

```
import pandas as pd
```

▼ Load data

```
# load the CSV file as a pandas DataFrame
df = pd.read_csv('/content/agaricus-lepiota.data', delimiter=',')
df.columns = ['class', 'cap-shape', 'cap-surface', 'cap-color', 'bruises?',
              'odor', 'gill-attachment', 'gill-spacing', 'gill-size',
              'gill-color', 'stalk-shape', 'stalk-root', 'stalk-surface-above-ring',
              'stalk-surface-below-ring', 'stalk-color-above-ring', 'stalk-color-below-ring',
              'veil-type', 'veil-color', 'ring-number', 'ring-type', 'spore-print-color',
              'population', 'habitat']

print(df)
```

```
↗
```

	class	cap-shape	cap-surface	...	spore-print-color	population	habitat
0	e	x	s	...	n	n	g
1	e	b	s	...	n	n	m
2	p	x	y	...	k	s	u
3	e	x	s	...	n	a	g
4	e	x	y	...	k	n	g
...
8118	e	k	s	...	b	c	l
8119	e	x	s	...	b	v	l
8120	e	f	s	...	b	c	l
8121	p	k	y	...	w	v	l
8122	e	x	s	...	o	c	l

```
[8123 rows x 23 columns]
```

▼ Pre-processing

```
# Label encoding
from sklearn.preprocessing import LabelEncoder

le = LabelEncoder()
df_encoded = df.apply(le.fit_transform)
print(df_encoded)
```

```
↗
```

	class	cap-shape	cap-surface	...	spore-print-color	population	habitat
0	0	5	2	...	3	2	1
1	0	0	2	...	3	2	3
2	1	5	3	...	2	3	5
3	0	5	2	...	3	0	1
4	0	5	3	...	2	2	1
...
8118	0	3	2	...	0	1	2
8119	0	5	2	...	0	4	2
8120	0	2	2	...	0	1	2
8121	1	3	3	...	7	4	2
8122	0	5	2	...	4	1	2

[8123 rows x 23 columns]

```
# Extract class column
classes = df_encoded['class']

# Remove class column and 'stalk-root' column (has missing values) from attribute dataframe
df_final = df_encoded.drop(labels=['class', 'stalk-root'], axis=1)

# Split data into train and test data in preparation for classification
from sklearn.model_selection import train_test_split

# where X is the attribute dataframe and Y is the classes dataframe
X_train, X_test, Y_train, Y_test = train_test_split(df_final, classes, test_size=0.2,
                                                    random_state=1)
```

▼ Naive Bayes

```
# Naive Bayes
from sklearn.naive_bayes import CategoricalNB

cNB = CategoricalNB()
cNB.fit(X_train, Y_train)
cNB_predictions = cNB.predict(X_test)
print(cNB_predictions)
#where 0 is edible and 1 is poisonous
```

```
☞ [0 0 1 ... 1 0 0]
```

▼ Support Vector Machine

```
from sklearn import svm
```

```
-svm = svm.SVC()
```

```
cSVM = svm.SVC()
cSVM.fit(X_train, Y_train)
cSVM_predictions = cSVM.predict(X_test)
print(cSVM_predictions)
#where 0 is edible and 1 is poisonous
```

```
↳ [0 1 1 ... 1 0 0]
```

▼ Analysis

```
# Accuracy
from sklearn.metrics import accuracy_score

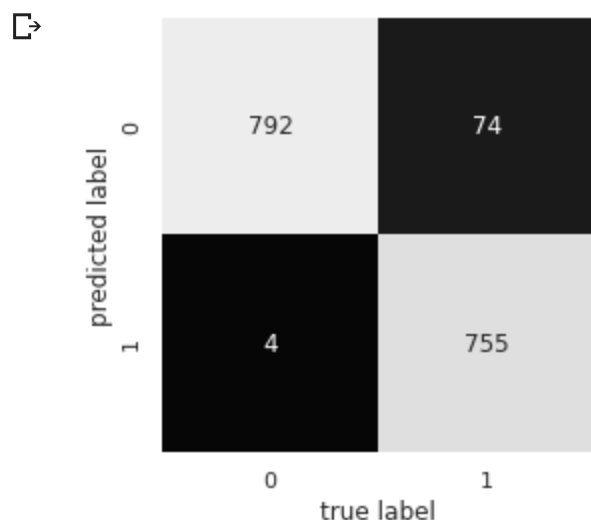
cNB_accuracy = accuracy_score(Y_test, cNB_predictions)
cSVM_accuracy = accuracy_score(Y_test, cSVM_predictions)

print("Naive Bayes Accuracy:", cNB_accuracy)
print("SVM Accuracy:", cSVM_accuracy)
```

```
↳ Naive Bayes Accuracy: 0.952
   SVM Accuracy: 0.9876923076923076
```

```
from sklearn.metrics import confusion_matrix
import matplotlib.pyplot as plt
import seaborn as sns; sns.set()

labels = [0, 1]
mat = confusion_matrix(Y_test, cNB_predictions)
sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False,
            xticklabels=labels, yticklabels=labels)
plt.xlabel('true label')
plt.ylabel('predicted label');
```



```
labels = [0, 1]
mat = confusion_matrix(Y_test, cSVM_predictions)
sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False,
             xticklabels=labels, yticklabels=labels)
plt.xlabel('true label')
plt.ylabel('predicted label');
```

