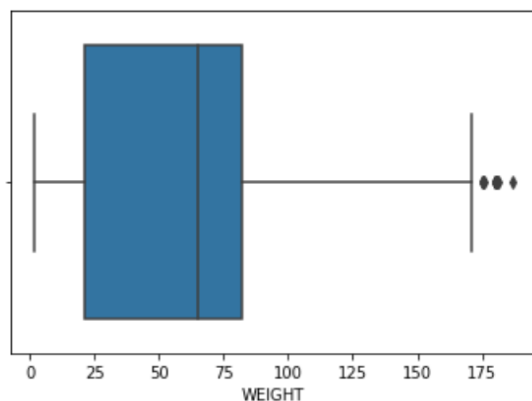
```
✓ [99] moreThanZeroWeight=EVitals.loc[(EVitals['WEIGHT'] > 0)&(EVitals['WEIGHT'] < 190)][['PATIENT_ID']].unique()  
0s
```

```
✓ [104] EVitals.loc[EVitals['WEIGHT'] > 190]  
0s
```

ENCOUNTER_ID	PATIENT_ID	HEIGHT	WEIGHT	TEMP	PULSE	RESPIRATORY_RATE	RES_RATE	BP_SYSTOLIC	BP_DIASTOLIC	READ_NDATE	READ_TIME
50429	252533	77042	163.0	923.0	36.7	88.0	0.0	NaN	141.0	86.0	2021-06-17 13:32:34
50849	252533	77042	163.0	923.0	NaN	NaN	0.0	NaN	0.0	0.0	2021-06-18 22:42:23
50987	252533	77042	163.0	923.0	NaN	NaN	0.0	NaN	0.0	0.0	2021-06-19 09:25:59



```
✓ [105] sns.boxplot(EVitals.loc[(EVitals['WEIGHT'] > 0) & (EVitals['WEIGHT'] < 190)][['WEIGHT']])  
0s
```

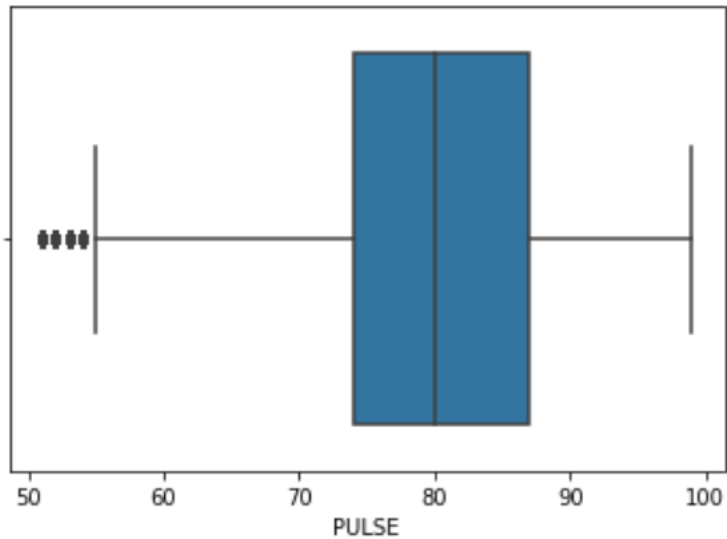




# Validating the “Pulse”

---

✓ 0s  `sns.boxplot(EVitals.loc[(EVitals['PULSE'] > 50)&(EVitals['PULSE'] < 100)][['PULSE']])`

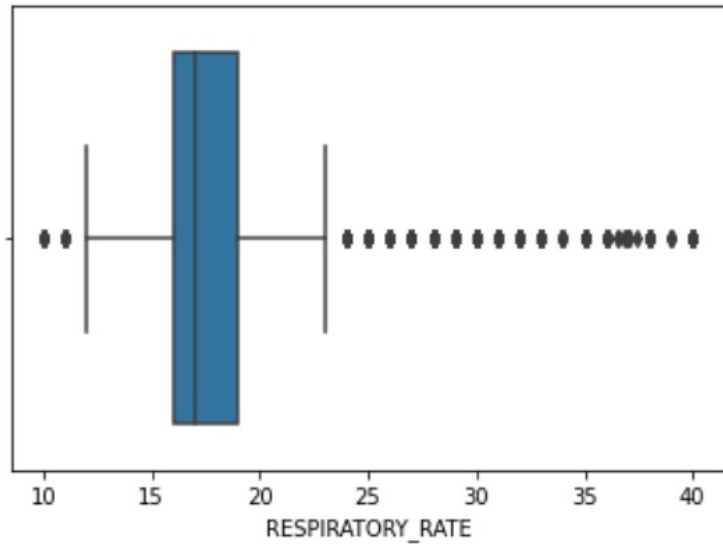







# Validating the “Respiratory Rate”

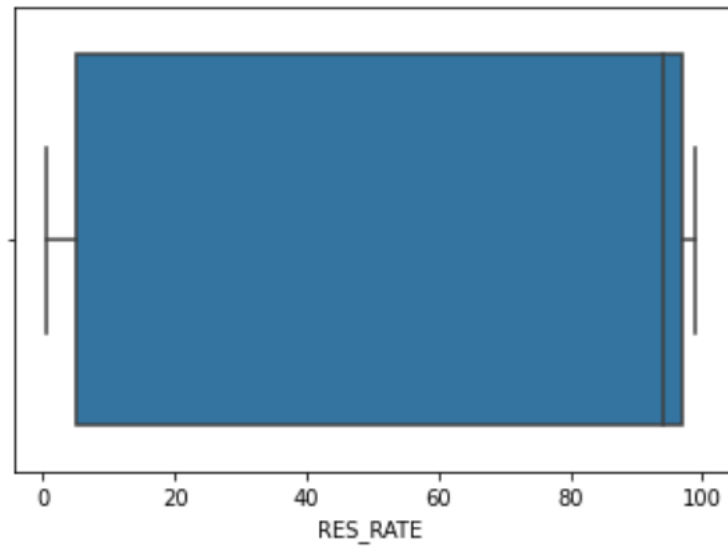
```
✓ [124] sns.boxplot(EVitals.loc[(EVitals['RESPIRATORY_RATE'] >= 10)&(EVitals['RESPIRATORY_RATE'] <= 40)]['RESPIRATORY_RATE'])
```





# Validating the “O2\_Saturation”

✓ 0s  `sns.boxplot(EVitals.loc[(EVitals['RES_RATE'] < 100)][['RES_RATE']])`

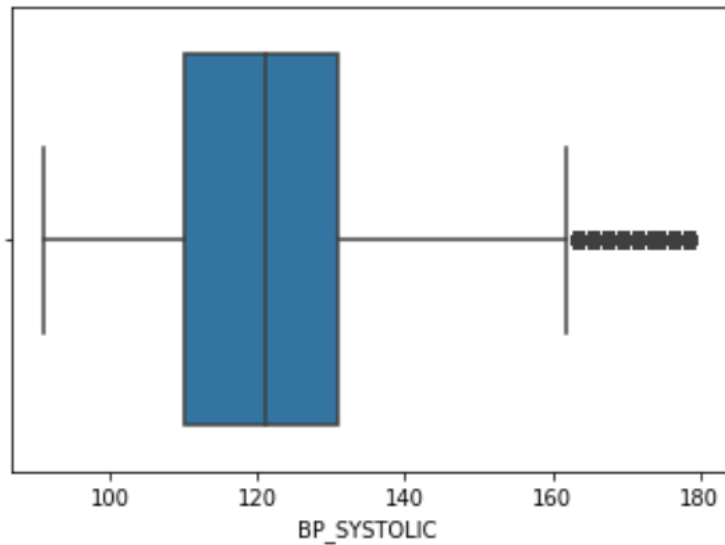




# Validating the “Blood Pressure”

✓  
0s

```
▶ sns.boxplot(EVitals.loc[(EVitals['BP_SYSTOLIC'] > 90)&(EVitals['BP_SYSTOLIC'] < 180))['BP_SYSTOLIC'])
```





# Mapping the “Product Code”

```
[ ] drug = pd.read_excel('/content/drive/MyDrive/Data-science-project/mappingData/Drug codes.xlsx')
```

```
[ ] drug = drug.astype({"Product Code": "int"})  
del drug['Standard Code']  
del drug['Usage Name']  
del drug['STOCK_BASE_UOM_DESC']
```

```
[ ] C2Med['Product Name'] = C2Med['Product Name'].str.lower()  
C2Med['Product Name'] = C2Med['Product Name'].str.strip()
```

```
[ ] drug['Product Name'] = drug['Product Name'].str.lower()  
drug['Product Name'] = drug['Product Name'].str.strip()
```

```
[ ] C2MedMapping = C2Med.merge(drug, on='Product Name', how='left')
```

```
[ ] C2MedMapping.info()
```

```
[ ] del C2MedMapping['Product Name']
```

# Data Management Plan

01

|

DATA  
DESCRIPTION,  
COLLECTION  
AND REUSING  
EXISTING  
DATA

02

|

DOCUMENTAT  
ION AND  
DATA  
QUALITY

03

|

LEGAL AND ETHICAL  
REQUIREMENTS

04

|

DATA SHARING  
AND LONG-TERM  
PRESERVATION

05

|

DATA  
MANAGEMENT  
RESPONSIBILITIES  
AND RESOURCES