

```

import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
sns.set()

from sklearn import preprocessing

from sklearn.linear_model import LogisticRegression
from sklearn.neural_network import MLPClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.svm import SVC
from xgboost import XGBClassifier

from sklearn.feature_selection import f_regression
from sklearn.model_selection import train_test_split, cross_val_score
from sklearn.model_selection import RandomizedSearchCV, GridSearchCV
from sklearn.metrics import confusion_matrix, classification_report
from sklearn.metrics import precision_score, recall_score, f1_score

import warnings
warnings.filterwarnings('ignore')

df = pd.read_csv("drug200.csv")
df.head()

```

	Age	Sex	BP	Cholesterol	Na_to_K	Drug	
0	23	F	HIGH	HIGH	25.355	DrugY	
1	47	M	LOW	HIGH	13.093	drugC	
2	47	M	LOW	HIGH	10.114	drugC	
3	28	F	NORMAL	HIGH	7.798	drugX	
4	61	F	LOW	HIGH	18.043	DrugY	

```

df['Drug'].unique()

array(['DrugY', 'drugC', 'drugX', 'drugA', 'drugB'], dtype=object)

df['Cholesterol'].unique()
df['BP'].unique()
df['Sex'].unique()

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

le1 = preprocessing.LabelEncoder()
df['Drug'] = le1.fit_transform(df['Drug'])
df['Cholesterol'] = le1.fit_transform(df['Cholesterol'])
df['BP'] = le1.fit_transform(df['BP'])
df['Sex'] = le1.fit_transform(df['Sex'])

len(df)
df["Drug"].value_counts().plot(kind="bar");
df["Cholesterol"].value_counts().plot(kind="bar");
df["Sex"].value_counts().plot(kind="bar");
df.info()

```

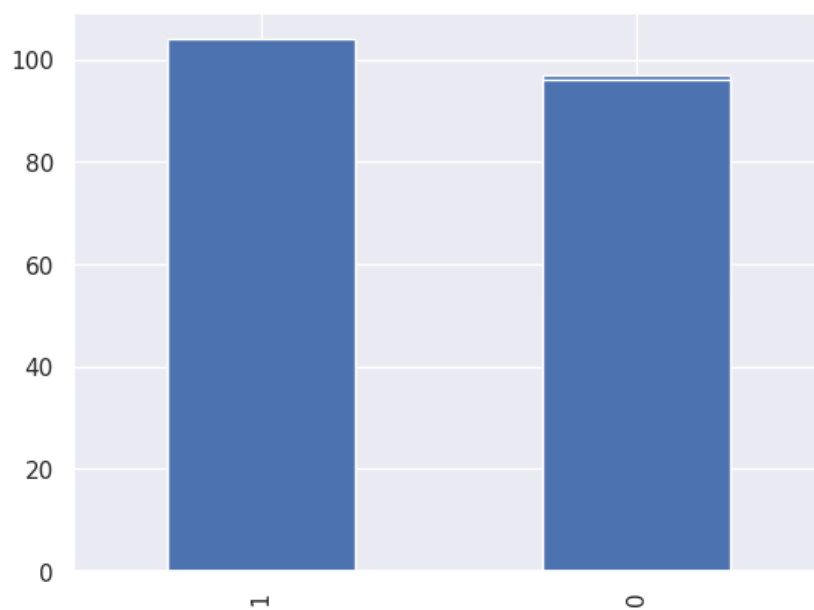
```
df.isna().sum()
df.describe()
pd.crosstab(df["Drug"], df["Sex"])
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 200 entries, 0 to 199
Data columns (total 6 columns):
#   Column          Non-Null Count  Dtype
---  ---
0   Age             200 non-null    int64
1   Sex             200 non-null    int64
2   BP              200 non-null    int64
3   Cholesterol      200 non-null    int64
4   Na_to_K         200 non-null    float64
5   Drug            200 non-null    int64
dtypes: float64(1), int64(5)
memory usage: 9.5 KB
```

```
Sex    0    1
```

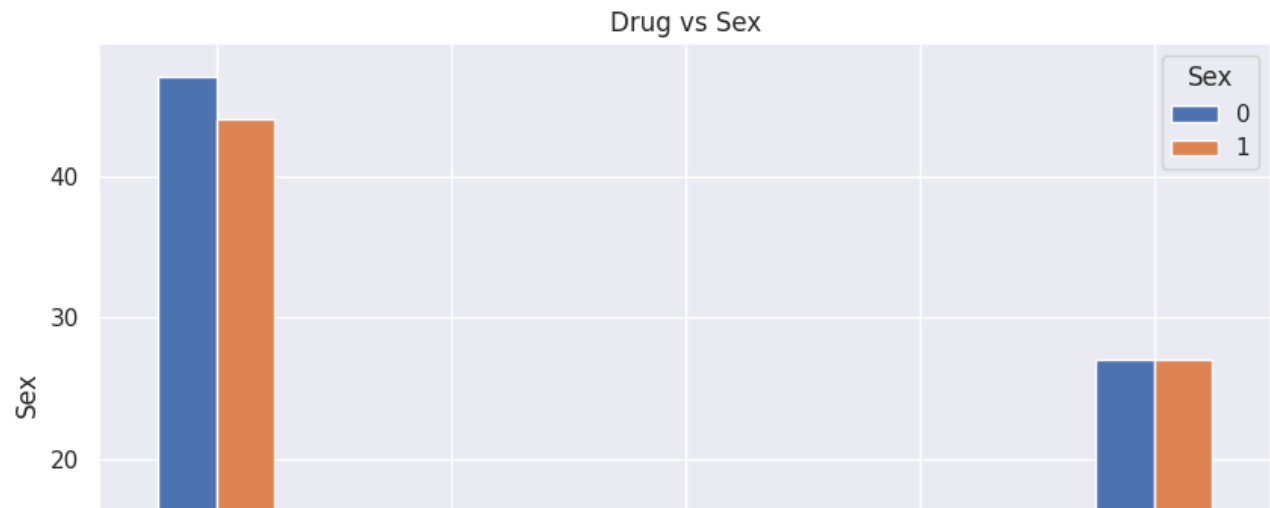
```
Drug
```

```
0    47  44
1     9  14
2     6  10
3     7   9
4    27  27
```

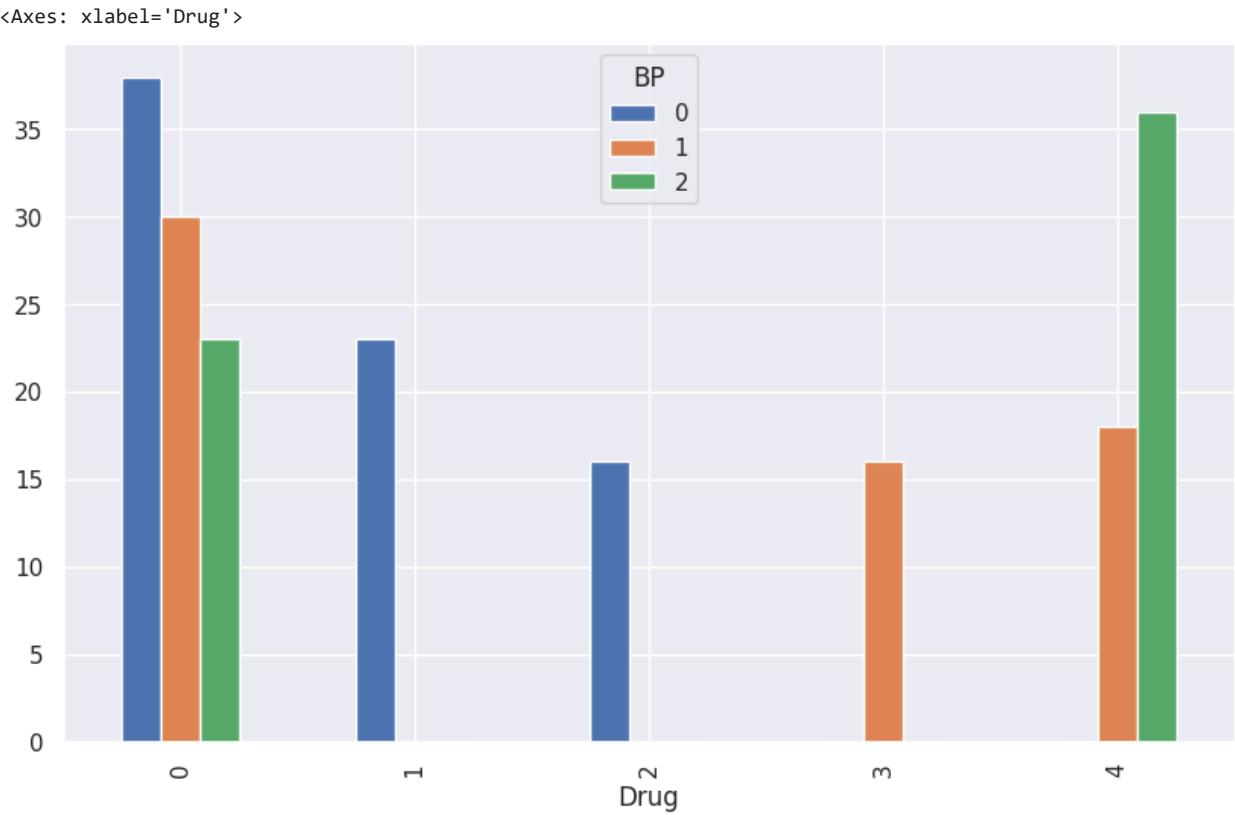


```
pd.crosstab(df["Drug"], df["Sex"]).plot(kind="bar", figsize=(10, 6))
plt.title("Drug vs Sex")
plt.xlabel("Drug Type")
plt.ylabel("Sex")
plt.xticks(rotation=0);
pd.crosstab(df["Drug"], df["BP"])
```


	BP	0	1	2
Drug				
0		38	30	23
1		23	0	0
2		16	0	0
3		0	16	0
4		0	18	36

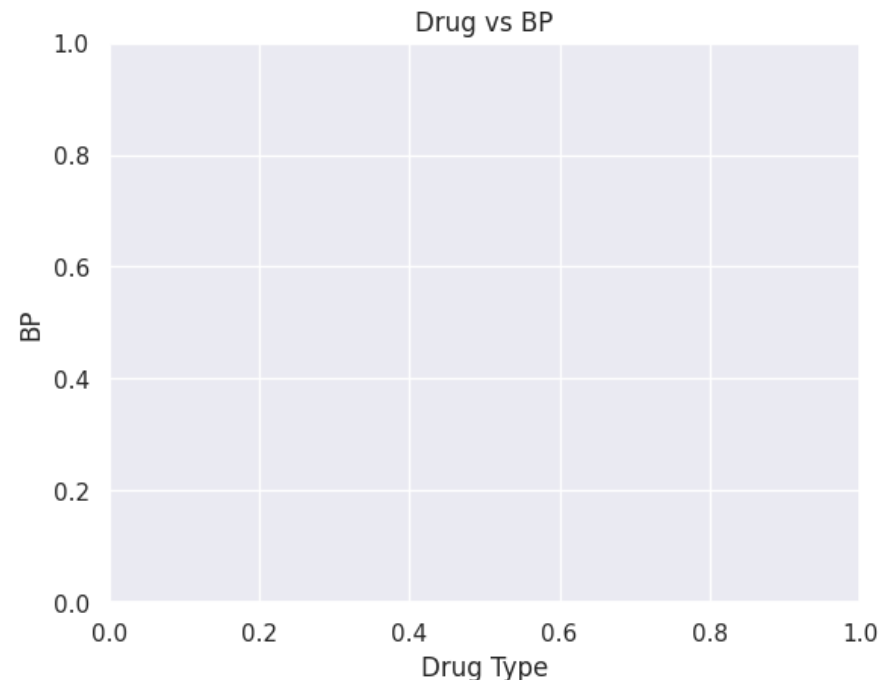


```
pd.crosstab(df["Drug"], df["BP"]).plot(kind="bar",
                                         figsize=(10, 6))
```



```
plt.title("Drug vs BP")
plt.xlabel("Drug Type")
plt.ylabel("BP")
plt.xticks(rotation=0);
pd.crosstab(df["Drug"], df["Cholesterol"])
```

Cholesterol	0	1	
Drug			
0	47	44	
1	12	11	
2	8	8	
3	16	0	
4	20	34	

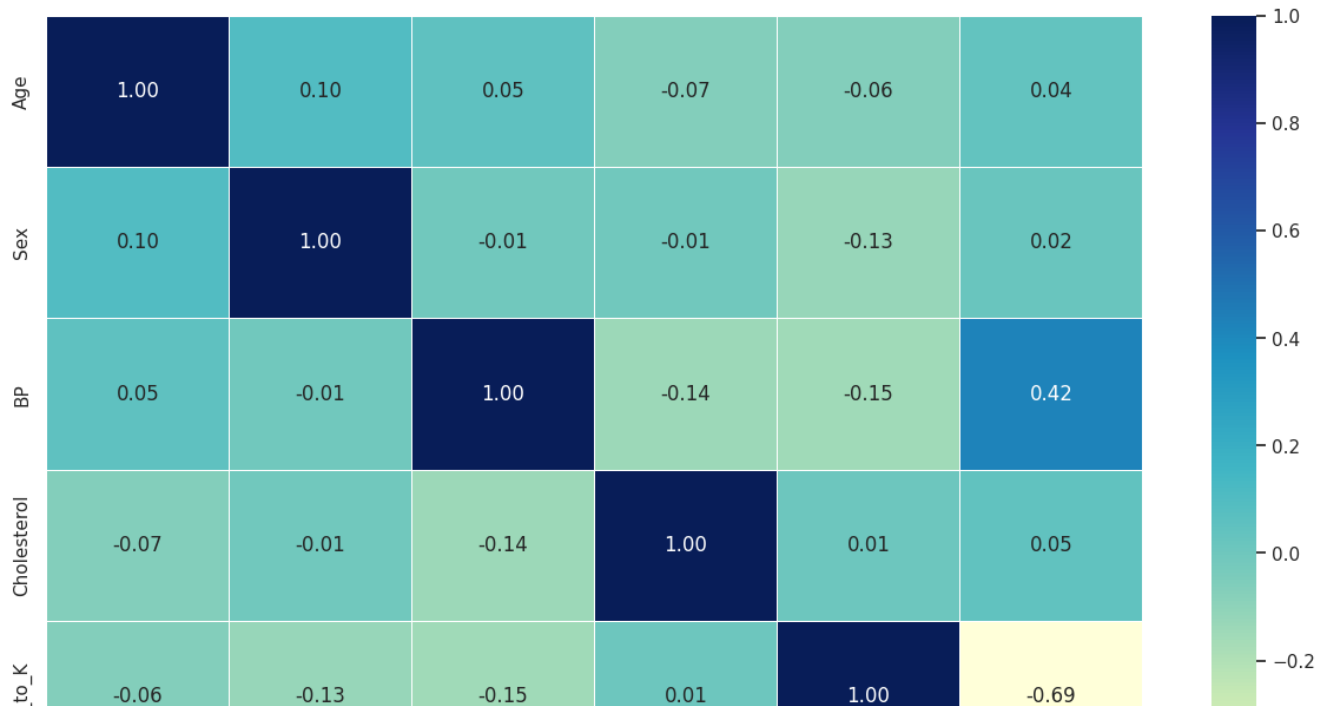


```
pd.crosstab(df["Drug"], df["Cholesterol"]).plot(kind="bar",
                                                    figsize=(10, 6))

plt.title("Drug vs Cholesterol")
plt.xlabel("Drug Type")
plt.ylabel("Cholesterol")
plt.xticks(rotation=0);
```

Drug vs Cholesterol

```
pd.crosstab(df["Sex"], df["Cholesterol"])
pd.crosstab(df["Sex"], df["BP"])
pd.crosstab(df["Cholesterol"], df["BP"])
corr_matrix = df.corr()
fig, ax = plt.subplots(figsize=(15, 10))
ax = sns.heatmap(corr_matrix,
                  annot=True,
                  linewidths=0.5,
                  fmt=".2f",
                  cmap="YlGnBu");
```



```
np.random.seed(42)
```

```
X_train, X_test, y_train, y_test = train_test_split(df.drop("Drug", axis = 1),
                                                    df["Drug"],
                                                    test_size=0.1)
```

```
x = df.drop("Drug", axis = 1)
y = df["Drug"]
```

```
p_values = f_regression(x,y)[1]
len(p_values)
l = []
```

```
col = list(df.columns)
iter_df = 0
```

```
for value in p_values:
    if col[iter_df] == "close":
        iter_df += 1

    l.append({col[iter_df] : value.round(3)})
    iter_df += 1
```

```
l
```

```
[{'Age': 0.556},
 {'Sex': 0.798},
```

```

    {'BP': 0.0},
    {'Cholesterol': 0.496},
    {'Na_to_K': 0.0}]

# Put models in a dictionary
models = {"ANN": MLPClassifier(),
          "Random Forest": RandomForestClassifier(),
          "SVM": SVC(),
          "XGBoost": XGBClassifier()}

# Create a function to fit and score models
def fit_and_score(models, X_train, X_test, y_train, y_test):
    """
    Fits and evaluates given machine learning models.
    models : a dict of different Scikit-Learn machine learning models
    X_train : training data (no labels)
    X_test : testing data (no labels)
    y_train : training labels
    y_test : test labels
    """
    # Set random seed
    np.random.seed(42)
    # Make a dictionary to keep model scores
    model_scores = {}
    # Loop through models
    for name, model in models.items():
        # Fit the model to the data
        model.fit(X_train, y_train)
        # Evaluate the model and append its score to model_scores
        model_scores[name] = model.score(X_test, y_test)
    return model_scores
model_scores = fit_and_score(models=models,
                              X_train=X_train,
                              X_test=X_test,
                              y_train=y_train,
                              y_test=y_test)

model_scores

{'ANN': 0.45, 'Random Forest': 1.0, 'SVM': 0.65, 'XGBoost': 1.0}

model_compare = pd.DataFrame(model_scores, index=["accuracy"])
model_compare.T.plot.bar();

```



```
rf_grid = {"n_estimators": np.arange(10, 1000, 50),
          "max_depth": [None, 3, 5, 10],
          "min_samples_split": np.arange(2, 20, 2),
          "min_samples_leaf": np.arange(1, 20, 2)}
```



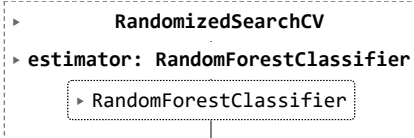
```
# Setup random seed
np.random.seed(42)
```



```
# Setup random hyperparameter search for RandomForestClassifier
rs_rf = RandomizedSearchCV(RandomForestClassifier(),
                           param_distributions=rf_grid,
                           cv=5,
                           n_iter=20,
                           verbose=True)
```

```
# Fit random hyperparameter search model for RandomForestClassifier()
rs_rf.fit(X_train, y_train)
```

Fitting 5 folds for each of 20 candidates, totalling 100 fits



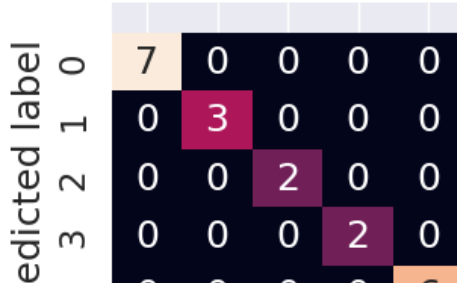
```
rs_rf.best_params_
rs_rf.score(X_test, y_test)
y_preds = rs_rf.predict(X_test)
sns.set(font_scale=1.5)
```

```
def plot_conf_mat(y_test, y_preds):
    """
    Plots a nice looking confusion matrix using Seaborn's heatmap()
    """
    fig, ax = plt.subplots(figsize=(3, 3))
    ax = sns.heatmap(confusion_matrix(y_test, y_preds),
                     annot=True,
                     cbar=False)
    plt.xlabel("True label")
    plt.ylabel("Predicted label")

    bottom, top = ax.get_ylim()
    ax.set_ylim(bottom + 0.5, top - 0.5)
```

```
plot_conf_mat(y_test, y_preds)
print(classification_report(y_test, y_preds))
import pickle
from sklearn.ensemble import RandomForestClassifier
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	7
1	1.00	1.00	1.00	3
2	1.00	1.00	1.00	2
3	1.00	1.00	1.00	2
4	1.00	1.00	1.00	6
accuracy			1.00	20
macro avg	1.00	1.00	1.00	20
weighted avg	1.00	1.00	1.00	20



```
!pip install nbconvert
```

Looking in indexes: <https://pypi.org/simple>, <https://us-python.pkg.dev/colab-wheels/public/simple/>

Requirement already satisfied: nbconvert in /usr/local/lib/python3.9/dist-packages (6.5.4)

Requirement already satisfied: tinycss2 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (1.2.1)

Requirement already satisfied: Jinja2>=3.0 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (3.1.2)

Requirement already satisfied: lxml in /usr/local/lib/python3.9/dist-packages (from nbconvert) (4.9.2)

Requirement already satisfied: nbformat>=5.1 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (5.8.0)

Requirement already satisfied: jupyterlab-pygments in /usr/local/lib/python3.9/dist-packages (from nbconvert) (0.2)

Requirement already satisfied: nbclient>=0.5.0 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (0.7.3)

Requirement already satisfied: BeautifulSoup4 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (4.11.2)

Requirement already satisfied: bleach in /usr/local/lib/python3.9/dist-packages (from nbconvert) (6.0.0)

Requirement already satisfied: MarkupSafe>=2.0 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (2.1.2)

Requirement already satisfied: pandocfilters>=1.4.1 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (1.4.1)

Requirement already satisfied: entrypoints>=0.2.2 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (0.4)

Requirement already satisfied: traitlets>=5.0 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (5.7.1)

Requirement already satisfied: mistune<2,>=0.8.1 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (0.8.4)

Requirement already satisfied: jupyter-core>=4.7 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (5.3.0)

Requirement already satisfied: packaging in /usr/local/lib/python3.9/dist-packages (from nbconvert) (23.1)

Requirement already satisfied: defusedxml in /usr/local/lib/python3.9/dist-packages (from nbconvert) (0.7.1)

Requirement already satisfied: pygments>=2.4.1 in /usr/local/lib/python3.9/dist-packages (from nbconvert) (2.14.0)

Requirement already satisfied: platformdirs>=2.5 in /usr/local/lib/python3.9/dist-packages (from jupyter-core>=4.7)

Requirement already satisfied: jupyter-client>=6.1.12 in /usr/local/lib/python3.9/dist-packages (from nbclient>=0.7.3)

Requirement already satisfied: fastjsonschema in /usr/local/lib/python3.9/dist-packages (from nbformat>=5.1->nbconvert)

Requirement already satisfied: jsonschema>=2.6 in /usr/local/lib/python3.9/dist-packages (from nbformat>=5.1->nbconvert)

Requirement already satisfied: soupsieve>1.2 in /usr/local/lib/python3.9/dist-packages (from BeautifulSoup4->nbconvert)

Requirement already satisfied: webencodings in /usr/local/lib/python3.9/dist-packages (from bleach->nbconvert) (0.5.1)

Requirement already satisfied: six>=1.9.0 in /usr/local/lib/python3.9/dist-packages (from bleach->nbconvert) (1.16.0)

Requirement already satisfied: attrs>=17.4.0 in /usr/local/lib/python3.9/dist-packages (from jsonschema>=2.6->nbconvert)

Requirement already satisfied: pyrsistent!=0.17.0,!0.17.1,!0.17.2,>=0.14.0 in /usr/local/lib/python3.9/dist-packages (from jsonschema>=2.6->nbconvert)

Requirement already satisfied: tornado>=4.1 in /usr/local/lib/python3.9/dist-packages (from jupyter-client>=6.1.12)

Requirement already satisfied: python-dateutil>=2.1 in /usr/local/lib/python3.9/dist-packages (from jupyter-client>=6.1.12)

Requirement already satisfied: pyzmq>=13 in /usr/local/lib/python3.9/dist-packages (from jupyter-client>=6.1.12->nbclient)

```
!jupyter nbconvert --to html Throid.ipynb
```

```
[NbConvertApp] Converting notebook Throid.ipynb to html
[NbConvertApp] Writing 860731 bytes to Throid.html
```


✓ 1s completed at 11:04 AM

