

## Track patient recovery in real-time by processing streaming data

#### BIOMEDICAL DATA DESIGN

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### Content

SAPS II: 17 features

Missing data processing

## 01 SAPS II: 17 features



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## Benchmarking deep learning models on large healthcare datasets

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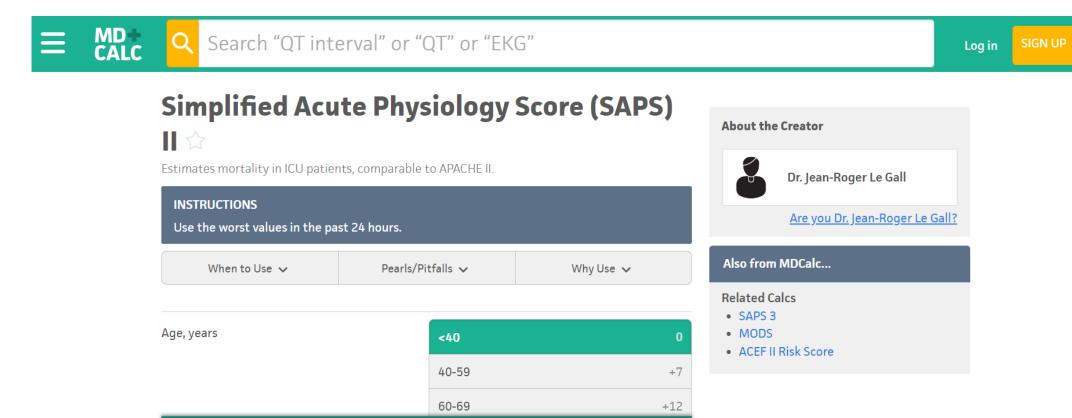
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#### 17 features

#### **How to get Physiology Score**

0 points

SAPS II Score



0.0 %

Copy Results 🗎

In-hospital mortality by SAPS II

Next Steps >>>>

# 02 Missing data processing

#### Data Preprocessing(Heart rate)

#### 1. Extract sub-categories patient id from cardiovascular import numpy as np import matplotlib.pyplot as plt import pandas as pd import data\_toolbox df\_diagnosis = pd.read\_csv('diagnosis.csv') df\_diagnosis.sort\_values(by=['patientunitstayid', 'diagnosisoffset'], inplace=True) # select cardiovascular patients df\_cardiovascular = df\_diagnosis[df\_diagnosis['diagnosisstring'].str.contains('cardiovascular')] # print(df\_cardiovascular) # get shock patient shock\_patient = df\_cardiovascular[df\_cardiovascular['diagnosisstring'].str.contains('shock')] # get ventricular patient ventricular\_patient = df\_cardiovascular[df\_cardiovascular['diagnosisstring'].str.contains('ventricular')] # get chest pain patient chest\_pain\_patient = df\_cardiovascular[df\_cardiovascular['diagnosisstring'].str.contains('chest pain')] # get arrhythmias patient arrhythmias\_patient = df\_cardiovascular[df\_cardiovascular['diagnosisstring'].str.contains('arrhythmias')] df wanted = pd.concat([shock patient, ventricular patient, chest pain patient, arrhythmias patient]) # print(df\_wanted) patient\_id\_all = df\_wanted['patientunitstayid'].unique() patient\_id\_all.sort() print(patient\_id\_all) 143870 151179 151900 ... 3351297 3352230 3352231]

```
# import patient.csv
df_patient = pd.read_csv('patient.csv')
df_patient.sort_values(by=['patientunitstayid'], inplace=True)
df_patient_buf = df_patient[df_patient['patientunitstayid'].isin(patient_id_all)]
df_1time_patient = df_patient_buf[df_patient_buf['unitvisitnumber']==1]
# print(df_1time_patient)

# select the patient id from df_1time_patient
patient_id = df_1time_patient['patientunitstayid'].unique()
print(f'Total number of patients: {len(patient_id)}')
Total number of patients: 915
```

915 Patients valid

#### Data Preprocessing(Heart rate)

```
# define heartrate preprocessing function
def normal_heartrate(num):
    """
    Function to normalize heart rate values.

Parameters:
        num: the originial input value
    Return:
        num: the normalized output value
    """

# Return null values directly
if pd.isna(num):
        return num

# Remove values out of range
elif num > 300 or num < 0:
        return np.nan
# Return normal values directly
else:
        return num</pre>
```

#### Filter the abnormal ones

```
# extract heart rate from df vitalPeriodic
    HR = df_vitalPeriodic[['patientunitstayid', 'observationoffset', 'heartrate']]
    print(f'First 5 rows of HR: \n{HR.head()}')
    # exclude abnormal heart rate values
    HR.loc[:, 'heartrate'] = HR['heartrate'].apply(normal_heartrate)
    # save HR to csv file (uncomment the code to save)
    # HR.to csv('HR.csv', index=False)
    value_position_dict = {}
    first_occurrences = []
    for idx, value in enumerate(HR['patientunitstayid']):
       # if the value is not in the dictionary, add it and create index
       if value not in value_position_dict:
           value position dict[value] = idx
           first_occurrences.append(idx)
    first occurrences.append(len(HR))
    # create first occurrence index for every patient
    HR index = pd.Series(first occurrences)
    print(f'First 5 rows of HR_index: \n{HR_index.head()}')
    # double check the index is correct
    # print(HR.iloc[HR_index].head())
First 5 rows of HR:
     patientunitstayid observationoffset heartrate
628
                143870
                                                44.0
                143870
                                       12
                                                42.0
                143870
                                       17
                                                41.0
                                       22
                143870
                                                41.0
519
                143870
                                       27
                                                41.0
First 5 rows of HR_index:
      158
     1017
     1708
     2501
dtype: int64
```

```
Example: how to use HR & HR_index
   i = 0
   print(f'HeartRate data for patient {i+1}: \n{HR.iloc[HR_index[i]:HR_index[i+1]]}'
 HeartRate data for patient 1:
     patientunitstavid observationoffset heartrate
 628
                143870
                                                44.0
 574
                143870
                                       12
                                                42.0
 543
                143870
                                                41.0
                143870
                                                41.0
 519
                143870
                                                41.0
614
                143870
                                                50.0
 584
                143870
                                                51.0
 578
                143870
                                      782
                                                48.0
                143870
                                      787
                                                48.0
                143870
                                      792
                                                49.0
 [158 rows x 3 columns]
```

Index to search certain patients

