

Thyroid Disease Detection

Problem Statement: Thyroid disease is a common cause of medical diagnosis and prediction, with an onset that is difficult to forecast in medical research. The thyroid gland is one of our body's most vital organs. Thyroid hormone releases are responsible for metabolic regulation. Hyperthyroidism and hypothyroidism are one of the two common diseases of the thyroid that releases thyroid hormones in regulating the rate of body's metabolism. The main goal is to predict the estimated risk on a patient's chance of obtaining thyroid disease or not.

```
In [1]: import pandas as pd
```

```
In [2]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import statsmodels.api as sm
import seaborn as sns

%matplotlib inline
import warnings
warnings.filterwarnings('ignore')
```

For classification problems

Check Null Values

Check Balance Data

```
In [3]: df = pd.read_csv('datafiles/hypothyroid.csv')
df.head()
```

Out[3]:

	age	sex	on thyroxine	query on thyroxine	on antithyroid medication	sick	pregnant	thyroid surgery	I131 treatment	hypoth
0	41	F	f	f	f	f	f	f	f	
1	23	F	f	f	f	f	f	f	f	
2	46	M	f	f	f	f	f	f	f	
3	70	F	t	f	f	f	f	f	f	
4	70	F	f	f	f	f	f	f	f	

5 rows × 30 columns

```
In [4]: df.shape
```

Out[4]: (3772, 30)

In [5]: `df.describe().T`

Out[5]:

	count	unique	top	freq
age	3772	94	59	95
sex	3772	3	F	2480
on thyroxine	3772	2	f	3308
query on thyroxine	3772	2	f	3722
on antithyroid medication	3772	2	f	3729
sick	3772	2	f	3625
pregnant	3772	2	f	3719
thyroid surgery	3772	2	f	3719
I131 treatment	3772	2	f	3713
query hypothyroid	3772	2	f	3538
query hyperthyroid	3772	2	f	3535
lithium	3772	2	f	3754
goitre	3772	2	f	3738
tumor	3772	2	f	3676
hypopituitary	3772	2	f	3771
psych	3772	2	f	3588
TSH measured	3772	2	t	3403
TSH	3772	288	?	369
T3 measured	3772	2	t	3003
T3	3772	70	?	769
TT4 measured	3772	2	t	3541
TT4	3772	242	?	231
T4U measured	3772	2	t	3385
T4U	3772	147	?	387
FTI measured	3772	2	t	3387
FTI	3772	235	?	385
TBG measured	3772	1	f	3772
TBG	3772	1	?	3772
referral source	3772	5	other	2201
binaryClass	3772	2	P	3481

Need to remove ? from the dataIn [6]: `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
```

In [7]: `df["binaryClass"].value_counts()`

```
Out[7]: binaryClass
P      3481
N       291
Name: count, dtype: int64
```

Need to generate some N values to balance the datasetIn [8]: `df["binaryClass"]=df["binaryClass"].map({"P":0,"N":1})`**replace t and f values with 1 and 0 respectively**

```
In [9]: df=df.replace({"t":1,"f":0})
```

replace ? with np.NAN in the dataset

```
In [10]: df=df.replace({"?":np.NAN})
```

```
In [11]: df.head()
```

```
Out[11]:
```

	age	sex	on thyroxine	query on thyroxine	on antithyroid medication	sick	pregnant	thyroid surgery	I131 treatment	hypoth
0	41	F	0	0	0	0	0	0	0	
1	23	F	0	0	0	0	0	0	0	
2	46	M	0	0	0	0	0	0	0	
3	70	F	1	0	0	0	0	0	0	
4	70	F	0	0	0	0	0	0	0	

5 rows × 30 columns

```
In [12]: len(df.columns),df.columns
```

```
Out[12]: (30,
```

```
Index(['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', 'TBG', 'referral source',
       'binaryClass'],
      dtype='object'))
```

```
In [13]: numerical_cols = df.select_dtypes(exclude='object').columns
categorical_cols = df.select_dtypes(include='object').columns
```

```
In [14]: len(numerical_cols), numerical_cols
```

```
Out[14]: (22,
```

```
Index(['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', 'TBG', 'binaryClass'],
      dtype='object'))
```

```
In [15]: len(categorical_cols), categorical_cols
```

```
Out[15]: (8,
```

```
Index(['age', 'sex', 'TSH', 'T3', 'TT4', 'T4U', 'FTI', 'referral source'], dtype=
object'))
```

```
In [16]: df.isnull().sum()
```

```
Out[16]: age                1
sex              150
on thyroxine     0
query on thyroxine 0
on antithyroid medication 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            369
T3 measured     0
T3            769
TT4 measured    0
TT4           231
T4U measured    0
T4U           387
FTI measured    0
FTI           385
TBG measured    0
TBG          3772
referral source 0
binaryClass     0
dtype: int64
```

```
In [17]: def printvalues(col):
          print(f"{col}")
          print(f"{df[col].value_counts()}")
          print(f"Total Null Values: {df[col].isnull().sum()}")
          print("*"*15)
```

```
In [18]: for cat in categorical_cols:
          printvalues(cat)
```

```
age
age
59      95
60      91
70      90
73      81
55      81
..
10       1
4         1
5         1
455       1
6         1
Name: count, Length: 93, dtype: int64
Total Null Values: 1
*****

sex
sex
F      2480
M      1142
Name: count, dtype: int64
Total Null Values: 150
*****

TSH
TSH
0.2      116
1.3      105
1.1       97
1.4       91
1.5       80
...
86         1
18.4        1
89         1
29         1
40         1
Name: count, Length: 287, dtype: int64
Total Null Values: 369
*****

T3
T3
2        238
1.8      207
2.2      201
1.9      189
2.1      184
...
6.7         1
7.3         1
4.6         1
5.2         1
6.6         1
Name: count, Length: 69, dtype: int64
Total Null Values: 769
*****

TT4
```

```
TT4
101    71
93     67
98     63
103    63
102    59
..
289     1
240     1
43      1
34      1
258     1
```

Name: count, Length: 241, dtype: int64

Total Null Values: 231

```
T4U
T4U
0.99    95
0.9     93
1.01    91
1       90
0.92    89
```

```
..
1.84     1
0.57     1
2.03     1
1.97     1
0.25     1
```

Name: count, Length: 146, dtype: int64

Total Null Values: 387

```
FTI
FTI
100     73
93      70
114     65
98      64
107     64
```

```
..
349     1
221     1
187     1
39      1
227     1
```

Name: count, Length: 234, dtype: int64

Total Null Values: 385

```
referral source
referral source
other    2201
SVI      1034
SVHC      386
STMW      112
SVHD       39
```

Name: count, dtype: int64

Total Null Values: 0

```
In [19]: df=df.replace({"F":1,"M":0})
```

```
In [20]: df["referral source"].value_counts()
```

```
Out[20]: referral source
other      2201
SVI        1034
SVHC       386
STMW       112
SVHD        39
Name: count, dtype: int64
```

```
In [21]: del df["referral source"]
```

```
In [22]: df["TBG measured"].value_counts()
```

```
Out[22]: TBG measured
0        3772
Name: count, dtype: int64
```

```
In [23]: df.dtypes
```

```
Out[23]: age                object
sex                float64
on thyroxine        int64
query on thyroxine  int64
on antithyroid medication int64
sick                int64
pregnant            int64
thyroid surgery     int64
I131 treatment      int64
query hypothyroid   int64
query hyperthyroid  int64
lithium             int64
goitre              int64
tumor               int64
hypopituitary       int64
psych               int64
TSH measured        int64
TSH                 object
T3 measured         int64
T3                  object
TT4 measured        int64
TT4                 object
T4U measured        int64
T4U                 object
FTI measured        int64
FTI                 object
TBG measured        int64
TBG                 float64
binaryClass         int64
dtype: object
```

```
In [24]: cols = df.columns[df.dtypes.eq('object')]
df[cols] = df[cols].apply(pd.to_numeric, errors='coerce')
df.dtypes
```

```
Out[24]: age                float64
sex                float64
on thyroxine       int64
query on thyroxine int64
on antithyroid medication int64
sick               int64
pregnant           int64
thyroid surgery    int64
I131 treatment     int64
query hypothyroid  int64
query hyperthyroid int64
lithium            int64
goitre             int64
tumor              int64
hypopituitary      int64
psych              int64
TSH measured       int64
TSH                float64
T3 measured        int64
T3                 float64
TT4 measured       int64
TT4                float64
T4U measured       int64
T4U                float64
FTI measured       int64
FTI                float64
TBG measured       int64
TBG                float64
binaryClass        int64
dtype: object
```

```
In [25]: df.isnull().sum()
```

```
Out[25]: age                1
        sex                150
        on thyroxine        0
        query on thyroxine  0
        on antithyroid medication  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                 369
        T3 measured         0
        T3                  769
        TT4 measured        0
        TT4                 231
        T4U measured        0
        T4U                 387
        FTI measured        0
        FTI                 385
        TBG measured        0
        TBG                 3772
        binaryClass         0
        dtype: int64
```

```
In [26]: df["TBG"].value_counts()
```

```
Out[26]: Series([], Name: count, dtype: int64)
```

```
In [27]: del df["TBG"]
```

```
In [28]: cols
```

```
Out[28]: Index(['age', 'TSH', 'T3', 'TT4', 'T4U', 'FTI'], dtype='object')
```

```
In [29]: from sklearn.impute import SimpleImputer

        imputer = SimpleImputer(strategy='mean')
```

```
In [30]: df['age'] = imputer.fit_transform(df[['age']])
        #
        df['T4U measured'] = imputer.fit_transform(df[['T4U measured']])

        df['TSH'] = imputer.fit_transform(df[['TSH']])
        df['T3'] = imputer.fit_transform(df[['T3']])
        df['TT4'] = imputer.fit_transform(df[['TT4']])
        df['T4U'] = imputer.fit_transform(df[['T4U']])
        df['FTI'] = imputer.fit_transform(df[['FTI']])
```

```
In [31]: df['sex'].fillna(df['sex'].mode()[0], inplace=True)
```

```
In [32]: df.isnull().sum().sum()
```

```
Out[32]: 0
```

```
In [33]: df.to_csv('datafiles/hypothyroid_train.csv', index=False)
```

```
In [34]: df = pd.read_csv('datafiles/hypothyroid_train.csv')
df.head()
```

```
Out[34]:
```

	age	sex	on thyroxine	query on thyroxine	on antithyroid medication	sick	pregnant	thyroid surgery	I131 treatment	hypoti
0	41.0	1.0	0	0	0	0	0	0	0	
1	23.0	1.0	0	0	0	0	0	0	0	
2	46.0	0.0	0	0	0	0	0	0	0	
3	70.0	1.0	1	0	0	0	0	0	0	
4	70.0	1.0	0	0	0	0	0	0	0	

5 rows × 28 columns

```
In [35]: df.isnull().sum().sum()
```

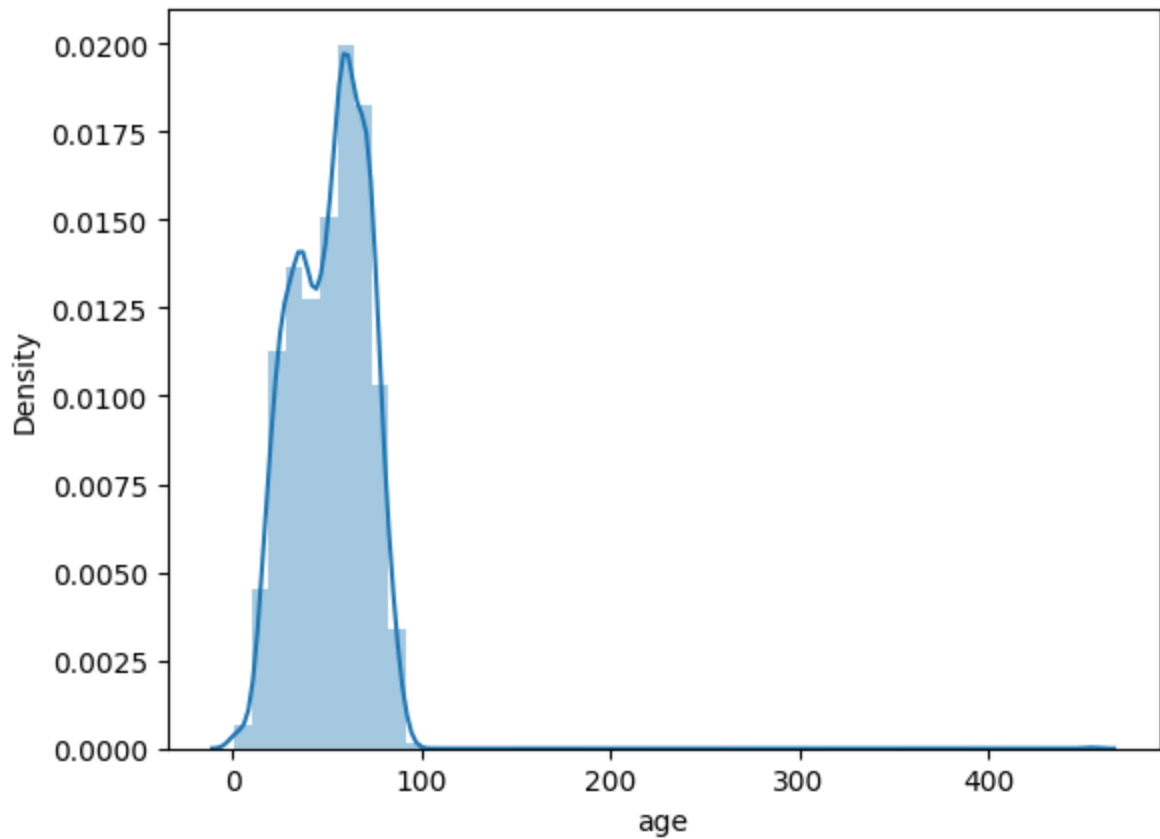
```
Out[35]: 0
```

```
In [36]: df.columns
```

```
Out[36]: Index(['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'],
              dtype='object')
```

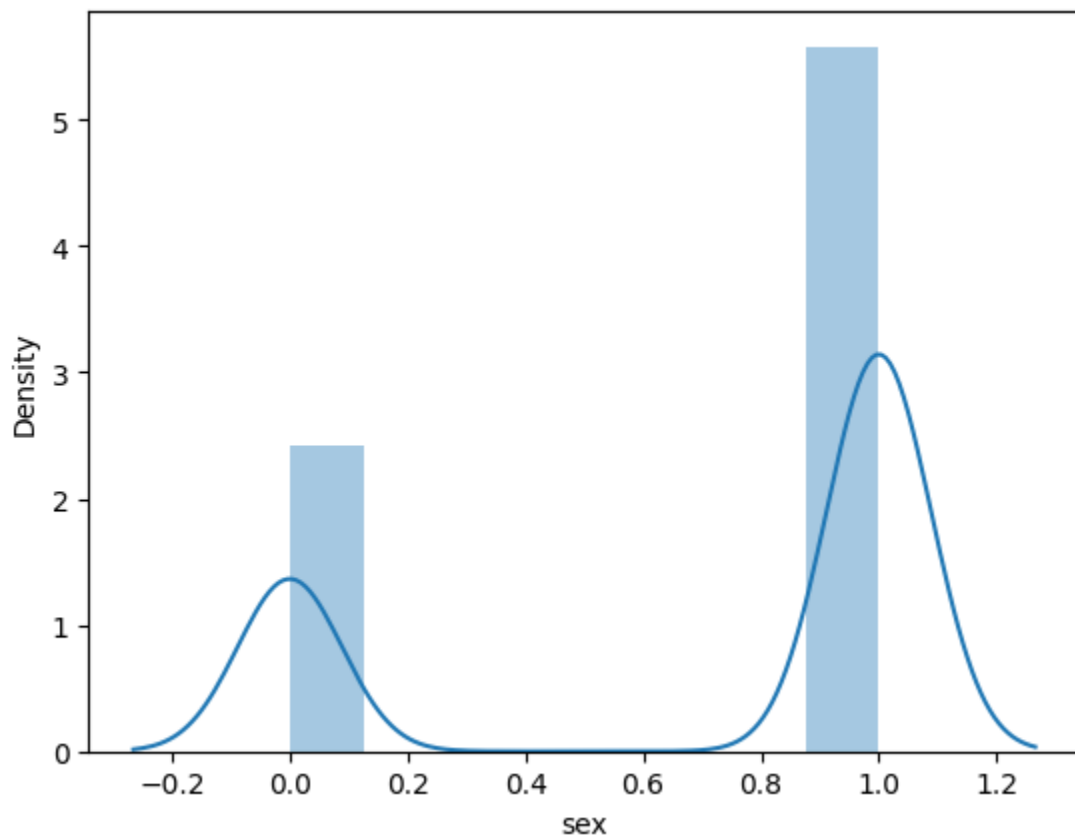
```
In [37]: sns.distplot(df['age'])
```

```
Out[37]: <Axes: xlabel='age', ylabel='Density'>
```



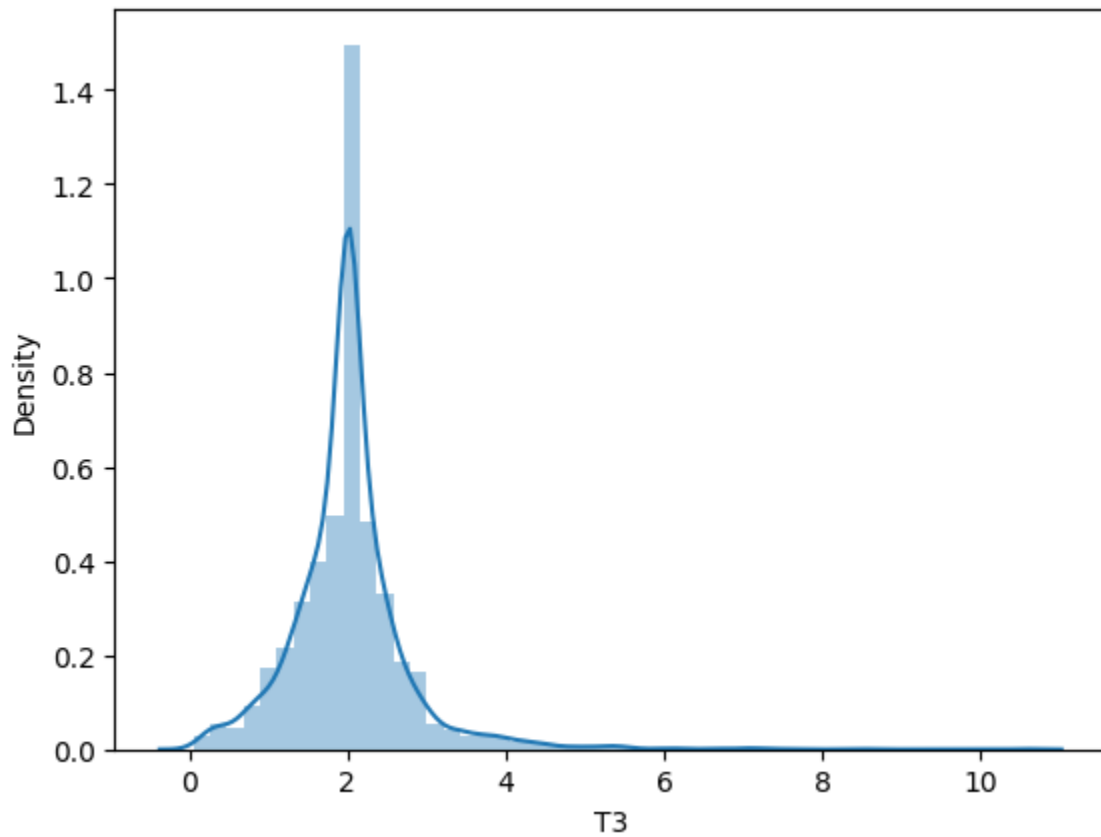
```
In [38]: sns.distplot(df['sex'])
```

```
Out[38]: <Axes: xlabel='sex', ylabel='Density'>
```



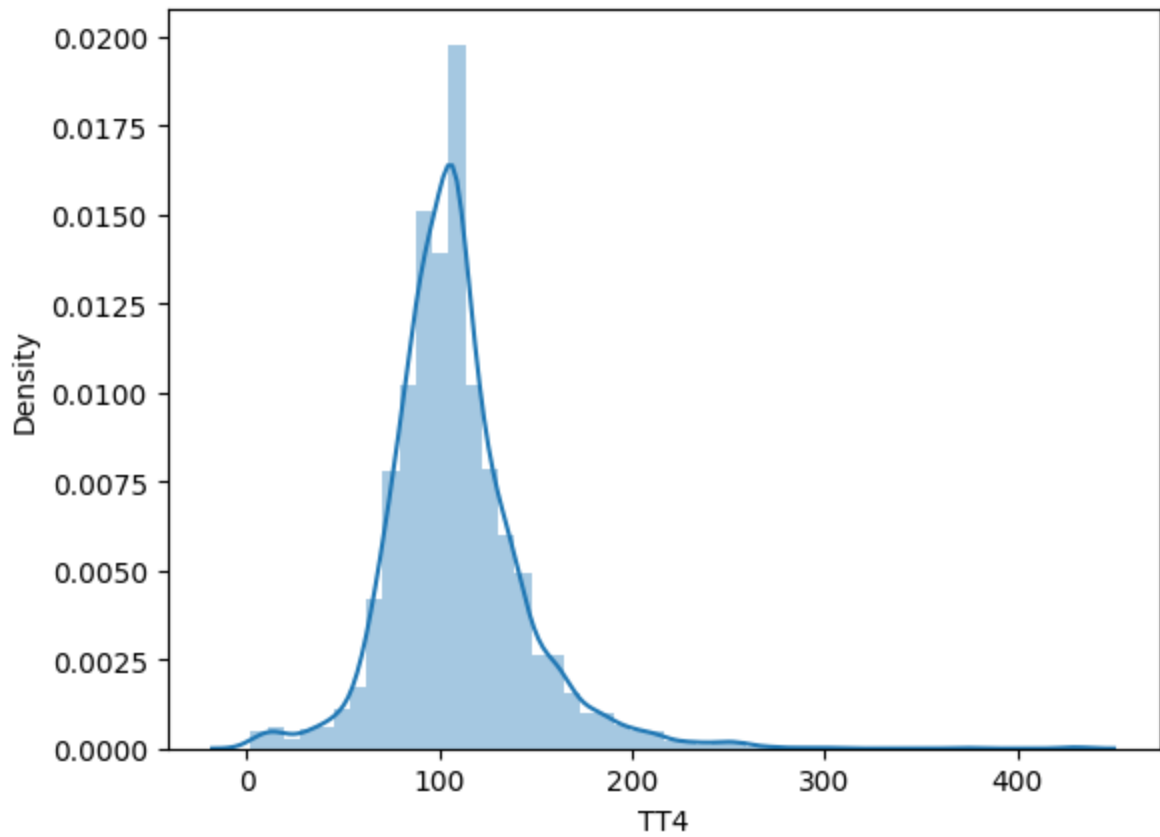
```
In [39]: sns.distplot(df['T3'])
```

```
Out[39]: <Axes: xlabel='T3', ylabel='Density'>
```



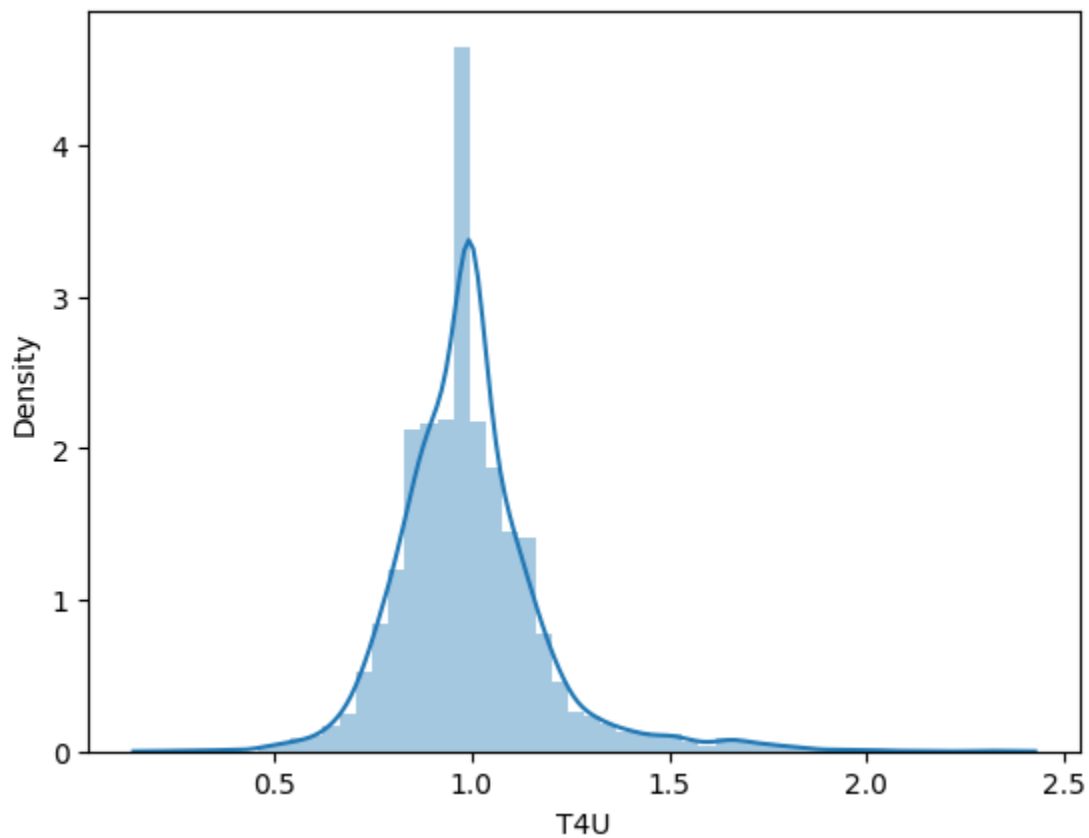
```
In [40]: sns.distplot(df['TT4'])
```

```
Out[40]: <Axes: xlabel='TT4', ylabel='Density'>
```



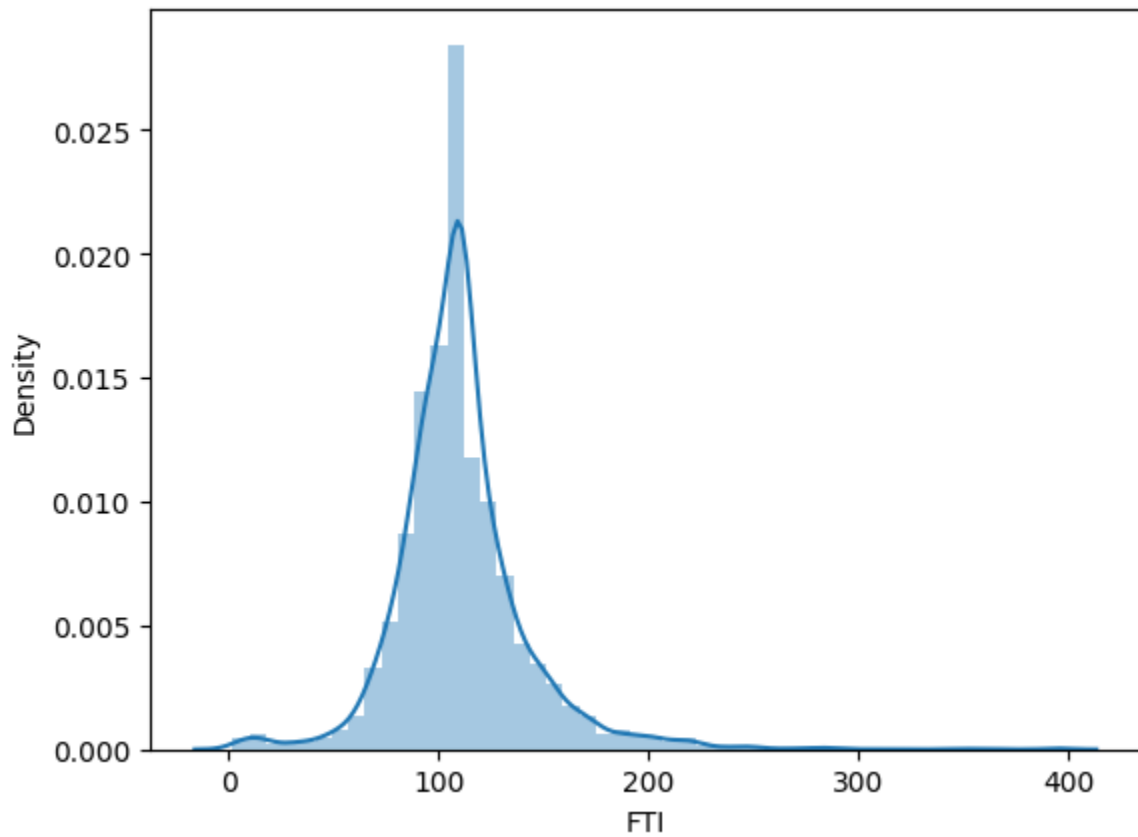
```
In [41]: sns.distplot(df['T4U'])
```

```
Out[41]: <Axes: xlabel='T4U', ylabel='Density'>
```



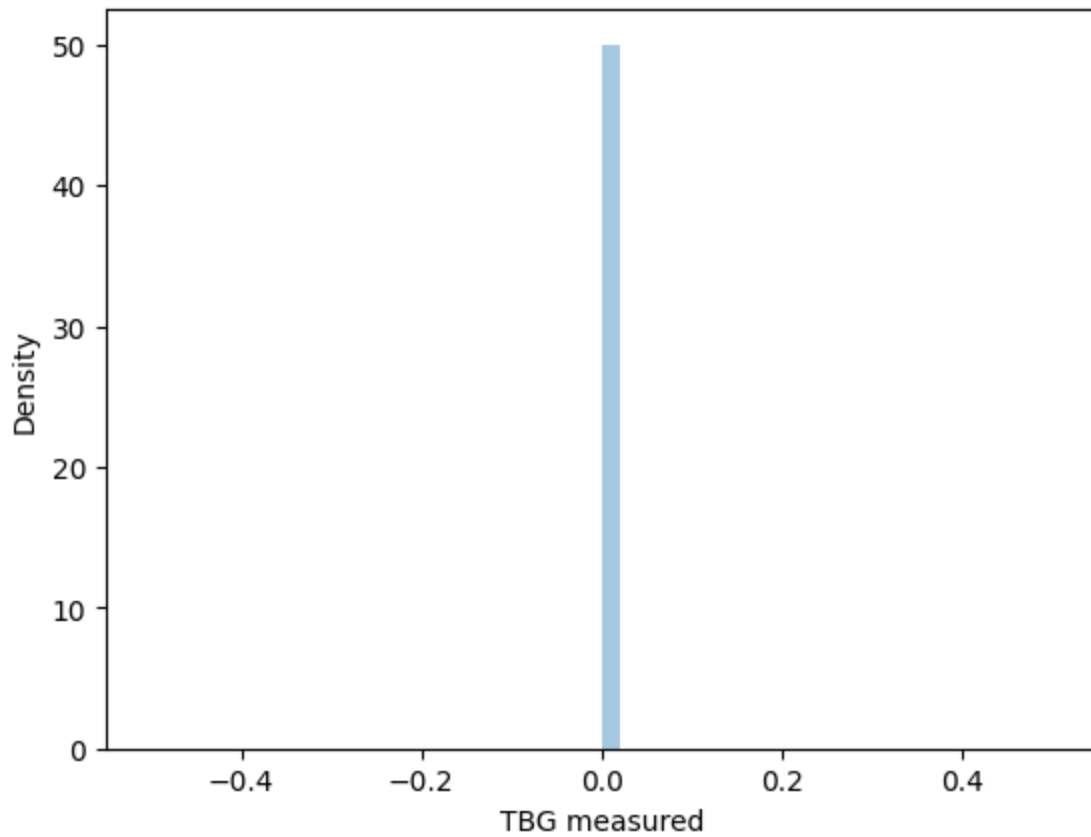
```
In [42]: sns.distplot(df['FTI'])
```

```
Out[42]: <Axes: xlabel='FTI', ylabel='Density'>
```



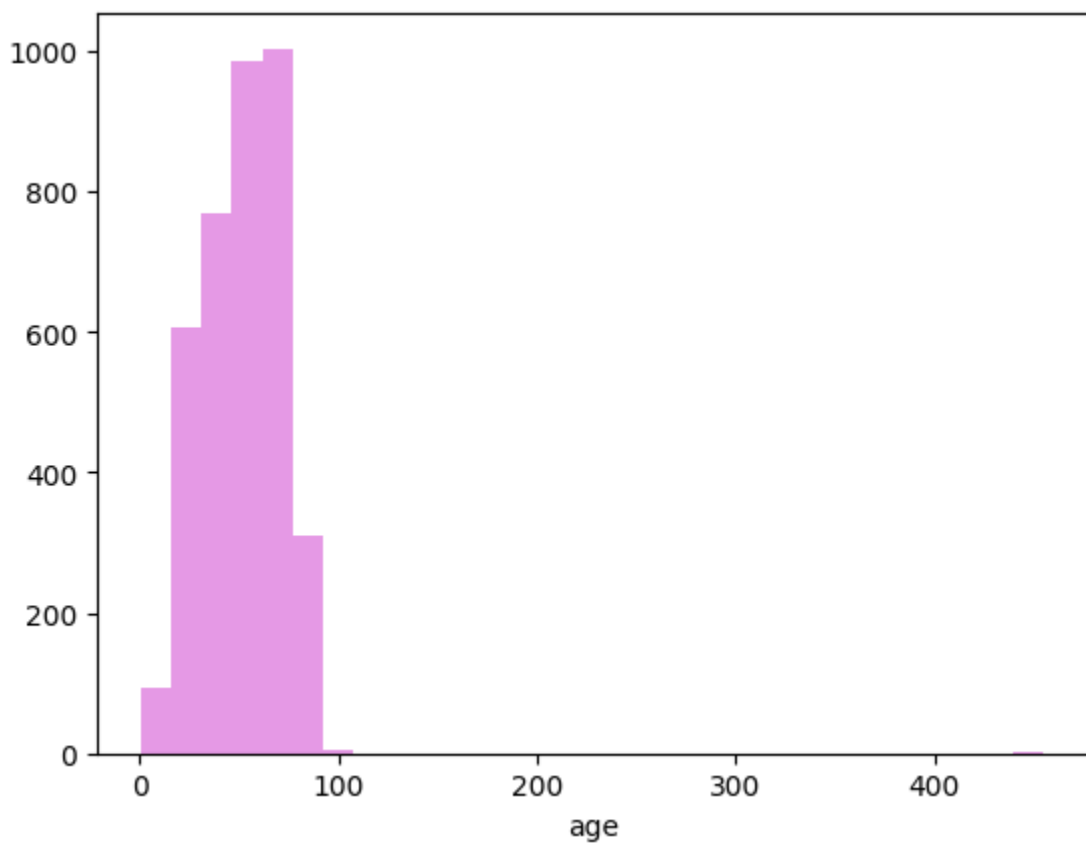
```
In [43]: sns.distplot(df['TBG measured'])
```

```
Out[43]: <Axes: xlabel='TBG measured', ylabel='Density'>
```

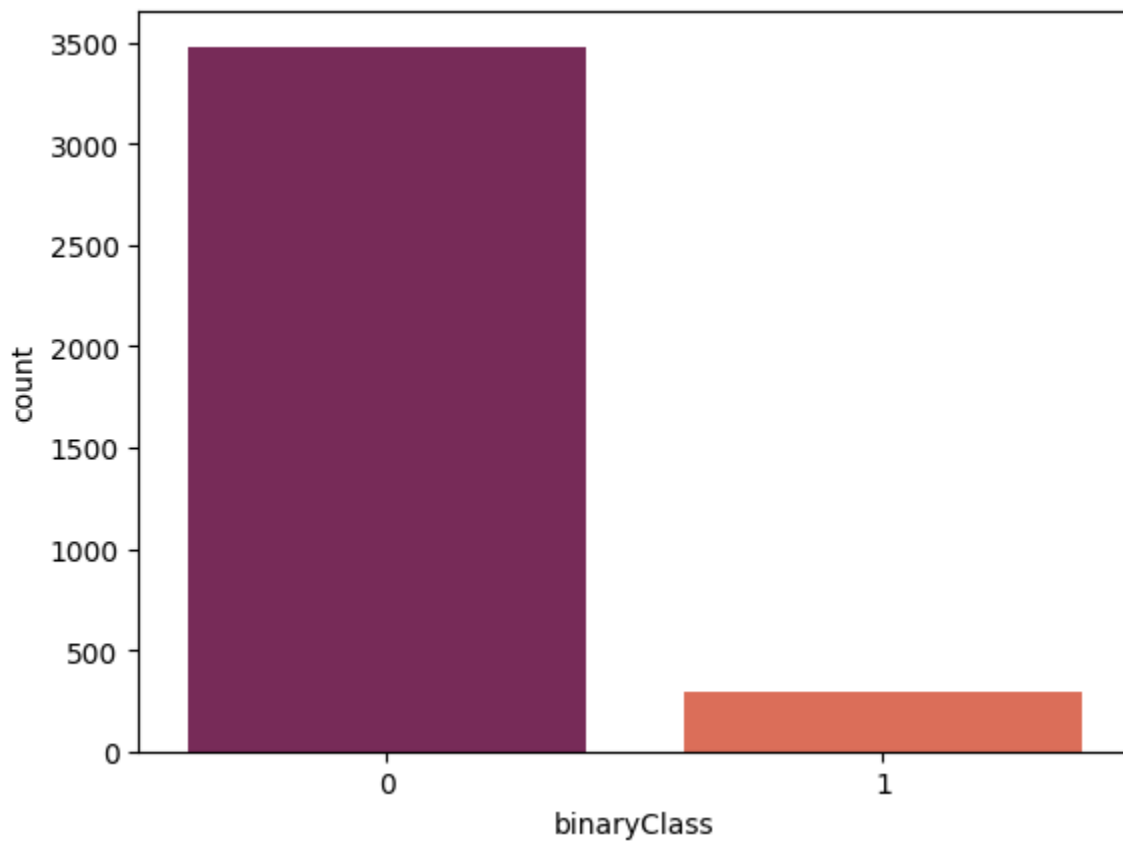
```
In [44]: sns.distplot(df['age'], kde=False, bins=30, color='m')
```

```
Out[44]: <Axes: xlabel='age'>
```



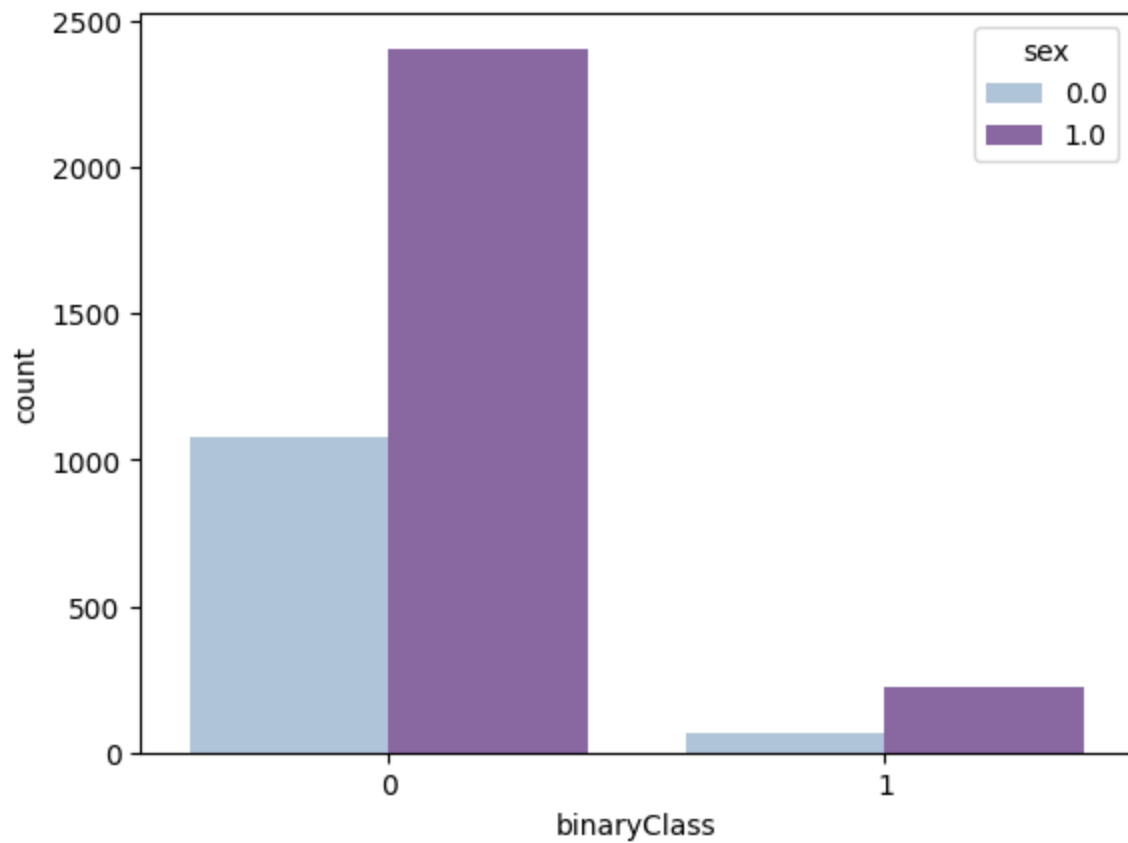
```
In [45]: sns.countplot(x='binaryClass', data=df, palette='rocket')
```

```
Out[45]: <Axes: xlabel='binaryClass', ylabel='count'>
```



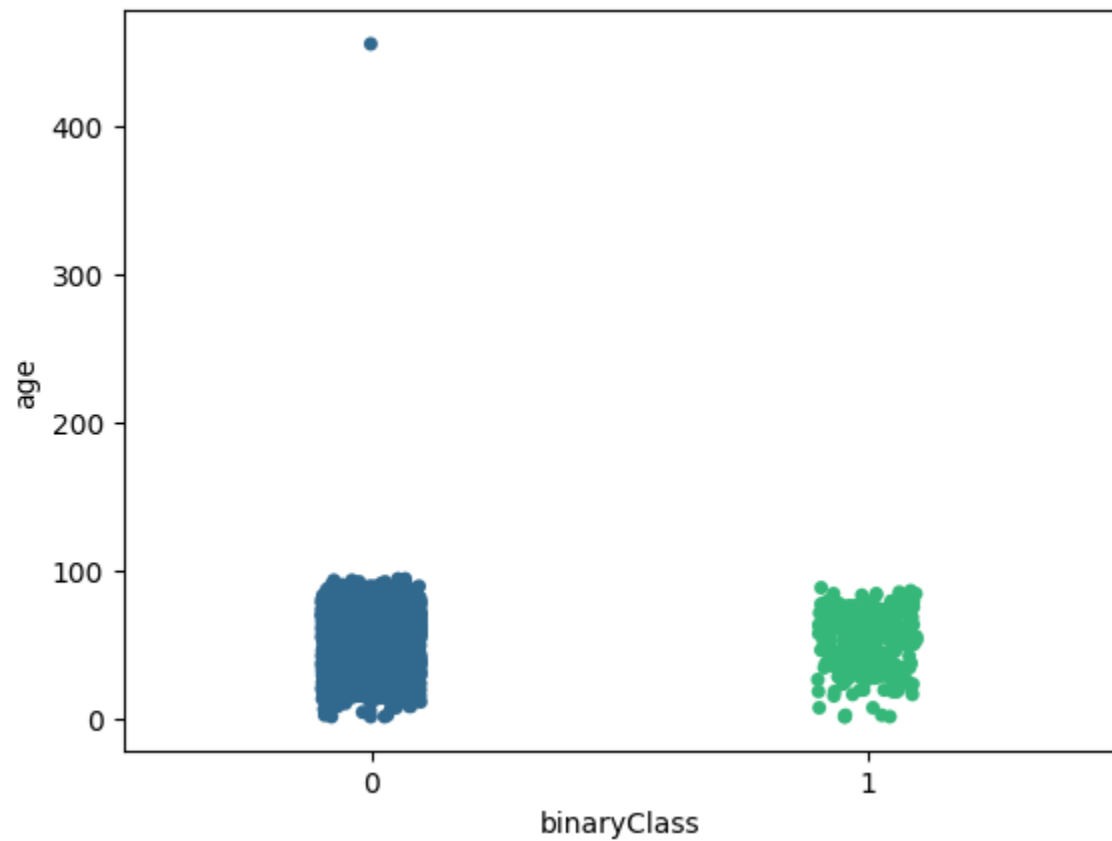
```
In [46]: sns.countplot(x='binaryClass', data=df, hue='sex', palette='BuPu')
```

```
Out[46]: <Axes: xlabel='binaryClass', ylabel='count'>
```



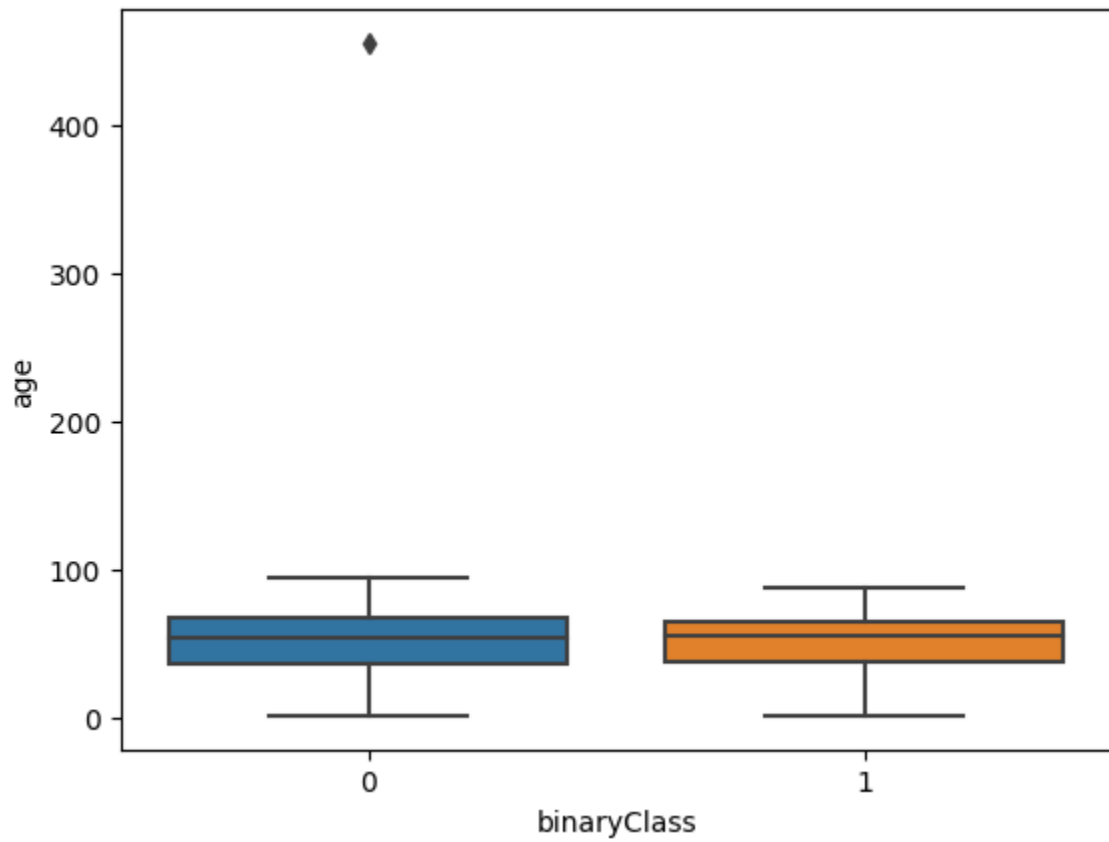
```
In [47]: sns.stripplot(x="binaryClass", y="age", data=df, palette="viridis")
```

```
Out[47]: <Axes: xlabel='binaryClass', ylabel='age'>
```



```
In [48]: sns.boxplot(x='binaryClass', y='age', data=df)
```

```
Out[48]: <Axes: xlabel='binaryClass', ylabel='age'>
```



```
In [49]: df_corr = df.corr()  
df_corr
```

Out[49]:

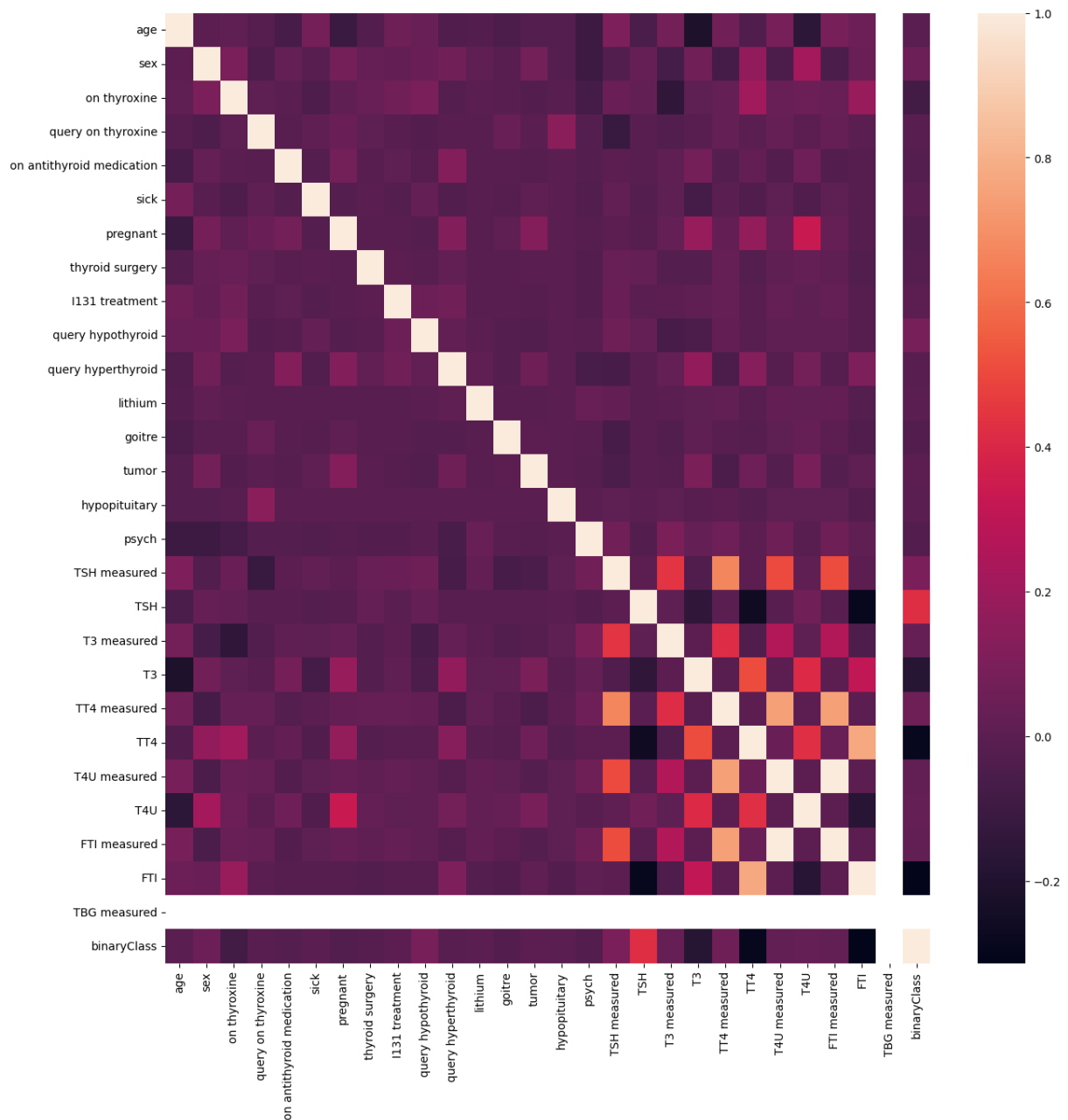
	age	sex	on thyroxine	query on thyroxine	on antithyroid medication	sick	pregnant
age	1.000000	0.000154	0.014563	-0.017870	-0.063881	0.079468	-0.113521
sex	0.000154	1.000000	0.095709	-0.044712	0.027278	-0.007438	0.078665
on thyroxine	0.014563	0.095709	1.000000	0.005995	-0.002201	-0.042053	0.010152
query on thyroxine	-0.017870	-0.044712	0.005995	1.000000	-0.012446	0.012594	0.045247
on antithyroid medication	-0.063881	0.027278	-0.002201	-0.012446	1.000000	-0.021624	0.072050
sick	0.079468	-0.007438	-0.042053	0.012594	-0.021624	1.000000	-0.024040
pregnant	-0.113521	0.078665	0.010152	0.045247	0.072050	-0.024040	1.000000
thyroid surgery	-0.029502	0.034543	0.037583	0.005858	-0.012819	-0.000762	-0.014251
I131 treatment	0.052704	0.022612	0.063373	-0.014610	0.006589	-0.025384	-0.015048
query hypothyroid	0.039562	0.042687	0.094412	-0.029808	-0.017264	0.027718	-0.021364
query hyperthyroid	-0.038054	0.063617	-0.023796	-0.010905	0.126566	-0.035206	0.117605
lithium	-0.030126	0.012138	-0.002509	-0.008026	-0.007436	-0.013944	-0.008266
goitre	-0.051830	-0.010417	-0.010098	0.038000	-0.010241	-0.019205	0.012447
tumor	-0.025037	0.073514	-0.029773	-0.004011	-0.017353	0.010949	0.123728
hypopituitary	-0.024927	-0.024712	-0.006099	0.140500	-0.001749	-0.003279	-0.001944
psych	-0.100116	-0.099896	-0.073571	-0.026247	-0.024318	-0.032883	-0.016577
TSH measured	0.105131	-0.036355	0.041818	-0.117891	0.001736	0.015588	0.001401
TSH	-0.056167	0.033615	0.017138	-0.009453	-0.010668	-0.022099	-0.019693
T3 measured	0.073111	-0.071355	-0.145061	-0.033409	0.010950	0.006695	0.026865
T3	-0.214925	0.064213	0.006485	-0.006466	0.079212	-0.076472	0.181147
TT4 measured	0.067509	-0.076857	0.024964	0.029603	-0.024649	-0.005700	0.021097
TT4	-0.037609	0.167546	0.212801	-0.004702	0.023811	-0.037006	0.172490
T4U measured	0.085361	-0.051662	0.038852	0.031550	-0.029532	0.009399	0.032942

	age	sex	on thyroxine	query on thyroxine	on antithyroid medication	sick	pregnant
T4U	-0.157523	0.221883	0.046368	0.000438	0.060365	-0.039069	0.334702
FTI measured	0.084534	-0.052532	0.038285	0.031420	-0.029788	0.009068	0.032808
FTI	0.050017	0.041900	0.185748	-0.003550	-0.016603	-0.021189	-0.016698
TBG measured	NaN	NaN	NaN	NaN	NaN	NaN	NaN
binaryClass	-0.003174	0.049960	-0.081060	-0.007448	-0.021689	-0.001749	-0.034516

28 rows × 28 columns

```
In [50]: plt.figure(figsize = (15,15))  
sns.heatmap(df_corr)
```

Out[50]: <Axes: >



```
In [51]: x = df.drop('binaryClass', axis=1)
         y = df['binaryClass']
```

```
In [52]: from sklearn.model_selection import train_test_split
         x_train, x_test, y_train, y_test = train_test_split(x, y, test_size=0.25, random_st
```

```
In [53]: from sklearn.preprocessing import StandardScaler

         sc = StandardScaler()
         sc.fit(x_train)
         x_train = sc.transform(x_train)
         x_test = sc.transform(x_test)
```

```
In [54]: x_train[0]
```


	precision	recall	f1-score	support
0	0.97	1.00	0.98	878
1	0.97	0.57	0.72	65
accuracy			0.97	943
macro avg	0.97	0.78	0.85	943
weighted avg	0.97	0.97	0.97	943

```
In [63]: scores = cross_val_score(LogisticRegression(), x_train, y_train, scoring='accuracy',
scores, scores.mean())
```

```
Out[63]: (array([0.96466431, 0.9540636 , 0.96113074, 0.96113074, 0.96466431,
0.96819788, 0.95759717, 0.97173145, 0.95053004, 0.96453901]),
0.9618249254442022)
```

```
In [64]: from sklearn.model_selection import GridSearchCV
```

```
In [65]: parameters = [{'penalty':['l1','l2']],
{'C':[1, 10, 100, 1000]}]
grid_search = GridSearchCV(LogisticRegression(),
param_grid = parameters,
scoring = 'accuracy',
cv = 5,
verbose=0)

best_model = grid_search.fit(x_train, y_train)
```

```
In [66]: best_model.best_params_
```

```
Out[66]: {'C': 100}
```

```
In [67]: model2 = LogisticRegression(C=100)
model2.fit(x_train, y_train)
```

```
Out[67]: ▾ LogisticRegression
LogisticRegression(C=100)
```

```
In [68]: predicted = model.predict(x_test)
```

```
In [69]: metrics.accuracy_score(y_test,predicted)
```

```
Out[69]: 0.9692470837751855
```

```
In [70]: from imblearn.over_sampling import SMOTE
```

```
sm = SMOTE(random_state = 2)
x_train_res, y_train_res = sm.fit_resample(x_train, y_train.ravel())
```

```
In [71]: sm = SMOTE(random_state = 2)
x_test_res, y_test_res = sm.fit_resample(x_test, y_test.ravel())
```

```
In [72]: x_train_res.shape, x_train.shape
```

```
Out[72]: ((5206, 27), (2829, 27))
```

```
In [73]: y_train_res.shape, y_train.shape
```

```
Out[73]: ((5206,), (2829,))
```

```
In [74]: zeros = 0
ones = 0
for item in y_train_res:
    if item == 1:
        ones += 1
    else:
        zeros += 1

print('zeros:', zeros)
print('ones:', ones)
```

```
zeros: 2603
```

```
ones: 2603
```

```
In [75]: lr = LogisticRegression()
model_res = lr.fit(x_train_res, y_train_res)
```

```
In [76]: scores = cross_val_score(lr, x_train_res, y_train_res, scoring='accuracy', cv=10)
scores, scores.mean()
```

```
Out[76]: (array([0.97504798, 0.99232246, 0.98848369, 0.98272553, 0.99424184,
                0.9865643 , 0.98653846, 0.98846154, 0.98076923, 0.98269231]),
         0.9857847335006642)
```

```
In [77]: predicted_res = model.predict(x_test_res)
```

```
In [78]: metrics.accuracy_score(y_test_res, predicted_res)
```

```
Out[78]: 0.8080865603644647
```

```
In [79]: params = [{'learning_rate': [0.01, 0.001],
                    'max_depth': [3, 5, 10],
                    'n_estimators': [10, 50, 100, 200]}]
```

```
In [80]: from sklearn.model_selection import GridSearchCV
from xgboost import XGBClassifier
Xbc = XGBClassifier()
Gcv = GridSearchCV(Xbc, params, scoring='accuracy', cv=5, n_jobs=3, verbose=3)
Gcv.fit(x_train_res, y_train_res)
```

Fitting 5 folds for each of 24 candidates, totalling 120 fits

```
[CV 1/5] END learning_rate=0.01, max_depth=3, n_estimators=10;; score=0.991 total ti
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time= 0.5s
[CV 5/5] END learning_rate=0.001, max_depth=10, n_estimators=200;; score=0.993 total
time= 0.4s

```

Out[80]:

```

  ▸ GridSearchCV
  ▸ estimator: XGBClassifier
    ▸ XGBClassifier

```

In [81]: `Gcv.best_params_`

Out[81]: `{'learning_rate': 0.01, 'max_depth': 5, 'n_estimators': 200}`

In [82]: `XBC = XGBClassifier(learning_rate=0.01,max_depth=5,n_estimators=200)`
`XBC.fit(x_train_res,y_train_res)`

Out[82]:

```

  ▾ XGBClassifier
  XGBClassifier(base_score=None, booster=None, callbacks=None,
                colsample_bylevel=None, colsample_bynode=None,
                colsample_bytree=None, early_stopping_rounds=None,
                enable_categorical=False, eval_metric=None, feature_types=None,
                gamma=None, gpu_id=None, grow_policy=None, importance_type=None,
                interaction_constraints=None, learning_rate=0.01, max_bin=None,

```

In [83]: `XBC.score(x_test_res,y_test_res)`

Out[83]: 0.9988610478359908

```
In [84]: y_pred = XBC.predict(x_test_res)
```

```
In [85]: metrics.accuracy_score(y_test_res,y_pred)
```

Out[85]: 0.9988610478359908

```
In [86]: from sklearn.ensemble import RandomForestRegressor
rfr = RandomForestRegressor()
rfr.fit(x_train,y_train)
```

Out[86]: ▼ RandomForestRegressor
RandomForestRegressor()

```
In [87]: print(f"Train_model_score: {rfr.score(x_train,y_train)}")
print(f"Test_model_score: {rfr.score(x_test,y_test)}")
```

Train_model_score: 0.9928308160427552

Test_model_score: 0.984167117574908

```
In [88]: grid_params = {"n_estimators" : [10,100],
                        "max_depth" : range(2,10,1),
                        "max_features" : ['log2']
                        }
```

```
In [89]: from sklearn.model_selection import GridSearchCV
grid_search = GridSearchCV(estimator=rfr,param_grid=grid_params,cv=5,n_jobs=-1,verb
```

```
In [90]: grid_search.fit(x_train,y_train)
```

Fitting 5 folds for each of 16 candidates, totalling 80 fits

```
In [ ]: grid_search.best_params_
```

Out[]: {'max_depth': 9, 'max_features': 'log2', 'n_estimators': 10}

```
In [ ]: rfr2 = RandomForestRegressor(n_estimators=100,max_depth=10,max_features='log2')
```

```
In [ ]: rfr2.fit(x_train,y_train)
```

Out[]: ▼ RandomForestRegressor
RandomForestRegressor(max_depth=10, max_features='log2')

```
In [ ]: rfr2.score(x_test,y_test)
```

Out[]: 0.8837919470333327

```
In [ ]: y_pred = rfr2.predict(x_test)
```



```
In [ ]: ##### DecisionTreeClassifier
```

```
In [ ]: from sklearn.tree import DecisionTreeClassifier
dtc = DecisionTreeClassifier()
dtc.fit(x_train,y_train)
dtc.score(x_test,y_test)
```

```
Out[ ]: 0.9968186638388123
```

```
In [ ]: y_pred = dtc.predict(x_test)
```

```
In [ ]: Accuracy_score = metrics.accuracy_score(y_test,y_pred)
Accuracy_score
```

```
Out[ ]: 0.9968186638388123
```

```
In [ ]: con_matrix = metrics.confusion_matrix(y_test,y_pred)
con_matrix
```

```
Out[ ]: array([[877,  1],
               [ 2, 63]], dtype=int64)
```

```
In [ ]: grid_param = {
    'criterion': ['gini', 'entropy'],
    'max_depth' : range(2,32,1),
    'splitter' : ['best', 'random']
}
```

```
In [ ]: from sklearn.model_selection import GridSearchCV
grid_search = GridSearchCV(estimator=dtc,
                           param_grid=grid_param,
                           cv=5,
                           n_jobs=-1)
```

```
In [ ]: grid_search.fit(x_train,y_train)
```

```
Out[ ]: ► GridSearchCV
        ► estimator: DecisionTreeClassifier
          ► DecisionTreeClassifier
```

```
In [ ]: best_parameters=grid_search.best_params_
print(best_parameters)

grid_search.best_score_
```

```
{'criterion': 'gini', 'max_depth': 4, 'splitter': 'best'}
```

```
Out[ ]: 0.9964651802745552
```

```
In [ ]: dtc2 = DecisionTreeClassifier(criterion = 'gini', max_depth =4, splitter = 'best')
dtc2.fit(x_train,y_train)
```

```
Out[ ]: ▼      DecisionTreeClassifier
      DecisionTreeClassifier(max_depth=4)
```

```
In [ ]: dtc2.score(x_test,y_test)
```

```
Out[ ]: 0.9989395546129375
```

```
In [ ]:
```

```
In [ ]:
```