B1-01



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Supervises and Leads Development of the Medical Assistant



Uppalapati Pranita - CED18I062Developer

Works on the development of the Backend of the Application



Hrishikesh R Menon - COE18B024 Lead Designer

Concept Designer and Main Developer of FrontEnd



Preety Banjare - COE17B038 Concept Designer

Concept and UI/UX Designer



Week 1 Progress

EPIONE(Minimum Viable Product)

Bill of Materials

EPIONE is a Virtual Medical Assistant that is available online for clients to access anywhere.

This webapp takes in manually inputted data and

- generates a summarized report when asked by the user for a fixed time basis.
- outputs the risk factor of diabetes based on certain minimum viable parameters as inputs and gives suitable advice on moving forward.

Name of Material	Status	Cost
Kaggle	Downloaded-Tested	Free
Flask	Downloaded and Ready	Free
Jupyter Notebook	Downloaded and Running	Free
HTML,CSS,Bootstrap	Downloaded and Running	Free
Python Libraries	Downloaded and Running	Free
Machine Learning Algorithm	Ready and Working	Free

Week 2 Progress

Project Plan with Appropriate Work Breakdown

1. Data Collection

a. Collecting Proper
 Datasets for our
 MVP,through open
 source websites.

2. Data Visualization

- a. Descriptive Statistics
- b. Statistical analysis
- c. Visualizing data distribution

3. Data Preprocessing

- a. Data Cleaning
- b. DataHandling(missing/c orrupted).

4. Model Creation

- a. Feature Selection
- b. Using Machine LearningAlgorithms we can
 - Risk index generation, predictive modelling.
 - ii. As well as for health care ratings.
 - iii. Performance testing.

5. Web App Development

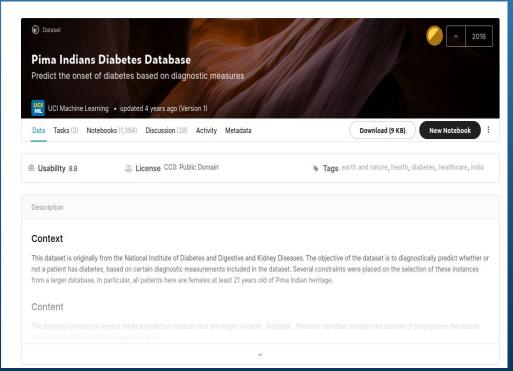
- using HTML,CSS,JavaScript to create a visually appealing web application to house our backend
- b. Using the Flask WebFramework we will build theMVP using our data models.

Week 3 Progress

Technical Schematic

Dataset Collection

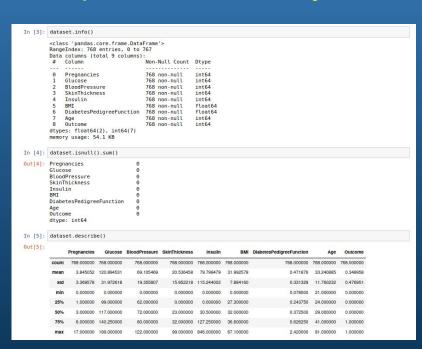




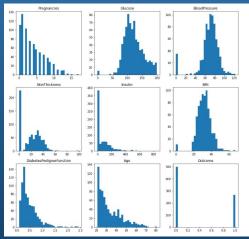
Week 4 Progress

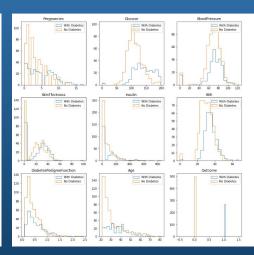
Now that the most suitable available dataset has been collected, week 4 was utilized to understand the dataset

Descriptive Statistics and Analysis



Dataset Visualisation



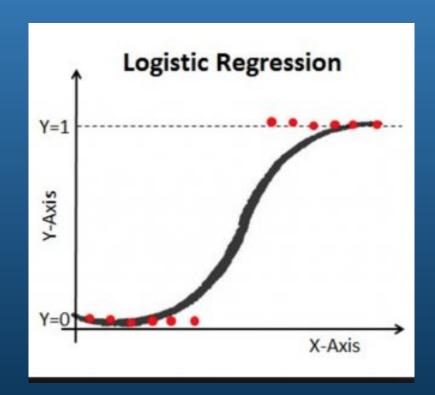


Research on Optimal Algorithm

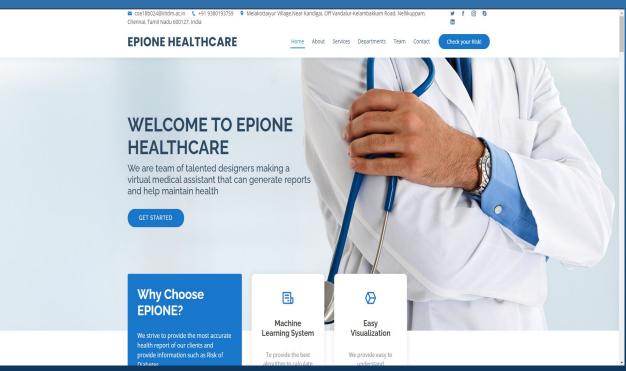
Logistic Regression

Logistic regression is a linear classification method that learns the probability of a sample belonging to a certain class. Logistic regression tries to find the optimal decision boundary that best separates the classes. Logistic regression directly models the posterior probability of P(y|x) by learning the input to output mapping by minimising the error. Logistic regression splits feature space linearly, and typically works reasonably well even when some of the variables are correlated.

Given the correlated feature space which is following a normal distribution after cleaning up of noisy samples, and given that logistic regression is a generalized linear model which works well with correlated features, <u>LOGISTIC REGRESSION</u> was chosen as a more compatible algorithm for the MVP dataset.



Prototyping Phase Conceptual Front End





Desktop Views

Data Cleaning and Handling

Dropping Duplicate Rows

```
In [11]:
    dataset = dataset.drop_duplicates(keep='first')
    dataset.shape

Out[11]: (768, 9)
```

Handling incomplete Data

```
In [12]: dataset = dataset.drop(dataset[dataset['BMI']==0].index)
    dataset = dataset.drop(dataset[dataset['BloodPressure']==0].index)
    dataset = dataset.drop(dataset[dataset['Insulin']==0].index)
    dataset = dataset.drop(dataset[dataset['Glucose']==0].index)
    dataset = dataset.drop(dataset[dataset['SkinThickness']==0].index)
    dataset.shape
```

Feature Selection

Feature Selection

The Pregnancies column contains information on number of pregnancies, but while there is a correlation between diabetes likelihood and number of pregnancies, our data-set does not actually distinguish between males and females. If, for instance, a male is more likely to have diabetes based on other indicators, we don't want the prediction for him having diabetes being marked down just because his number of Pregnancies is 0. To avoid the problem of the predictive model being confounded by this, we will get rid of the Pregnancies column.

There is also the fact that we have very limited understanding on how the DiabetesPedigreeFunction column's values are determined. This column measures an individual's hereditary predisposition to diabetes. Realistically, it is not going to be possible for an individual to input this measurement into our predictive model. We'll thus drop this column as well

```
features = cols.copy()
features.remove('Outcome')
features.remove('DiabetesPedigreeFunction')
features.remove('Pregnancies')

print(features)
['Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI', 'Age']
```

^ Final Features that were selected

Predictive Modelling - Logistic Regression

```
warnings.filterwarnings("ignore")
from sklearn.linear model import LogisticRegression
from sklearn.model selection import cross val score
from sklearn.metrics import accuracy score
from sklearn.model selection import KFold
# Instantiate the model.
weight = {
   1:1.05.
   0:1
log = LogisticRegression(class_weight = weight)
kf = KFold(n splits=6)
score = cross_val_score(log, dataset[features], dataset['Outcome'], cv=kf, scoring='accuracy')
print(score)
print("The mean accuracy is:",score.mean())
[0.81818182 0.71212121 0.72307692 0.83076923 0.87692308 0.78461538]
The mean accuracy is: 0.790947940947941
```

Risk Index Generation

def diabetes_risk_prediction(glucose, bp, skinthickness, insulin, bmi, age):
 indicator list = [glucose, bp, skinthickness, insulin, bmi, age]

We calculate the Posterior probability to evaluate the risk through a scaled metric.

ML Model - Performance testing and Accuracy

Re-instantiate the logistics regression and model and re-fit the data.

```
def diabetes_risk_prediction2(glucose, bp, skinthickness, insulin, bmi, age):
    weight = {
    1:1.05,
    0:1}

    log_model2 = LogisticRegression(class_weight = weight)
    log_model2.fit(dataset[features], dataset['Outcome'])

    indicator_list = [glucose, bp, skinthickness, insulin, bmi, age]
    predictions = log_model2.predict_proba(np.array(indicator_list).reshape(1, -1))
    risk = predictions[0,1]
    return risk

diabetes_risk_prediction2(100,80,30,120,30,45) #Test array.

0.22443666622161737
```

```
kf = KFold(n_splits=6)
score = cross_val_score(log, dataset[features], dataset['Outcome'], cv=kf, scoring='accuracy')
print(score)
print("The mean accuracy is:",score.mean())

[0.81818182 0.71212121 0.72307692 0.83076923 0.87692308 0.78461538]
The mean accuracy is: 0.790947940947941

Using KFold cross validation with slightly more weight placed on a positive check for Diabetes, with 6 folds, we get an accuracy of around 79%.
```



50	110
10	100
13	24
ou are probably in good heal	th, keep it up.

DEMONSTRATION

EPIONE Check your Risk!
Fill in Details so we can calculate Health Risk Index
200
150
50
140
29
25
See a doctor as soon as you can . You might be on the way to developing diabetes if you don't change your lifestyle. Your Diabetes Risk Index is 41.00/50.

Desktop Views

Mobile View **B1-01**

Design Pipeline

·Data Collection and Preprocessing

(Status: Completed) --20/01/2021

· Data Analysis and Visualization

(Status: Completed) -- 27/01/2021

· Data Trimming and Cleaning

(Status: Completed)--03/02/2021

· Algorithm Selection

(Status: Completed)--03/02/2021

FrontEnd Work

(Status: Completed)--03/03/2021

·Data Model Training and Testing

(Status: Completed)--17/02/2021

·Performance Testing and Bug Identifying

(Status: Completed)--24/02/2021

·Backend Implementation and UI/UX Design

(Status: Completed)--10/03/2021

•Front to Back Integration

(Status: Completed)--17/03/2021

•Testing, Optimizing and Deployment

(Status: Completed)--24/03/2021

Click Here For Detailed MVP Report