

# wine-prediction-ultimate

June 21, 2024

```
[1]: import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt

df = pd.read_csv('winequalityN.csv')
```

```
[2]: df.head()
```

```
[2]:      type  fixed acidity  volatile acidity  citric acid  residual sugar  \
0  white           7.0           0.27           0.36           20.7
1  white           6.3           0.30           0.34           1.6
2  white           8.1           0.28           0.40           6.9
3  white           7.2           0.23           0.32           8.5
4  white           7.2           0.23           0.32           8.5

      chlorides  free sulfur dioxide  total sulfur dioxide  density  pH  \
0         0.045             45.0             170.0     1.0010  3.00
1         0.049             14.0             132.0     0.9940  3.30
2         0.050             30.0              97.0     0.9951  3.26
3         0.058             47.0             186.0     0.9956  3.19
4         0.058             47.0             186.0     0.9956  3.19

      sulphates  alcohol  quality
0         0.45      8.8        6
1         0.49      9.5        6
2         0.44     10.1        6
3         0.40      9.9        6
4         0.40      9.9        6
```

```
[3]: df.tail()
```

```
[3]:      type  fixed acidity  volatile acidity  citric acid  residual sugar  \
6492  red           6.2           0.600           0.08           2.0
6493  red           5.9           0.550           0.10           2.2
6494  red           6.3           0.510           0.13           2.3
6495  red           5.9           0.645           0.12           2.0
```

6496	red	6.0	0.310	0.47	3.6
------	-----	-----	-------	------	-----

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH \
6492	0.090	32.0	44.0	0.99490	3.45
6493	0.062	39.0	51.0	0.99512	3.52
6494	0.076	29.0	40.0	0.99574	3.42
6495	0.075	32.0	44.0	0.99547	3.57
6496	0.067	18.0	42.0	0.99549	3.39

	sulphates	alcohol	quality
6492	0.58	10.5	5
6493	NaN	11.2	6
6494	0.75	11.0	6
6495	0.71	10.2	5
6496	0.66	11.0	6

[4]: df

[4]:

	type	fixed acidity	volatile acidity	citric acid	residual sugar \
0	white	7.0	0.270	0.36	20.7
1	white	6.3	0.300	0.34	1.6
2	white	8.1	0.280	0.40	6.9
3	white	7.2	0.230	0.32	8.5
4	white	7.2	0.230	0.32	8.5
...	...	...	...	...	...
6492	red	6.2	0.600	0.08	2.0
6493	red	5.9	0.550	0.10	2.2
6494	red	6.3	0.510	0.13	2.3
6495	red	5.9	0.645	0.12	2.0
6496	red	6.0	0.310	0.47	3.6

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH \
0	0.045	45.0	170.0	1.00100	3.00
1	0.049	14.0	132.0	0.99400	3.30
2	0.050	30.0	97.0	0.99510	3.26
3	0.058	47.0	186.0	0.99560	3.19
4	0.058	47.0	186.0	0.99560	3.19
...	...	...	...	...	...
6492	0.090	32.0	44.0	0.99490	3.45
6493	0.062	39.0	51.0	0.99512	3.52
6494	0.076	29.0	40.0	0.99574	3.42
6495	0.075	32.0	44.0	0.99547	3.57
6496	0.067	18.0	42.0	0.99549	3.39

	sulphates	alcohol	quality
0	0.45	8.8	6
1	0.49	9.5	6

2	0.44	10.1	6
3	0.40	9.9	6
4	0.40	9.9	6
...	...	...	...
6492	0.58	10.5	5
6493	NaN	11.2	6
6494	0.75	11.0	6
6495	0.71	10.2	5
6496	0.66	11.0	6

[6497 rows x 13 columns]

```
[5]: df.loc[df['quality'] >= 7]
```

```
[5]:
```

	type	fixed acidity	volatile acidity	citric acid	residual sugar	\
13	white	6.6	0.16	0.40	1.5	
15	white	6.6	0.17	0.38	1.5	
17	white	NaN	0.66	0.48	1.2	
20	white	6.2	0.66	0.48	1.2	
21	white	6.4	0.31	0.38	2.9	
...	...	...	...	...	...	
6439	red	7.4	0.25	0.29	2.2	
6442	red	8.4	0.37	0.43	2.3	
6447	red	7.4	0.36	0.30	1.8	
6453	red	7.0	0.56	0.17	1.7	
6482	red	6.7	0.32	0.44	2.4	

	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	\
13	0.044	48.0	143.0	0.99120	3.54	
15	0.032	28.0	112.0	0.99140	3.25	
17	0.029	29.0	75.0	0.98920	3.33	
20	0.029	29.0	75.0	0.98920	3.33	
21	0.038	19.0	102.0	0.99120	3.17	
...	...	...	...	...	...	
6439	0.054	19.0	49.0	0.99666	3.40	
6442	0.063	12.0	19.0	0.99550	3.17	
6447	0.074	17.0	24.0	0.99419	3.24	
6453	0.065	15.0	24.0	0.99514	3.44	
6482	0.061	24.0	34.0	0.99484	3.29	

	sulphates	alcohol	quality
13	0.52	12.40	7
15	0.55	11.40	7
17	0.39	12.80	8
20	0.39	12.80	8
21	0.35	11.00	7
...	...	...	...

6439	0.76	10.90	7
6442	0.81	11.20	7
6447	0.70	11.40	8
6453	0.68	10.55	7
6482	0.80	11.60	7

[1277 rows x 13 columns]

```
[6]: df.isnull().sum()
```

```
[6]: type                0
fixed acidity          10
volatile acidity       8
citric acid            3
residual sugar         2
chlorides              2
free sulfur dioxide     0
total sulfur dioxide    0
density               0
pH                    9
sulphates              4
alcohol               0
quality               0
dtype: int64
```

```
[7]: df.dropna(inplace=True)
```

```
[8]: df.isnull().sum()
```

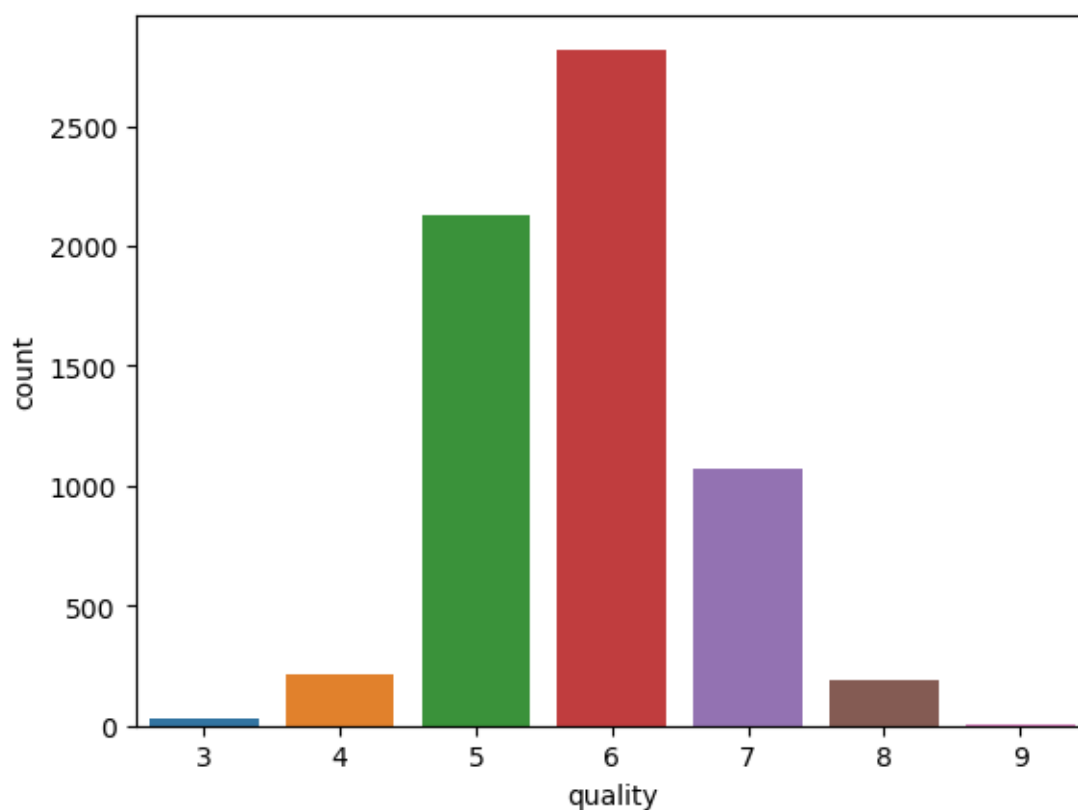
```
[8]: type                0
fixed acidity           0
volatile acidity        0
citric acid             0
residual sugar          0
chlorides               0
free sulfur dioxide     0
total sulfur dioxide    0
density                0
pH                     0
sulphates              0
alcohol                0
quality                0
dtype: int64
```

```
[9]: df['type'].value_counts()
```

```
[9]: type
      white    4870
      red      1593
      Name: count, dtype: int64
```

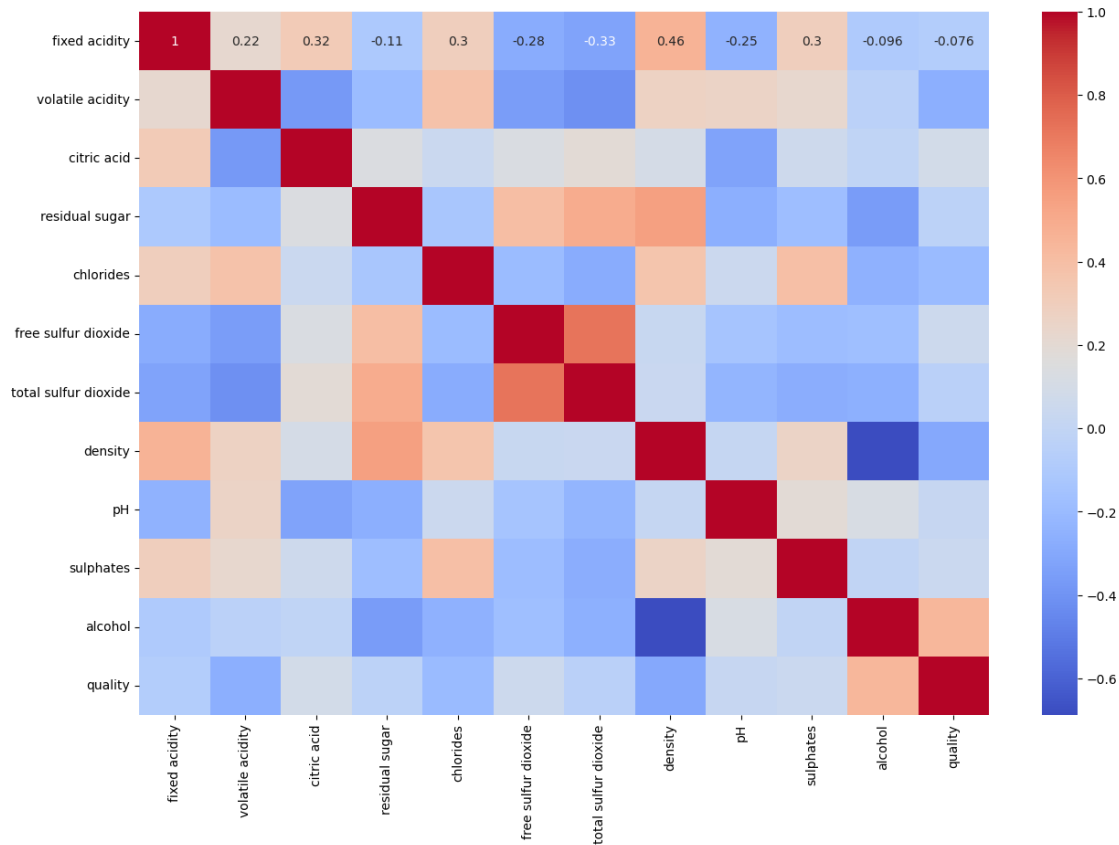
```
[10]: sns.countplot(x='quality', data=df)
```

```
[10]: <Axes: xlabel='quality', ylabel='count'>
```



```
[11]: plt.figure(figsize = (15,10))
      sns.heatmap(df.corr(numeric_only=True), cmap = 'coolwarm', annot = True)
```

```
[11]: <Axes: >
```



## 1 Feature Engineering

```
[12]: # Alcohol is mediumly +ve correlated to quality
# Density is mediumly -ve correlated to quality
# A possible feature could be alcohol/density which also remodensityves heavy
# correlation between alc and den
# Also remove free sulfur dioxide as its very correlated to total sulfur dioxide
# Removing former not latter because latter is more correlated to quality

df_new = df.drop('free sulfur dioxide', axis = 1)
```

```
[13]: df_new['alcohol density'] = (df_new['alcohol']**5)/df_new['density']
```

```
[14]: df_new.head()
```

```
[14]:   type  fixed acidity  volatile acidity  citric acid  residual sugar  \
0  white           7.0           0.27           0.36           20.7
1  white           6.3           0.30           0.34            1.6
2  white           8.1           0.28           0.40            6.9
```

3	white	7.2	0.23	0.32	8.5
4	white	7.2	0.23	0.32	8.5

	chlorides	total sulfur dioxide	density	pH	sulphates	alcohol \
0	0.045	170.0	1.0010	3.00	0.45	8.8
1	0.049	132.0	0.9940	3.30	0.49	9.5
2	0.050	97.0	0.9951	3.26	0.44	10.1
3	0.058	186.0	0.9956	3.19	0.40	9.9
4	0.058	186.0	0.9956	3.19	0.40	9.9

	quality	alcohol density
0	6	52720.471209
1	6	77845.164738
2	6	105618.535836
3	6	95519.289865
4	6	95519.289865

```
[15]: df_ml = pd.get_dummies(df_new, drop_first=True)
```

```
[16]: df_ml.head()
```

```
[16]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides \
0	7.0	0.27	0.36	20.7	0.045
1	6.3	0.30	0.34	1.6	0.049
2	8.1	0.28	0.40	6.9	0.050
3	7.2	0.23	0.32	8.5	0.058
4	7.2	0.23	0.32	8.5	0.058

	total sulfur dioxide	density	pH	sulphates	alcohol	quality \
0	170.0	1.0010	3.00	0.45	8.8	6
1	132.0	0.9940	3.30	0.49	9.5	6
2	97.0	0.9951	3.26	0.44	10.1	6
3	186.0	0.9956	3.19	0.40	9.9	6
4	186.0	0.9956	3.19	0.40	9.9	6

	alcohol density	type_white
0	52720.471209	True
1	77845.164738	True
2	105618.535836	True
3	95519.289865	True
4	95519.289865	True

```
[17]: df_ml = df_ml.drop(['density', 'alcohol'], axis = 1)
```

```
[18]: df_ml.head()
```

```
[18]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides \
0	7.0	0.27	0.36	20.7	0.045
1	6.3	0.30	0.34	1.6	0.049
2	8.1	0.28	0.40	6.9	0.050
3	7.2	0.23	0.32	8.5	0.058
4	7.2	0.23	0.32	8.5	0.058

	total sulfur dioxide	pH	sulphates	quality	alcohol density	type_white
0	170.0	3.00	0.45	6	52720.471209	True
1	132.0	3.30	0.49	6	77845.164738	True
2	97.0	3.26	0.44	6	105618.535836	True
3	186.0	3.19	0.40	6	95519.289865	True
4	186.0	3.19	0.40	6	95519.289865	True

```
[19]: df_ml.isnull().sum()
```

```
[19]: fixed acidity      0
volatile acidity      0
citric acid           0
residual sugar        0
chlorides              0
total sulfur dioxide  0
pH                    0
sulphates             0
quality               0
alcohol density       0
type_white            0
dtype: int64
```

```
[20]: Y = df_ml['quality'].apply(lambda y: 1 if y>=6 else 0)
Y
```

```
[20]: 0      1
1      1
2      1
3      1
4      1
..
6491   1
6492   0
6494   1
6495   0
6496   1
Name: quality, Length: 6463, dtype: int64
```

```
[21]: X = df_ml.drop('quality', axis = 1)
X.head()
```



```
[21]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides \
0	7.0	0.27	0.36	20.7	0.045
1	6.3	0.30	0.34	1.6	0.049
2	8.1	0.28	0.40	6.9	0.050
3	7.2	0.23	0.32	8.5	0.058
4	7.2	0.23	0.32	8.5	0.058

	total sulfur dioxide	pH	sulphates	alcohol density	type_white
0	170.0	3.00	0.45	52720.471209	True
1	132.0	3.30	0.49	77845.164738	True
2	97.0	3.26	0.44	105618.535836	True
3	186.0	3.19	0.40	95519.289865	True
4	186.0	3.19	0.40	95519.289865	True

```
[22]: # Standardize feature values so that high valued feautres don't influence ↵
      ↪ others
```

```
from sklearn.preprocessing import StandardScaler

scaler = StandardScaler()
scaler.fit(X)
X_standard = scaler.transform(X)

X_standard #standardised numpy array of features
```

```
[22]: array([[ -0.16778609, -0.42270958,  0.2839587 , ..., -0.5449872 ,
        -1.03523983,  0.5719307 ],
        [-0.70715516, -0.2404789 ,  0.14625658, ..., -0.27635393,
        -0.75614416,  0.5719307 ],
        [ 0.67979387, -0.36196602,  0.55936296, ..., -0.61214551,
        -0.44762587,  0.5719307 ],
        ...,
        [-0.70715516,  1.03513588, -1.29961576, ...,  1.46976231,
         0.1757951 , -1.74846359],
        [-1.01536606,  1.85517396, -1.36846682, ...,  1.20112905,
        -0.38884046, -1.74846359],
        [-0.93831333, -0.17973534,  1.0413204 , ...,  0.86533746,
         0.1762463 , -1.74846359]])
```

```
[23]: X = X_standard
```

```
[24]: from sklearn.model_selection import train_test_split
```

```
[25]: X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.1, ↵
      ↪ random_state = 25)
```

```
[26]: from sklearn.ensemble import RandomForestClassifier
```

```
rfc = RandomForestClassifier()  
rfc.fit(X_train, Y_train)  
rfc_pred = rfc.predict(X_test)
```

```
[27]: from sklearn.metrics import accuracy_score, classification_report,  
      ↪confusion_matrix
```

```
[28]: accuracy_score(Y_test,rfc_pred)
```

```
[28]: 0.8408037094281299
```

```
[29]: print(classification_report(Y_test, rfc_pred))
```

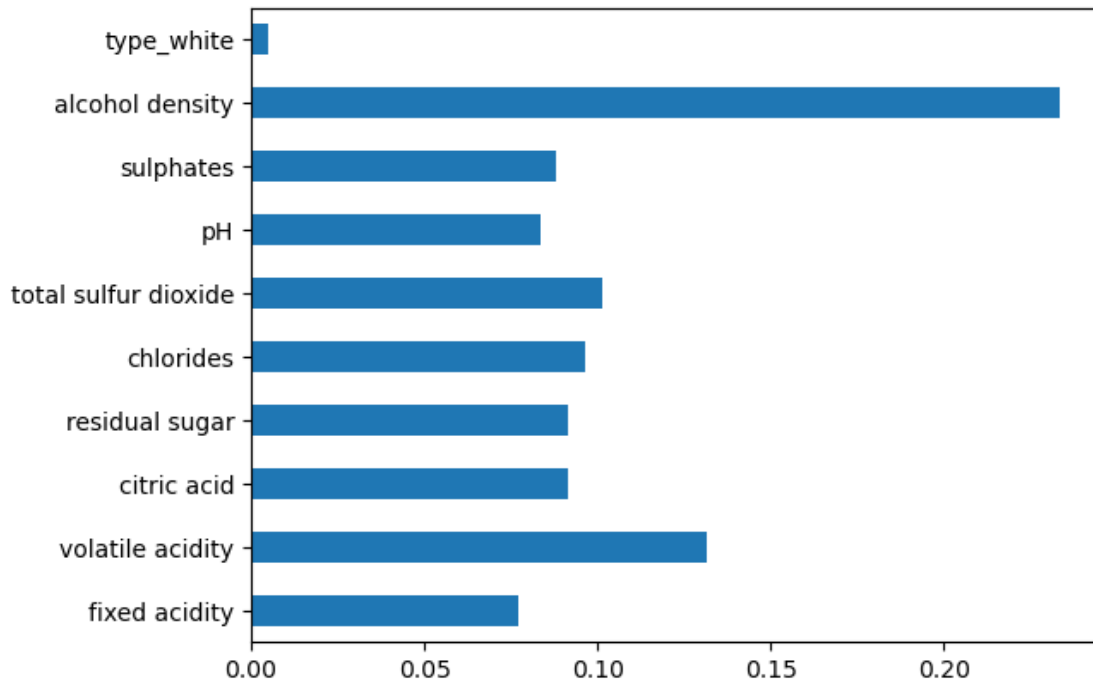
	precision	recall	f1-score	support
0	0.82	0.74	0.78	247
1	0.85	0.90	0.88	400
accuracy			0.84	647
macro avg	0.84	0.82	0.83	647
weighted avg	0.84	0.84	0.84	647

```
[30]: print(confusion_matrix(Y_test, rfc_pred))
```

```
[[183  64]  
 [ 39 361]]
```

```
[526]: pd.Series(rfc.feature_importances_, index=df_ml.drop('quality', axis = 1).  
      ↪columns).plot(kind = 'barh')
```

```
[526]: <Axes: >
```



```
[527]: from sklearn.ensemble import GradientBoostingClassifier
```

```
[528]: gbc = GradientBoostingClassifier().fit(X_train, Y_train)
```

```
[529]: gbc_pred = gbc.predict(X_test)
```

```
[530]: accuracy_score(Y_test, gbc_pred)
```

```
[530]: 0.7511591962905718
```

```
[531]: print(classification_report(Y_test, gbc_pred))
```

	precision	recall	f1-score	support
0	0.70	0.58	0.63	242
1	0.77	0.85	0.81	405
accuracy			0.75	647
macro avg	0.74	0.72	0.72	647
weighted avg	0.75	0.75	0.75	647

```
[542]: i_fixed_acidity = float(input("Enter fixed acidity [3.8-15.9]: "))
       i_volatile_acidity = float(input("Enter volatile acidity [0.08-1.58]: "))
```

```

i_citric_acid = float(input("Enter citric acid [0.0-1.66]: "))
i_residual_sugar = float(input("Enter residual sugar [0.6-65.8]: "))
i_chlorides = float(input("Enter chlorides [0.009-0.611]: "))
i_total_sulfur_dioxide = float(input("Enter total sulfur dioxide [6.0-440.0]: \n
↪"))
i_density = float(input("Enter density [0.98-1.04]: "))
i_ph = float(input("Enter pH [2.72-4.01]: "))
i_sulphates = float(input("Enter sulphates [0.22-2.0]: "))
i_type_white = bool(input("Enter 1 if wine is white, 0 if red: "))
i_alcohol = float(input("Enter alcohol percentage [8.0-14.9]: "))

i_predict = np.array([i_fixed_acidity, i_volatile_acidity, i_citric_acid, \n
↪i_residual_sugar, i_chlorides, i_total_sulfur_dioxide, i_ph, i_sulphates, \n
↪((i_alcohol*5)/i_density), i_type_white]).reshape(1,-1)
scaler.fit(i_predict)

i_predict_std = scaler.transform(i_predict)

predict_output = rfc.predict(i_predict_std)
predict_output_int = predict_output[0]
print(predict_output_int)

```

```

Enter fixed acidity [3.8-15.9]: 4
Enter volatile acidity [0.08-1.58]: 1
Enter citric acid [0.0-1.66]: 1
Enter residual sugar [0.6-65.8]: 2
Enter chlorides [0.009-0.611]: 0.5
Enter total sulfur dioxide [6.0-440.0]: 22
Enter density [0.98-1.04]: 1
Enter pH [2.72-4.01]: 2
Enter sulphates [0.22-2.0]: 1.2
Enter 1 if wine is white, 0 if red: 1
Enter alcohol percentage [8.0-14.9]: 17

```

1

```

[ ]: #6.7      0.23      0.31      2.1      0.046      30.0      0.
↪99260      3.33      0.64      10.7

```

[1]:

```

-----
NameError                                Traceback (most recent call last)
Cell In[1], line 1
----> 1 predict_output_int = int(predict_output[0])
      2 print(predict_output_int)

```

```
NameError: name 'predict_output' is not defined
```

```
[ ]:
```