

```
import pandas as pd
import matplotlib as plt
import seaborn as sns
```

```
df = pd.read_csv(r"C:\Users\susmi\Desktop\test_counter\ml_code\
hypothyroid.csv")
```

```
df.head()
```

	age	sex	on thyroxine	query on thyroxine	on antithyroid medication
sick \					
0	41	F	f	f	f
f					
1	23	F	f	f	f
f					
2	46	M	f	f	f
f					
3	70	F	t	f	f
f					
4	70	F	f	f	f
f					

	pregnant	thyroid surgery	I131 treatment	query hypothyroid	... TT4
measured \					
0	f	f	f	f	...
t					
1	f	f	f	f	...
t					
2	f	f	f	f	...
t					
3	f	f	f	f	...
t					
4	f	f	f	f	...
t					

	TT4	T4U measured	T4U	FTI measured	FTI	TBG measured	TBG	referral
source \								
0	125	t	1.14	t	109	f	?	
SVHC								
1	102	f	?	f	?	f	?	
other								
2	109	t	0.91	t	120	f	?	
other								
3	175	f	?	f	?	f	?	
other								
4	61	t	0.87	t	70	f	?	
SVI								

```
binaryClass
0 P
```

```
1      P
2      P
3      P
4      P
```

```
[5 rows x 30 columns]
```

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 3772 entries, 0 to 3771
```

```
Data columns (total 30 columns):
```

#	Column	Non-Null Count	Dtype
0	age	3772 non-null	object
1	sex	3772 non-null	object
2	on thyroxine	3772 non-null	object
3	query on thyroxine	3772 non-null	object
4	on antithyroid medication	3772 non-null	object
5	sick	3772 non-null	object
6	pregnant	3772 non-null	object
7	thyroid surgery	3772 non-null	object
8	I131 treatment	3772 non-null	object
9	query hypothyroid	3772 non-null	object
10	query hyperthyroid	3772 non-null	object
11	lithium	3772 non-null	object
12	goitre	3772 non-null	object
13	tumor	3772 non-null	object
14	hypopituitary	3772 non-null	object
15	psych	3772 non-null	object
16	TSH measured	3772 non-null	object
17	TSH	3772 non-null	object
18	T3 measured	3772 non-null	object
19	T3	3772 non-null	object
20	TT4 measured	3772 non-null	object
21	TT4	3772 non-null	object
22	T4U measured	3772 non-null	object
23	T4U	3772 non-null	object
24	FTI measured	3772 non-null	object
25	FTI	3772 non-null	object
26	TBG measured	3772 non-null	object
27	TBG	3772 non-null	object
28	referral source	3772 non-null	object
29	binaryClass	3772 non-null	object

```
dtypes: object(30)
```

```
memory usage: 884.2+ KB
```

```
df['TBG'].unique()
```

```
array(['?'], dtype=object)
```

```
df = df.drop('referral source', axis =1)
```

```
df.head()
```

```
age sex on thyroxine query on thyroxine on antithyroid medication
sick \
0 41 F f f f
1 23 F f f f
2 46 M f f f
3 70 F t f f
4 70 F f f f
```

```
pregnant thyroid surgery I131 treatment query hypothyroid ... T3
\
0 f f f f ... 2.5
1 f f f f ... 2
2 f f f f ... ?
3 f f f f ... 1.9
4 f f f f ... 1.2
```

```
TT4 measured TT4 T4U measured T4U FTI measured FTI TBG measured
TBG \
0 t 125 t 1.14 t 109 f
?
1 t 102 f ? f ? f
?
2 t 109 t 0.91 t 120 f
?
3 t 175 f ? f ? f
?
4 t 61 t 0.87 t 70 f
?
```

```
binaryClass
0 P
1 P
2 P
3 P
4 P
```

```
[5 rows x 29 columns]
```

```

from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import accuracy_score, confusion_matrix,
classification_report

numeric_columns = ['age', 'TSH', 'T3', 'TT4', 'T4U', 'FTI', 'TBG']
df[numeric_columns] = df[numeric_columns].apply(pd.to_numeric, errors
= 'coerce')

binary_columns = ['on thyroxine', 'query on thyroxine', 'on
antithyroid medication', 'sick', 'pregnant', 'thyroid surgery',
'I131 treatment', 'query hypothyroid', 'query
hyperthyroid', 'lithium', 'goitre', 'tumor',
'hypopituitary', 'psych', 'TSH measured', 'T3
measured', 'TT4 measured', 'T4U measured',
'FTI measured', 'TBG measured', 'binaryClass']

le = LabelEncoder()
df[binary_columns] = df[binary_columns].apply(le.fit_transform)

df['sex'] = le.fit_transform(df['sex'])

x = df.drop('binaryClass', axis =1)
y = df['binaryClass']

x_train, x_test, y_train, y_test = train_test_split(x, y,
test_size=0.2, random_state=42)

x_train.head()

```

	age	sex	on thyroxine	query on thyroxine	on antithyroid medication
2613	60.0	2	0	0	0
249	25.0	1	0	0	0
2663	19.0	1	0	0	0
3068	71.0	2	1	0	0
1705	52.0	1	0	0	0

	sick	pregnant	thyroid surgery	I131 treatment	query hypothyroid
2613	0	0	0	0	0
249	0	0	0	0	0
2663	0	0	0	0	0

```

1 ...
3068 0 0 0 0
1 ...
1705 0 0 0 0
1 ...

      TSH  T3 measured  T3  TT4 measured  TT4  T4U measured
T4U \
2613 0.005          1  7.1          1  184.0          1
1.03
249  1.600          1  5.4          1  152.0          1
1.50
2663 3.700          1  1.1          1  126.0          1
0.91
3068 25.000         1  0.3          1   31.0          1
0.68
1705 35.000         1  2.1          1   77.0          1
1.10

```

```

      FTI measured  FTI  TBG measured
2613          1  177.0          0
249          1  102.0          0
2663          1  138.0          0
3068          1   46.0          0
1705          1   70.0          0

```

[5 rows x 27 columns]

```
df.head()
```

```

      age  sex  on thyroxine  query on thyroxine  on antithyroid
medication \
0  41.0    1          0          0
0
1  23.0    1          0          0
0
2  46.0    2          0          0
0
3  70.0    1          1          0
0
4  70.0    1          0          0
0

      sick  pregnant  thyroid surgery  I131 treatment  query hypothyroid
... \
0  0          0          0          0          0
...
1  0          0          0          0          0
...
2  0          0          0          0          0

```

```

...
3      0      0      0      0      0
...
4      0      0      0      0      0
...

```

```

      T3  TT4 measured  TT4  T4U measured  T4U  FTI measured
FTI  \
0  2.5      1  125.0      1  1.14      1  109.0
1  2.0      1  102.0      0  NaN      0  NaN
2  NaN      1  109.0      1  0.91      1  120.0
3  1.9      1  175.0      0  NaN      0  NaN
4  1.2      1  61.0      1  0.87      1  70.0

```

```

      TBG measured  TBG  binaryClass
0      0  NaN      1
1      0  NaN      1
2      0  NaN      1
3      0  NaN      1
4      0  NaN      1

```

```
[5 rows x 29 columns]
```

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 3772 entries, 0 to 3771
```

```
Data columns (total 29 columns):
```

```

#      Column                                Non-Null Count  Dtype
---  -
0      age                                3771 non-null      float64
1      sex                                3772 non-null      int32
2      on thyroxine                       3772 non-null      int32
3      query on thyroxine                 3772 non-null      int32
4      on antithyroid medication          3772 non-null      int32
5      sick                               3772 non-null      int32
6      pregnant                           3772 non-null      int32
7      thyroid surgery                    3772 non-null      int32
8      I131 treatment                     3772 non-null      int32
9      query hypothyroid                  3772 non-null      int32
10     query hyperthyroid                 3772 non-null      int32
11     lithium                            3772 non-null      int32
12     goitre                             3772 non-null      int32
13     tumor                              3772 non-null      int32
14     hypopituitary                     3772 non-null      int32

```

15	psych	3772	non-null	int32
16	TSH measured	3772	non-null	int32
17	TSH	3403	non-null	float64
18	T3 measured	3772	non-null	int32
19	T3	3003	non-null	float64
20	TT4 measured	3772	non-null	int32
21	TT4	3541	non-null	float64
22	T4U measured	3772	non-null	int32
23	T4U	3385	non-null	float64
24	FTI measured	3772	non-null	int32
25	FTI	3387	non-null	float64
26	TBG measured	3772	non-null	int32
27	TBG	0	non-null	float64
28	binaryClass	3772	non-null	int32

dtypes: float64(7), int32(22)

memory usage: 530.6 KB

```
df = df.drop('TBG', axis =1)
```

```
df = df.dropna()
```

```
clf = RandomForestClassifier(random_state = 42)
```

```
clf.fit(x_train, y_train)
```

```
RandomForestClassifier(random_state=42)
```

```
y_pred = clf.predict(x_test)
```

```
accuracy = accuracy_score(y_test, y_pred)
```

```
print(f'accuracy_test: {accuracy}')
```

```
accuracy_test: 0.9927404718693285
```

```
report = classification_report(y_test, y_pred)
```

```
print(f'Classification Report:\n{report}')
```

Classification Report:

	precision	recall	f1-score	support
0	0.90	1.00	0.95	37
1	1.00	0.99	1.00	514
accuracy			0.99	551
macro avg	0.95	1.00	0.97	551
weighted avg	0.99	0.99	0.99	551

```
cm = confusion_matrix(y_test, y_pred)
```

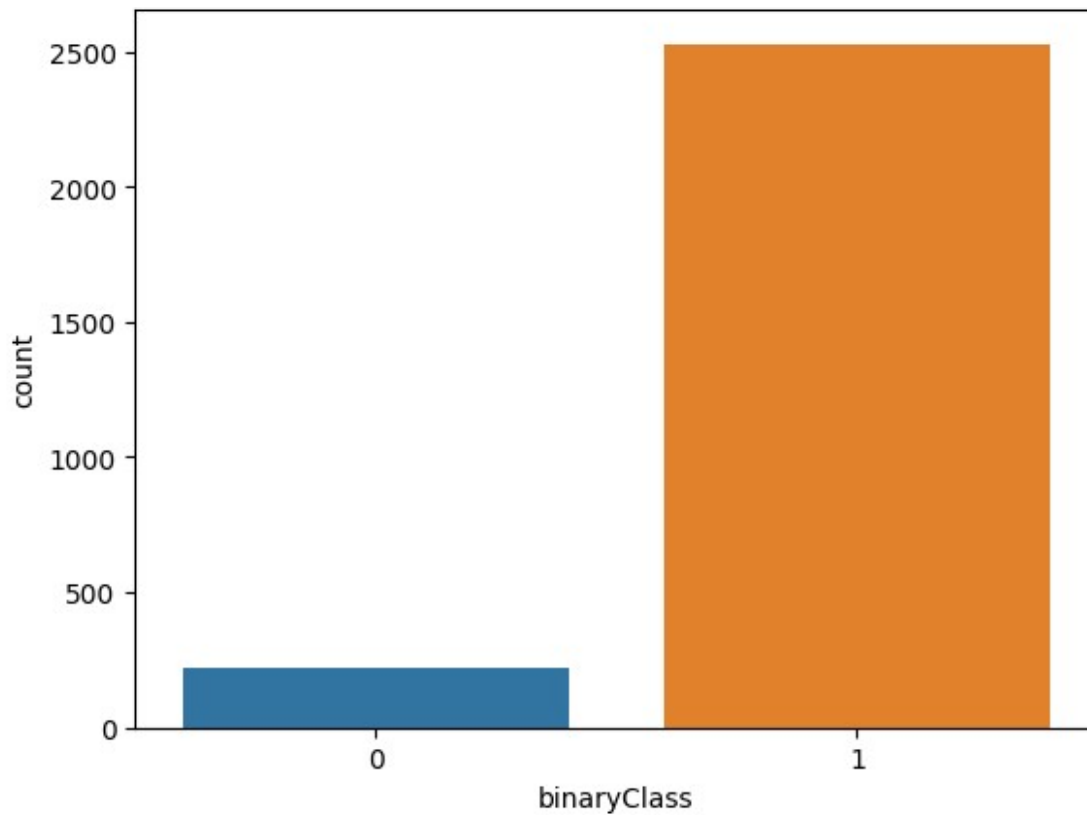
```
print(f'Confusion Matrix:\n{cm}')
```

Confusion Matrix:

```
[[ 37  0]
 [ 4 510]]
```

```
sns.countplot(x= 'binaryClass', data = df)
```

```
<Axes: xlabel='binaryClass', ylabel='count'>
```



```
df['binaryClass'].value_counts()
```

```
binaryClass
```

```
1    2528
```

```
0     223
```

```
Name: count, dtype: int64
```

```
affected_rate = 223/2751
```

```
affected_rate
```

```
0.08106143220647037
```


DATA_VISUALIZATION

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
Index: 2751 entries, 0 to 3771
```

```
Data columns (total 28 columns):
```

#	Column	Non-Null Count	Dtype
0	age	2751 non-null	float64
1	sex	2751 non-null	int32
2	on thyroxine	2751 non-null	int32
3	query on thyroxine	2751 non-null	int32
4	on antithyroid medication	2751 non-null	int32
5	sick	2751 non-null	int32
6	pregnant	2751 non-null	int32
7	thyroid surgery	2751 non-null	int32
8	I131 treatment	2751 non-null	int32
9	query hypothyroid	2751 non-null	int32
10	query hyperthyroid	2751 non-null	int32
11	lithium	2751 non-null	int32
12	goitre	2751 non-null	int32
13	tumor	2751 non-null	int32
14	hypopituitary	2751 non-null	int32
15	psych	2751 non-null	int32
16	TSH measured	2751 non-null	int32
17	TSH	2751 non-null	float64
18	T3 measured	2751 non-null	int32
19	T3	2751 non-null	float64
20	TT4 measured	2751 non-null	int32
21	TT4	2751 non-null	float64
22	T4U measured	2751 non-null	int32
23	T4U	2751 non-null	float64
24	FTI measured	2751 non-null	int32
25	FTI	2751 non-null	float64
26	TBG measured	2751 non-null	int32
27	binaryClass	2751 non-null	int32

```
dtypes: float64(6), int32(22)
```

```
memory usage: 386.9 KB
```

```
df.head(1)
```

	age	sex	on thyroxine	query on thyroxine	on antithyroid medication \
0	41.0	1	0	0	0

	sick	pregnant	thyroid surgery	I131 treatment	query hypothyroid
...	\				
0	0	0	0	0	0

...

	T3 measured	T3	TT4 measured	TT4	T4U measured	T4U	FTI
measured \							
0	1	2.5	1	125.0	1	1.14	
1							

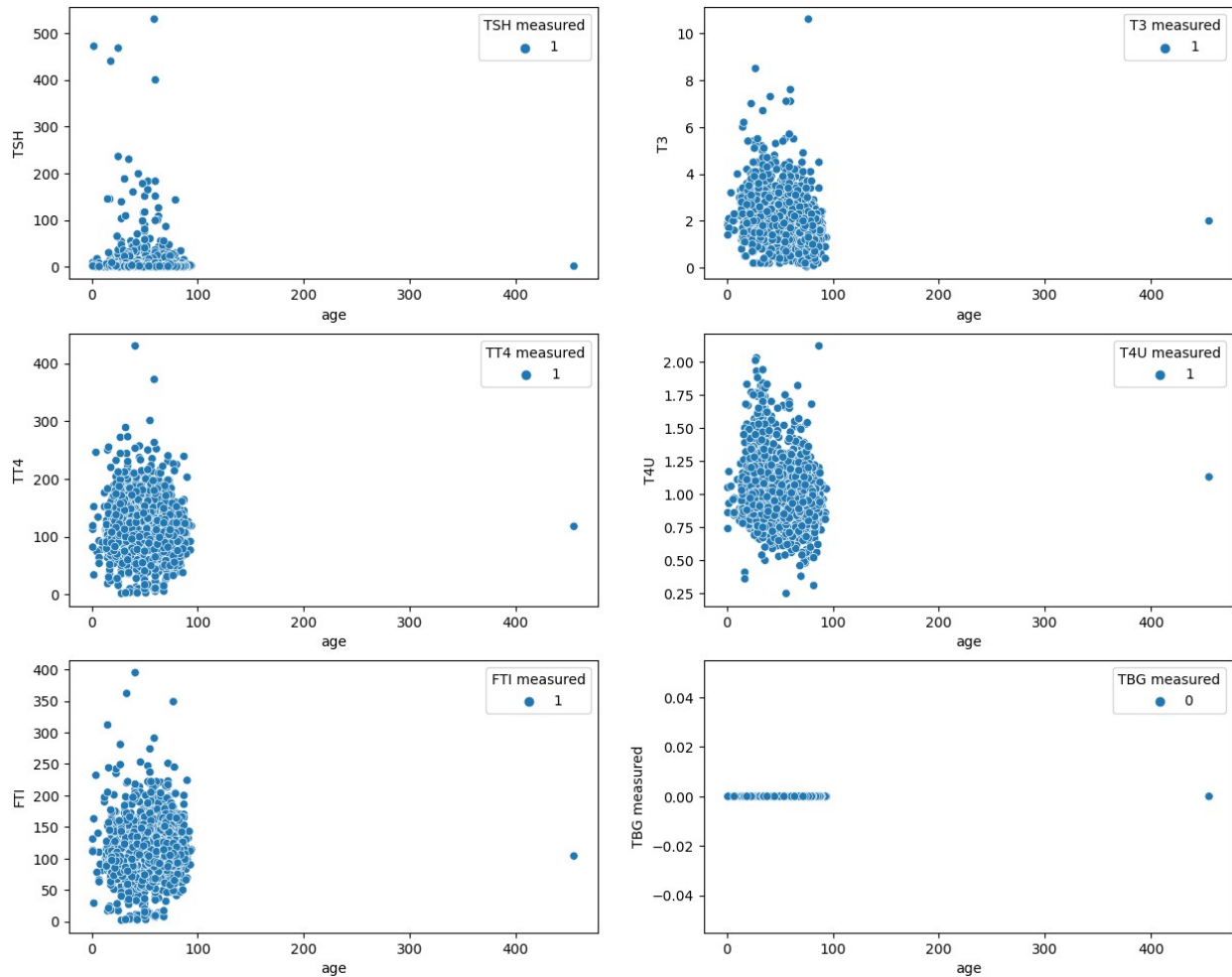
	FTI	TBG measured	binaryClass
0	109.0	0	1

[1 rows x 28 columns]

```
thyroid_data = df[['TSH measured', 'TSH', 'T3 measured', 'T3', 'TT4 measured', 'TT4', 'T4U measured', 'T4U', 'FTI measured', 'FTI', 'TBG measured', 'age']]
```

```
def func_plot(col):  
    fig, axes = plt.subplots(nrows=3, ncols=2, figsize=(15, 12))  
    sns.scatterplot(x=col, y='TSH', hue='TSH measured',  
data=thyroid_data, ax=axes[0, 0])  
    sns.scatterplot(x=col, y='T3', hue='T3 measured',  
data=thyroid_data, ax=axes[0, 1])  
    sns.scatterplot(x=col, y='TT4', hue='TT4 measured',  
data=thyroid_data, ax=axes[1, 0])  
    sns.scatterplot(x=col, y='T4U', hue='T4U measured',  
data=thyroid_data, ax=axes[1, 1])  
    sns.scatterplot(x=col, y='FTI', hue='FTI measured',  
data=thyroid_data, ax=axes[2, 0])  
    sns.scatterplot(x=col, y='TBG measured', hue='TBG measured',  
data=thyroid_data, ax=axes[2, 1])
```

```
func_plot("age")  
plt.show()
```



```
df.head(0)
```

Empty DataFrame

Columns: [age, sex, on thyroxine, query on thyroxine, on antithyroid medication, sick, pregnant, thyroid surgery, I131 treatment, query hypothyroid, query hyperthyroid, lithium, goitre, tumor, hypopituitary, psych, TSH measured, TSH, T3 measured, T3, TT4 measured, TT4, T4U measured, T4U, FTI measured, FTI, TBG measured, binaryClass]

Index: []

[0 rows x 28 columns]