

Thyroid Disease Classification Using ML

TEAM SIZE : 4

TEAM LEADER : KRISHNAMOORTHY D

TEAM MEMBERS : MUGILA R

SHANMUGAPRIYA T

SANJAY S

INTRODUCTION

Overview

The Thyroid gland is a vascular gland and one of the most important organs of the human body. This gland secretes two hormones which help in controlling the metabolism of the body.

The two types of Thyroid disorders are Hyperthyroidism and Hypothyroidism. When this disorder occurs in the body, they release certain types of hormones into the body which imbalances the body's metabolism. A thyroid-related Blood test is used to detect this disease but it is often blurred and noise will be present. Data cleansing methods were used to make the data primitive enough for the analytics to show the risk of patients getting this disease. Machine Learning plays a very deciding role in disease prediction. Machine Learning algorithms, SVM - support vector machine, Random Forest Classifier, XGB Classifier and ANN - Artificial Neural Networks are used to predict the patient's risk of getting thyroid disease. The web app is created to get data from users to predict the type of disease.

Purpose

The Thyroid gland is a vascular gland and one of the most important organs of the human body. This gland secretes two hormones which help in controlling the metabolism of the body.

The two types of Thyroid disorders are Hyperthyroidism and Hypothyroidism. When this disorder occurs in the body, they release certain types of hormones into the body which imbalances the body's metabolism. A thyroid-related Blood test is used to detect this disease but it is often blurred and noise will be present. Data cleansing methods were used to make the data primitive enough for the analytics to show the risk of patients getting this disease. Machine Learning plays a very deciding role in disease prediction. Machine Learning algorithms, SVM - support vector machine, Random Forest Classifier, XGB Classifier and ANN - Artificial Neural Networks are used to predict the patient's risk of getting thyroid disease. The web app is created to get data from users to predict the type of disease.

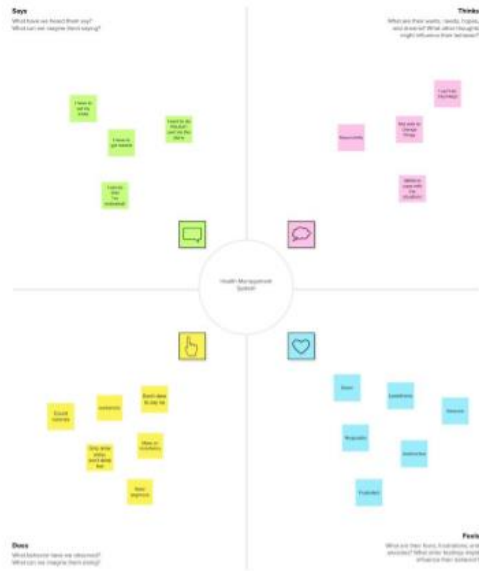
Problem definition & design thinking

Empathy Map

Empathy map

Use this framework to develop a deep, shared understanding and empathy for other people. An empathy map helps describe the aspects of a user's experience, needs and pain points, to quickly understand your users' experience and mindset.

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Ideation & Brainstorming Map

Ideation & Brainstorming Map

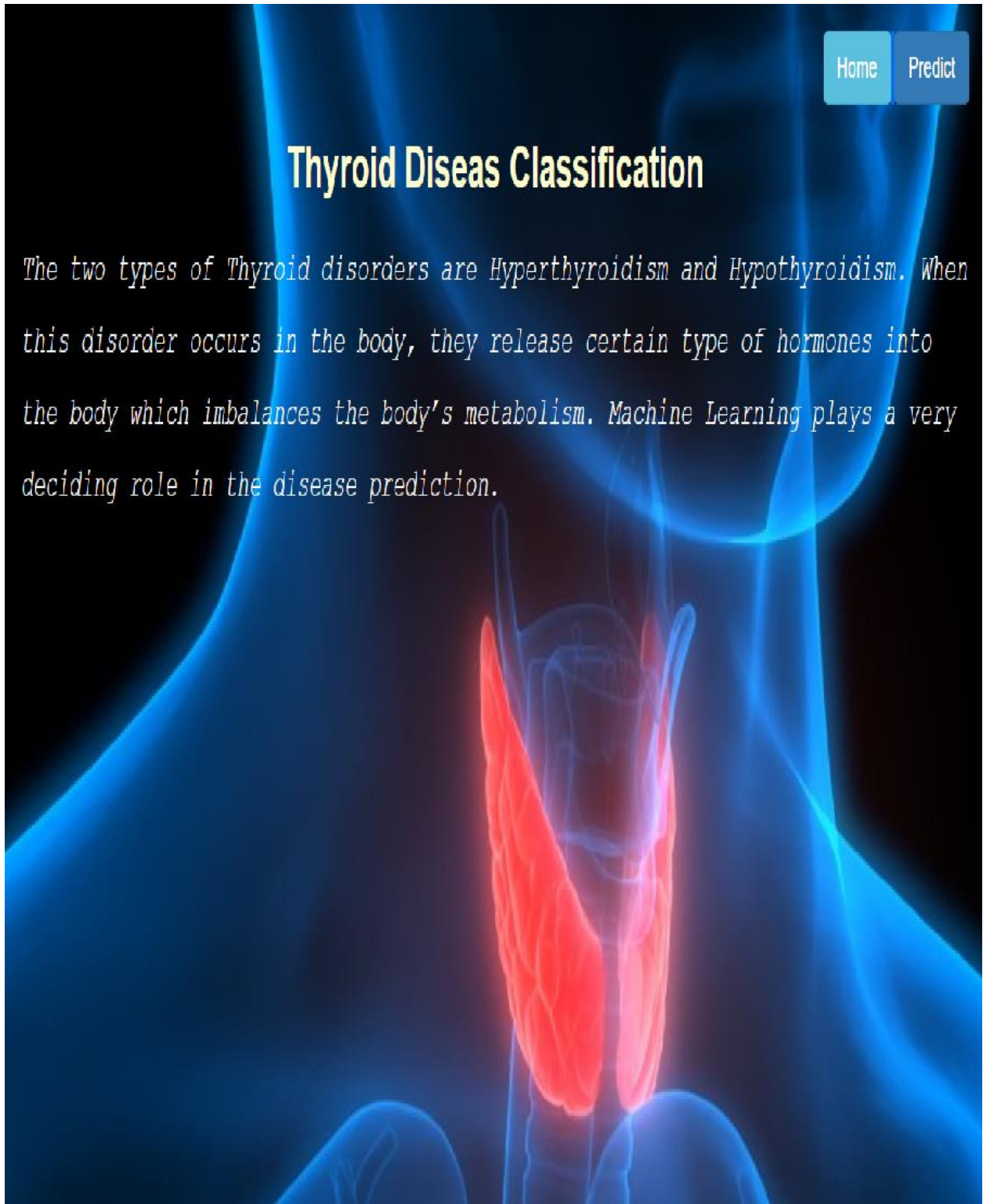


RESULT

[Home](#)[Predict](#)

Thyroid Diseas Classification

The two types of Thyroid disorders are Hyperthyroidism and Hypothyroidism. When this disorder occurs in the body, they release certain type of hormones into the body which imbalances the body's metabolism. Machine Learning plays a very deciding role in the disease prediction.



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Thyroid Disease Classification

goitre

Male ▾

tumor

Male ▾

hypopituitary

Male ▾

psych

Male ▾

TSH

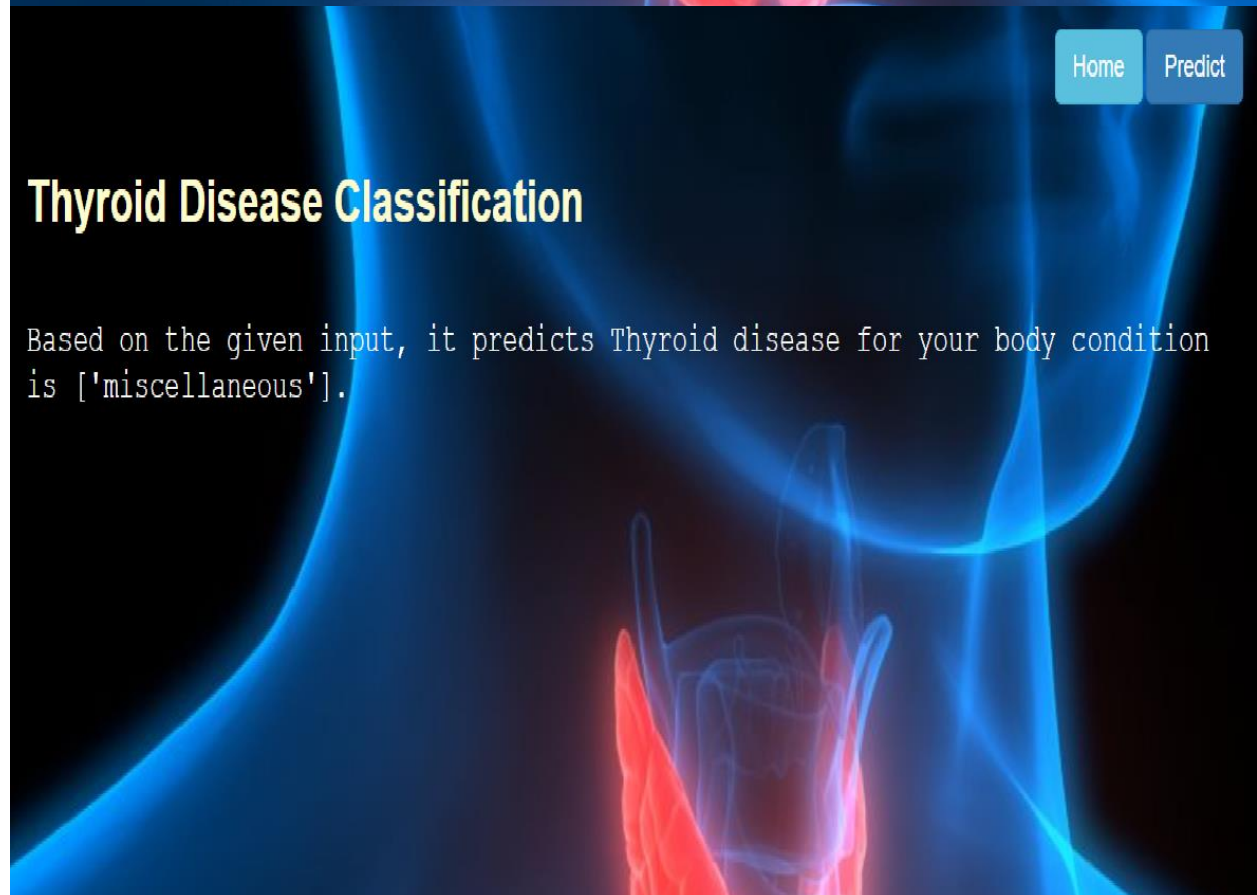
TSH

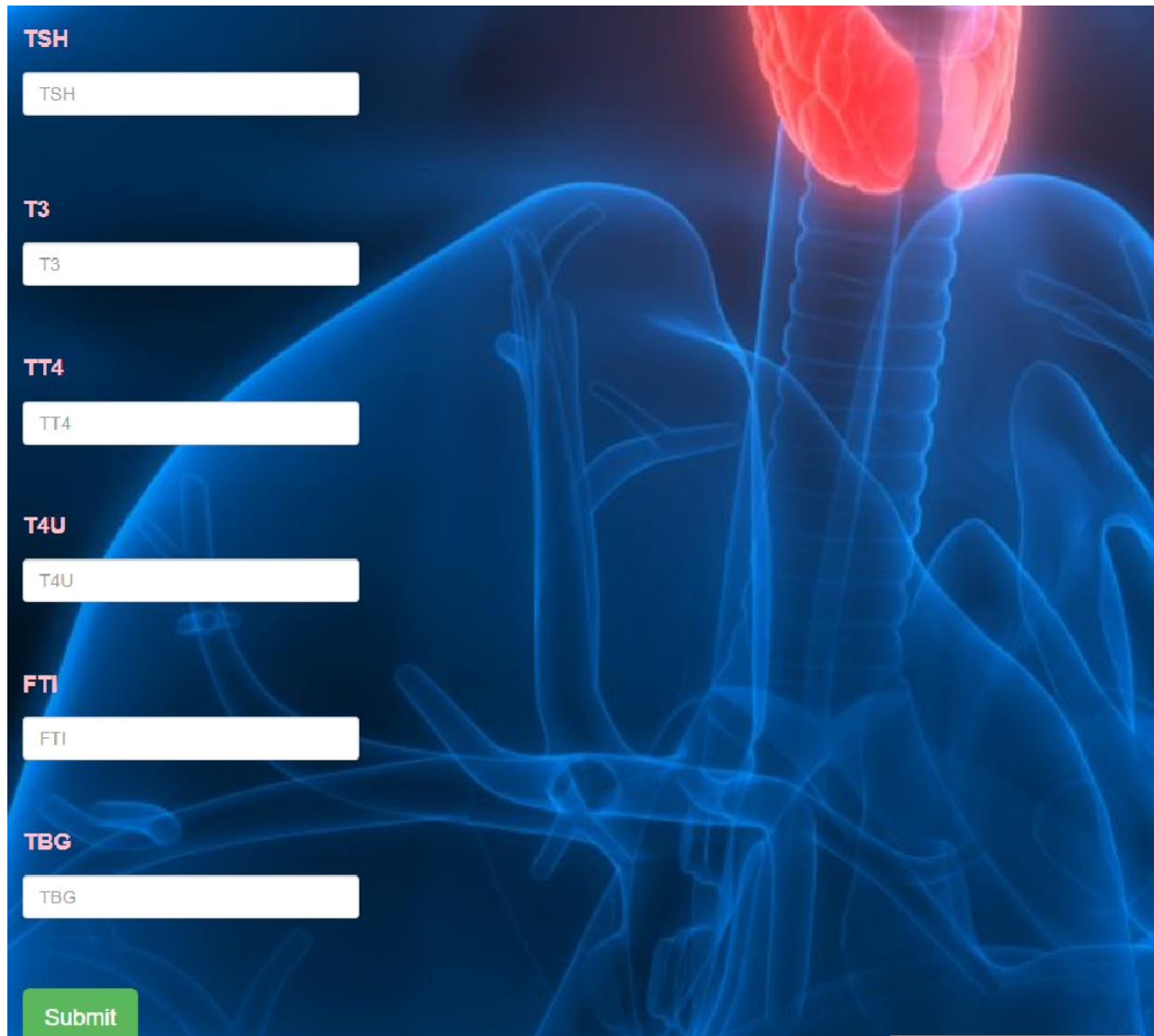


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Thyroid Disease Classification

Based on the given input, it predicts Thyroid disease for your body condition is ['miscellaneous'].





TSH

T3

TT4

T4U

FTI

TBG

Submit

ADVANTAGES & DISADVANTAGES

Advantages:

- The thyroid gland is one of the body's most visible endocrine glands. Its size is determined by the individual's age, gender, and physiological

states, such as pregnancy or lactation. It is divided into two lobes (right and left) by an isthmus (a band of tissue).

- It is imperceptible in everyday life yet can be detected when swallowing. The thyroid hormones T4 and T3 are needed for normal thyroid function. These hormones have a direct effect on the body's metabolic rate. It contributes to the stimulation of glucose, fatty acid, and other molecule consumption.
- Additionally, it enhances oxygen consumption in the majority of the body's cells by assisting in the processing of uncoupling proteins, which contributes to an improvement in the rate of cellular respiration.

Disadvantages :

- Thyroid conditions are difficult to detect in test results, and only trained professionals can do so. However, reading such extensive reports and predicting future results is difficult.
- The thyroid disease can then be easily identified based on the symptoms in the patient's history. Currently, models are evaluated using accuracy metrics on a validation dataset that is accessible.

- **Social Impact:-** Untreated/undetected thyroid disease is more dangerous at times it can lead to fatal of the person. So, we can detect it at the earliest then people can get treatment and get cured.
- **Business Model/Impact:-** We can make this application public, offer services as a subscription based or can collaborate with healthcare centres or specialists.

APPLICATION

- Open anaconda prompt from the start menu
- Navigate to the folder where your python script is.
- Now type “python app.py” command
- Navigate to the localhost where you can view your web page.
- Click on the predict button from the top right corner, enter the inputs, click on the submit button, and see the result/prediction on the web

CONCLUSION

Saving the best model after comparing its performance using different evaluation metrics means selecting the model with the highest performance and saving its weights and configuration. This can be useful in avoiding the need to retrain the model every time it is needed and also to be able to use it in the future.

FUTURE SCOPE

In this section, we will be building a web application that is integrated to the model we built. A UI is provided for the uses where he has to enter the values for predictions. The enter values are given to the saved model and prediction is showcased on the UI.

This section has the following tasks

- Building HTML Pages
- Building server side script

APPENDIX

SOURCE CODE :

```
data = pd.read_csv("data.csv")
```

```
data.head()
```

	age	sex	on_thyroxine	query_on_thyroxine	on_antithyroid_meds	sick	pregnant	thyroid_surgery	I131_treatment	query_hypothyroid	...	TT4	T4U_measu
0	29	F	f		f	f	f	f	f	f	t ...	NaN	
1	29	F	f		f	f	f	f	f	f	f ...	128.0	
2	41	F	f		f	f	f	f	f	f	f ...	NaN	
3	36	F	f		f	f	f	f	f	f	f ...	NaN	
4	32	F	f		f	f	f	f	f	f	f ...	NaN	

5 rows × 31 columns

```
data.shape
```

(9172, 31)

```
#re-mapping target values to diagnostic group
diagnoses = {'A': 'hyperthyroid conditions',
             'B': 'hyperthyroid conditions',
             'C': 'hyperthyroid conditions',
             'D': 'hyperthyroid conditions',
             'E': 'hypothyroid conditions',
             'F': 'hypothyroid conditions',
             'G': 'hypothyroid conditions',
             'H': 'hypothyroid conditions',
             'I': 'binding protein',
             'J': 'binding protein',
             'K': 'general health',
             'L': 'replacement therapy',
             'M': 'replacement therapy',
             'N': 'replacement therapy',
             'O': 'antithyroid treatment',
             'P': 'antithyroid treatment',
             'Q': 'antithyroid treatment',
             'R': 'miscellaneous',
             'S': 'miscellaneous',
             'T': 'miscellaneous'}

data['target'] = data['target'].map(diagnoses) #remapping
```

```
#Removing Redundant attributes from dataset
```

```
data.drop(['TSH_measured', 'T3_measured', 'TT4_measured', 'T4U_measured', 'FTI_measured', 'TBG_measured', 'referral_source', 'patient_id'], axis=1, inplace=True)
```

```
data.dropna(subset=['target'],inplace=True)
```

```
data['target'].value_counts()
```

```
hypothyroid conditions    593  
general health            436  
binding protein          376  
replacement therapy      336  
miscellaneous            281  
hyperthyroid conditions  182  
antithyroid treatment    33  
Name: target, dtype: int64
```

```
x['sex'].unique()
```

```
array(['F', 'M', nan], dtype=object)
```

```
x['sex'].replace(np.nan, 'F', inplace=True)
```

```
x['sex'].value_counts()
```

```
F    1701  
M     536  
Name: sex, dtype: int64
```

```
data.isnull().sum()
```

```
age          0
sex          307
on_thyroxine  0
query_on_thyroxine  0
on_antithyroid_meds  0
sick         0
pregnant     0
thyroid_surgery  0
I131_treatment  0
query_hypothyroid  0
query_hyperthyroid  0
lithium      0
goitre       0
tumor        0
hypopituitary  0
psych        0
TSH_measured  0
TSH          842
T3_measured   0
T3           2604
TT4_measured  0
TT4          442
T4U_measured  0
T4U          809
FTI_measured  0
FTI          802
TBG_measured  0
TBG          8823
referral_source  0
target       0
patient_id   0
dtype: int64
```

```
#Checking whether the age above 100
data[data.age>100]
```

```
age  sex  on_thyroxine  query_on_thyroxine  on_antithyroid_meds  sick  pregnant  thyroid_surgery  I131_treatment  query_hypothyroid  ...  tumor  hypopituitary
```

```
0 rows x 23 columns
```

x

	age	sex	on_thyroxine	query_on_thyroxine	on_antithyroid_meds	sick	pregnant	thyroid_surgery	l131_treatment	query_hypothyroid	...	goitre	tumor
4	32.0	F	f	f	f	f	f	f	f	f	f ...	f	f
18	63.0	F	t	f	f	t	f	f	f	f	f ...	f	f
32	41.0	M	f	f	f	f	f	f	f	f	f ...	f	f
33	71.0	F	t	f	f	f	f	f	f	f	f ...	f	f
39	55.0	F	t	f	f	f	f	f	f	f	t ...	f	f
...
9153	64.0	M	f	f	f	f	f	f	f	f	f ...	f	f
9157	60.0	M	f	f	t	f	f	f	f	f	f ...	f	f
9158	64.0	M	f	f	f	f	f	f	f	f	t ...	f	f
9162	36.0	F	f	f	f	f	f	f	f	f	f ...	f	f
9169	69.0	M	f	f	f	f	f	f	f	f	f ...	f	f

2237 rows × 22 columns



```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import tensorflow
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Layer, Dense, Dropout
```