

A Mini- Project Report
on
“Diabetes Prediction Tool”

Submitted to the
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In partial fulfillment for the award of the Degree of
Bachelor of Engineering
in
Information Technology
by

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CERTIFICATE

This is to certify that the project report entitled

DIABETES PREDICTION TOOL

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is a bonafide work carried out by them under the supervision of Prof. R. R.Chhajed and it is approved for the partial fulfillment of the requirement of **Computer Laboratory -X** for the award of the Degree of Bachelor of Engineering (Information Technology).

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ABSTRACT

The recent covid-19 pandemic has taken its toll on the population across the world. Especially, the elderly and the people with comorbidities like Asthma and Diabetes are especially at risk. Having diabetes also increase the chances of contracting the Mucormycosis, a.k.a. the black fungus. Having such comorbidities also delay the healing procedure of covid-19 infected patients and hence, should not go unknown and untreated.

Therefore, through this project, we wish to allow the general population to gain an insight into their diabetes situation to help them manage the disease better; and effectively help them lead a better lifestyle.

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TOPICS

1. Introduction

COVID-19 remains the biggest pandemic the world has ever witnessed. India is at the forefront with a huge number of cases and COVID-related deaths. As per reports, elderly people are highly susceptible to the Corona virus and those who have diabetes are even more prone. In general, people with diabetes are more likely to have more severe symptoms and complications when infected with any virus. Your risk of getting very sick from COVID-19 is likely to be lower if your diabetes is well-managed.

Therefore, finding out if a person has diabetes could be a very important step in understanding how severe would your COVID infection be, if you are ever affected by the virus. Lab methods generally take a while to deliver reports stating your diabetic status.

Hence, to provide a better intuition about the same problem, we propose a diabetes prediction tool. This tool takes 8 different inputs from the user and on the basis of that tells the user whether he or she is possibly diabetic. This tool makes use of Random Forest, which is a supervised machine learning technique for classifying whether the person has diabetes or not. As the prediction is purely based on numbers, this tool cannot be used to obtain a final verdict and in doubt the user is advised to contact a physician.

2. Scope and Objective

Scope of our project is to create a web application with considerable accuracy and speed to predict the condition of diabetes in a patient based on the outcomes of his results on certain parameters. The web application will also provide insights into the data gathered by the exploration of the dataset.

Objective of the project is to educate the populace and provide insights into the probability of one having diabetes and the related conditions. The insights into the data explored will also help in understanding the typical signs of one having diabetes and thus, enabling them to tackle the issue earlier.

3. System architecture and flow

We have considered using the Flask framework for the creation of the website. This mainly resorts to the fact that using the same environment of Python for data model and website ensures high cohesion. Flask is a light-weight micro web framework written in Python, which makes use of additional extensions if needed; to provide extra features like form validations, database abstraction layers, etc. It also contains its own light development server and debugger, support for Unit testing, RESTful service dispatching with support for client-side secure cookies.

We have created a website of two pages. First page of the site is an information page, which shows the insights gained after exploration of the data, and a link to the predictor on the second page. First page is available at decorator (/) which redirects to (/home). Second page contains a form which will accept the values input by the user under various heads, which are the features of our dataset, namely Skin Thickness, Blood pressure, BMI, Diabetes Pedigree function, age, etc. This form submits to the function predictor (decorator /predictor). In this function, the form data (the features input by the user) are extracted.

A pickle dump file (.pkl) was created for the trained model. This file basically contains the pretrained model. The values submitted through the form are scaled with one-hot encoding which is defined in the pickle dump file.

These features are then used to predict the value of the output, which outputs 1 for diabetic and 0 for non-diabetic values.

The final value is shown on the same page, just below the submit button.

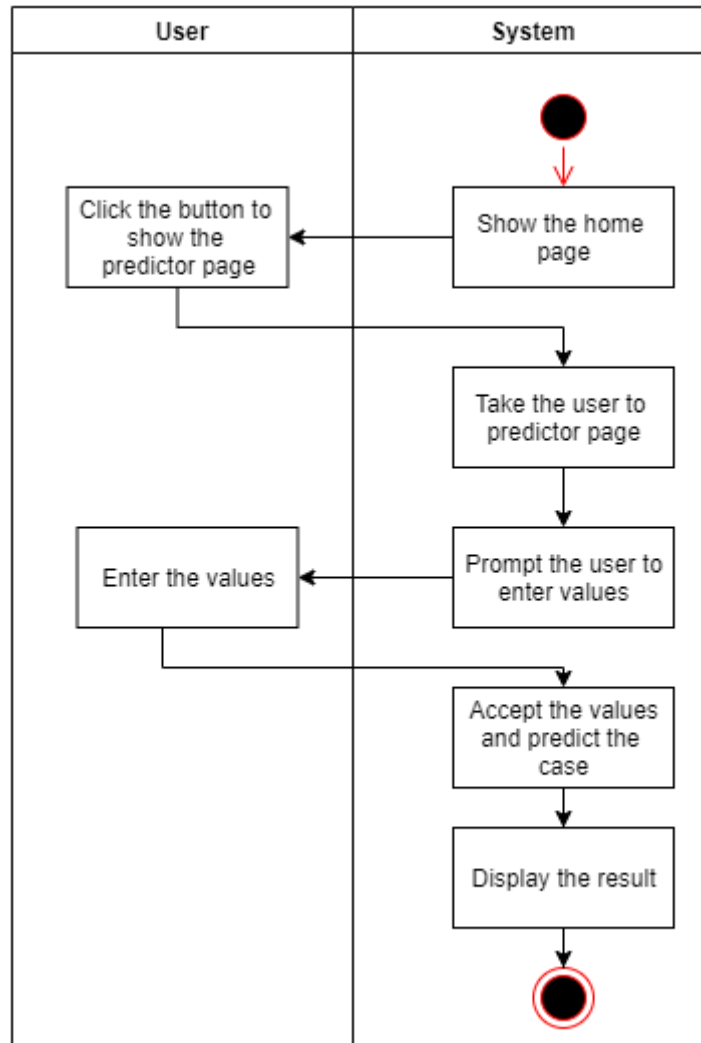


Figure 1: Flow of the system

4. Code and snapshots

Code for data model:

4.1 Reading data

```
[ ] from google.colab import files
    uploaded_data = files.upload()

Choose Files No file chosen Upload widget is only available when the cell has been executed in the current browser session. Please rerun this cell to enable.
Saving diabetes_pima.csv to diabetes_pima (1).csv

[ ] diabetes_df = pd.read_csv(io.BytesIO(uploaded_data['diabetes_pima.csv']))

[ ] diabetes_df.head()
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	23.3	0.672	32	1
3	1	89	66	23	94	28.1	0.167	21	0

4.2 Preprocessing

4.2.1 Replacing 0's in columns with average value

```
[ ] nonzerocols = ['Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI']

[ ] diabetes_df[nonzerocols] = diabetes_df[nonzerocols].replace(0, np.NaN)

[ ] diabetes_df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 768 entries, 0 to 767
Data columns (total 9 columns):
#   Column                Non-Null Count  Dtype  
---  -
0   Pregnancies            768 non-null   int64  
1   Glucose                763 non-null   float64 
2   BloodPressure          733 non-null   float64 
3   SkinThickness          541 non-null   float64 
4   Insulin                394 non-null   float64 
5   BMI                    757 non-null   float64 
6   DiabetesPedigreeFunction 768 non-null   float64 
7   Age                    768 non-null   int64  
8   Outcome                768 non-null   int64  
dtypes: float64(6), int64(3)
memory usage: 54.1 KB
```

```
[ ] for column in nonzerocols:
    diabetes_df[column] = diabetes_df[column].replace(np.NaN, int(diabetes_df[column].mean(skipna = True)))
```

4.2.2 Detecting and removing outliers

```
[ ] def thresholdLimits(df,var):
    quartile1 = df[var].quantile(0.25)
    quartile3 = df[var].quantile(0.75)
    interquartile_range = quartile3 - quartile1
    up_limit = quartile3 + (1.5 * interquartile_range)
    low_limit = quartile1 - (1.5 * interquartile_range)
    return low_limit, up_limit

[ ] def doOutliersExist(df, var):
    low_limit, up_limit = thresholdLimits(df, var)
    if df[(df[var] < low_limit) | (df[var] > up_limit)].any(axis = None):
        print(var, "yes")
    else:
        print(var, "no")

[ ] def replaceWithThresholds(df, numeric_columns):
    for column in numeric_columns:
        low_limit, up_limit = thresholdLimits(df,column)
        df.loc[(df[column] < low_limit), column] = low_limit
        df.loc[(df[column] > up_limit), column] = up_limit

[ ] for column in diabetes_df.columns:
    doOutliersExist(diabetes_df, column)
```

```
[ ] for column in diabetes_df.columns:
    doOutliersExist(diabetes_df, column)

Pregnancies yes
Glucose no
BloodPressure yes
SkinThickness yes
Insulin yes
BMI yes
DiabetesPedigreeFunction yes
Age yes
Outcome no

[ ] replaceWithThresholds(diabetes_df,diabetes_df.columns)

[ ] for column in diabetes_df.columns:
    doOutliersExist(diabetes_df, column)

Pregnancies no
Glucose no
BloodPressure no
SkinThickness no
Insulin no
BMI no
DiabetesPedigreeFunction no
Age no
Outcome no
```

4.2.3 Converting BMI into a categorical column

```
[ ] BMICategory = pd.Series(["Underweight", "Normal", "Overweight", "Obesity 1", "Obesity 2", "Obesity 3"], dtype = "category")

diabetes_df["BMICategory"] = BMICategory

diabetes_df.loc[diabetes_df["BMI"] < 18.5, "BMICategory"] = BMICategory[0]
diabetes_df.loc[(diabetes_df["BMI"] > 18.5) & (diabetes_df["BMI"] <= 24.9), "BMICategory"] = BMICategory[1]
diabetes_df.loc[(diabetes_df["BMI"] > 24.9) & (diabetes_df["BMI"] <= 29.9), "BMICategory"] = BMICategory[2]
diabetes_df.loc[(diabetes_df["BMI"] > 29.9) & (diabetes_df["BMI"] <= 34.9), "BMICategory"] = BMICategory[3]
diabetes_df.loc[(diabetes_df["BMI"] > 34.9) & (diabetes_df["BMI"] <= 39.9), "BMICategory"] = BMICategory[4]
diabetes_df.loc[diabetes_df["BMI"] > 39.9, "BMICategory"] = BMICategory[5]

[ ] diabetes_df.head()
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome	BMICategory
0	6.0	148.0	72.0	35.0	155.0	33.6	0.627	50.0	1.0	Obesity 1
1	1.0	85.0	66.0	29.0	155.0	26.6	0.351	31.0	0.0	Overweight
2	8.0	183.0	64.0	29.0	155.0	23.3	0.672	32.0	1.0	Normal
3	1.0	89.0	66.0	23.0	94.0	28.1	0.167	21.0	0.0	Overweight
4	0.0	137.0	40.0	35.0	168.0	43.1	1.200	33.0	1.0	Obesity 3

4.2.4 One-hot encoding BMI categories

```
[ ] def oneHotEncoder(dataframe, categorical_columns):
    original_columns = list(dataframe.columns)
    dataframe = pd.get_dummies(dataframe, columns = categorical_columns, drop_first = True)
    new_columns = [col for col in dataframe.columns if col not in original_columns]
    return dataframe, new_columns

[ ] diabetes_df, new_cols_ohc = oneHotEncoder(diabetes_df, ["BMICategory"])
diabetes_df = diabetes_df.drop(["BMI"], axis = 1)
new_cols_ohc

['BMICategory_Obesity 1',
 'BMICategory_Obesity 2',
 'BMICategory_Obesity 3',
 'BMICategory_Overweight',
 'BMICategory_Underweight']

[ ] diabetes_df.head()
```

	Age	Outcome	BMICategory_Obesity 1	BMICategory_Obesity 2	BMICategory_Obesity 3	BMICategory_Overweight	BMICategory_Underweight
0	50.0	1.0	1	0	0	0	0
1	31.0	0.0	0	0	0	1	0
2	32.0	1.0	0	0	0	0	0
3	21.0	0.0	0	0	0	1	0
4	33.0	1.0	0	0	0	0	0

4.3 Hyperparameter-tuned Model Training and Result

```
model = RandomForestClassifier(n_estimators = 50,max_depth = 12.0,min_samples_split=2,random_state=31)
model.fit(X_train, y_train)
y_pred = model.predict(X_test)
print(accuracy_score(y_test, y_pred) * 100)
cm = confusion_matrix(y_test, y_pred)
print("\n", cm)
```

83.11688311688312

```
[[94 13]
 [13 34]]
```

BACKEND

4.4 Code for extracting the values through the form:

Note: This is the code inside the predictor function.

```
if request.method == "POST" and form.validate():
    pregnancies = form.pregnancies.data
    glucose = form.glucose.data
    bp = form.bp.data
    st = form.st.data
    insulin = form.insulin.data
    bmi = form.bmi.data
    pedigreeFunc = form.diabetesPedigree.data
    age = form.bmi.data
```

4.5 Code for loading the pickle file and predict the output:

```
scalerLoad = pickle.load(open("scaler.pkl", "rb"))

modelLoad = pickle.load(open("randomForestModel.sav", "rb"))
featureList = np.array([pregnancies, glucose, bp, st, insulin, pedigreeFunc, age]).reshape(1, -1)

featureList = scalerLoad.transform(featureList)

finalFeatures = np.concatenate((featureList, BMIVar), axis = 1)
value = modelLoad.predict(finalFeatures)
print("Value = {}".format(value))
```

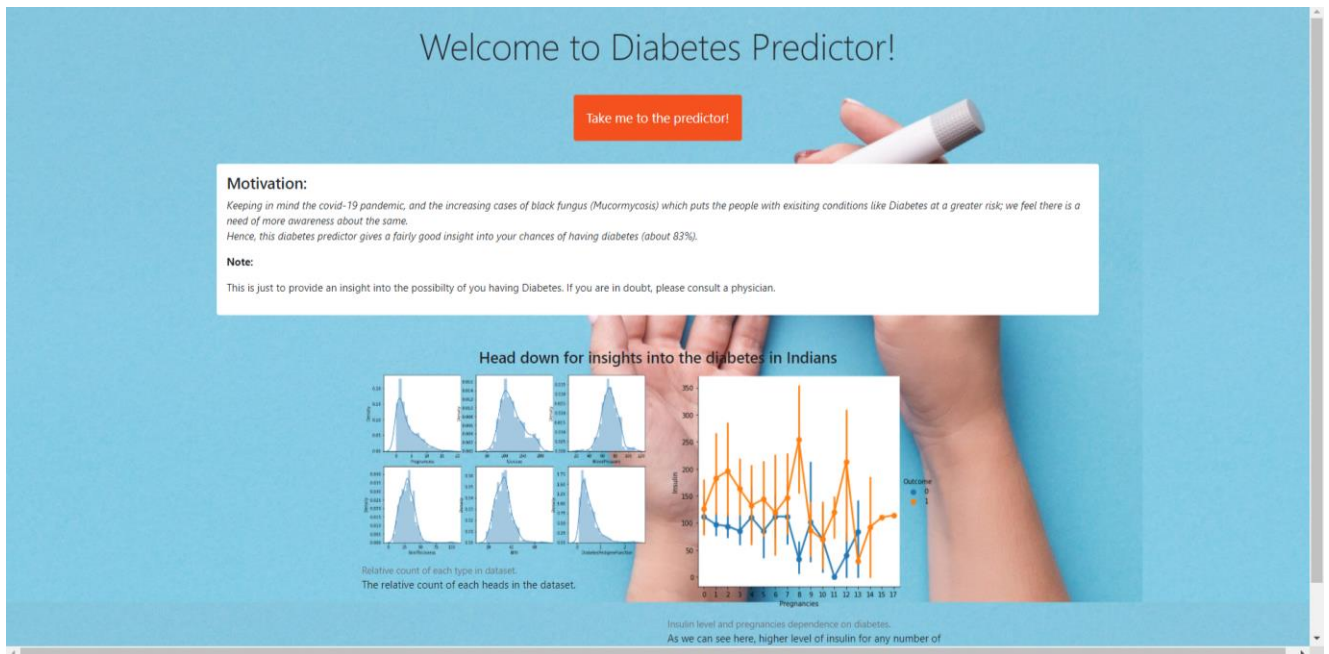
The modelLoad is the loaded model from the pickle file; whereas scalerLoad variable is the scalar defined and trained for the dataset using sklearn.

Finally, the numpy array of features; finalFeatures; is used to predict the outcome from the given data.

Snapshots of the project:

4.6 UI

4.6.1 Homepage



4.6.2 Predictor page

The screenshot shows the predictor page of the 'Diabetes Predictor' application. The background is a light blue gradient with a hand holding a white marker. The main heading is 'Enter your values for prediction here!'. Below it is a form with input fields for various features: 'Number of pregnancies' (1), 'Your glucose level' (148), 'Your blood pressure' (80), 'Your Skin Thickness' (30), 'Insulin' (160), 'Diabetes Pedigree function' (0.627), 'BMI' (29.32), and 'Age' (50). An orange 'Submit' button is located below the form. Below the form, a red message says 'Congratulations!! There is a 83% chance you are not a diabetic!'. At the bottom, there is a red button that says 'Go back to home page!'. The footer text reads 'Made with ❤️ by Vedant and Sahil!'.

5. Results

To get an estimate of the highest accuracy achievable on our dataset, we implemented a number of machine learning algorithms for the same. Out of all the algorithms tried, Random Forest performed the best with a test accuracy of 83%. The open-source dataset we used is called the PIMA Indians Diabetes Dataset, which is a standard dataset used for diabetes related problem statements.

Model	Test Accuracy
Naive Bayes	33.11%
Decision Tree	70.12%
K-Nearest Neighbors	79.22%
XGBoost	79.8%
SVM	80.5%
Logistic Regression	81.16%
Random Forest	83.11%

Following is an empirical analysis of several supervised machine learning algorithms we tried:

As Random Forest performed the best as compared to others, we used other metrics to get a better idea about the model's classification ability, for which we used confusion matrix.

Confusion Matrix:

	Predicted True	Predicted False
Actually True	94 (TP)	13 (FN)
Actually False	13 (FP)	34 (TN)

$$\text{Precision} = \text{TP}/(\text{TP}+\text{FP}) = 94/(94+13) = 0.878$$

$$\text{Recall} = \text{TP}/(\text{TP}+\text{FN}) = 94/(94+13) = 0.878$$

$$\text{F1-score} = 2*\text{precision}*\text{recall}/(\text{precision} + \text{recall}) = 0.878$$

6. Conclusion and Future Scope

Hence, as COVID-19 remains the biggest health problem faced by the world today, knowing whether or not a person has diabetes can help him assess his health and be prepared in advance for steps to be taken, as diabetics face more complications than non-diabetics. To get a numerical prediction, we employed several supervised machine learning algorithms, out of which Random Forest proved to deliver the highest accuracy of 83%. This tool uses traditional machine learning. If more data is available, using neural networks for classification would be a better solution in the future. Additionally, also making an integrated app for obtaining preliminary predictions of several diseases related to kidney, liver, heart, etc. would prove to provide a one-stop solution to a number of problems.

GitHub link for project: <https://github.com/ItsNaSa/Diabetes-Predictor>

7. References

1. <https://machinelearningmastery.com/>
2. <https://scikit-learn.org/stable/>
3. <https://www.datacamp.com/community/tutorials/pickle-python-tutorial>
4. <https://www.kaggle.com/uciml/pima-indians-diabetes-database>
5. <https://flask.palletsprojects.com/en/1.1.x/pdf/>
6. [Deploy a machine learning model using flask](#)