Project_3_Weihua_PAN1

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IE 7300: Statistical learning for Engineering

Project check point #3

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Percentage of Effort Contributed by Student : 100% Signature of Student : Weihua Pan Submission

Date: 11/28/2023

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

pd.set_option('display.max_columns',30)
```

1 a) Problem Statement

1.1 Clearly define the problem statement

Problem statement: To determine whether some features of Mushroom, such as cap diameter, cap shape and cap color, can be used to predict its edibility.

1.2 State your hypothesis

Hypothesis: Mushrooms with specify feature characteristic are more likely to be edible or poisonous.

2 Dataset

2.1 Present the dataset and include a data dictionary

```
[]: df = pd.read_csv("MushroomDataset/secondary_data.csv",sep=";")
    df.info()
    df.head()

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

RangeIndex: 61069 entries, 0 to 61068

Data	columns (total 21 col	umns):	
#	Column	Non-Null Count	Dtype
0	class	61069 non-null	object
1	cap-diameter	61069 non-null	float64
2	cap-shape	61069 non-null	object
3	cap-surface	46949 non-null	object
4	cap-color	61069 non-null	object
5	does-bruise-or-bleed	61069 non-null	object

6 gill-attachment 51185 non-null object 7 gill-spacing 36006 non-null object 8 gill-color 61069 non-null object stem-height 61069 non-null float64 stem-width float64 10 61069 non-null

11 stem-root 9531 non-null object 12 stem-surface 22945 non-null object

13 stem-color 61069 non-null object 14 veil-type 3177 non-null object

15 veil-color 7413 non-null object 16 has-ring 61069 non-null object

17 ring-type 58598 non-null object

18 spore-print-color 6354 non-null object 19 habitat 61069 non-null object

20 season 61069 non-null object

dtypes: float64(3), object(18)

У

memory usage: 9.8+ MB

4

[]:		class	cap-dia	ameter	cap-	shape	cap-	-surfac	ce caj	p-color	doe	s-bruise-	or-bleed	\	
	0	p		15.26		x			g	0			f		
	1	p		16.60		x			g	0			f		
	2	p		14.07		x			g	0			f		
	3	p		14.17		f			h	е			f		
	4	p		14.64		X			h	0			f		
		øill−at	tachme	nt øill	-spa	cing	rill-	-color	ster	m-height	. s	tem-width	stem-ro	ot.	\
	0	8		е е		NaN	5	W		16.95		17.09		s	•
	1			e		NaN		W		17.99		18.19		S	
	2			e		NaN		W		17.80		17.74		S	
	3			e		NaN		W		15.77		15.98		S	
	4			е		NaN		W		16.53		17.20		s	
		stem-su	rface :	stem-co	olor	veil-	type	veil-	color	has-rin	ng r	ing-type	\		
	0		У		W		u		W		t	g			
	1		У		W		u		W		t	g			
	2		У		W		u		W		t	g			
	3		У		W		u		W		t	р			

u

t

p

```
spore-print-color habitat season
0
                                       d
                        {\tt NaN}
                                                  W
                                       d
1
                        {\tt NaN}
                                                  u
2
                        {\tt NaN}
                                       d
                                                  W
3
                        {\tt NaN}
                                       d
                                                  W
4
                        {\tt NaN}
                                       d
                                                  W
```

The dataset contains 61069 rows and 21 columns. The target variable is class (e or p).

```
[]: df.select_dtypes(include='object')
```

[]:		class	cap-s	shape	cap-s	surface	cap-	color	doe	es-bruis	se-c	or-bleed	\		
	0	p	•	х	•	g	•	0				f			
	1	p		х		g		0				f			
	2	p		х		g		0				f			
	3	p		f		h		е				f			
	4	p		x		h		0				f			
	•••		•••		•••	•••				•••					
	61064	р		s		s		У				f			
	61065	р		f		s		У				f			
	61066	р		s		s		У				f			
	61067	р		f		s		У				f			
	61068	p		s		s		У				f			
														,	
	^	gill-a	attach		gill-	-	-	T-coTc		stem-roo		stem-surf		\	
	0			е		NaN			W		s		У		
	1			е		NaN			W		s		У		
	2			е		NaN			W		S		У		
	3			е		NaN			W		S		У		
	4			е		NaN	J		W		s		У		
			•							NT -	 - NT		NT - NT		
	61064			f		f			f	Na			NaN		
	61065			f		f			f	Na			NaN		
	61066			f		f			f	Na			NaN		
	61067			f		f			f	Na			NaN		
	61068			f		f	:		f	Na	aN		NaN		
		stem-c	color	veil-	-type	veil-co	olor	has-ri	ng	ring-ty	ype	spore-pr	int-	color	\
	0		W		u		W		t		g			${\tt NaN}$	
	1		W		u		W		t		g			NaN	
	2		W		u		W		t		g			${\tt NaN}$	
	3		W		u		W		t		р			${\tt NaN}$	
	4		W		u		W		t		p			NaN	
	•••	•		•••			•••	••				•••			
	61064		У		NaN		NaN		f		f			${\tt NaN}$	
	61065		У		NaN		NaN		f		f			NaN	

61066	У	NaN	NaN	f	f	NaN
61067	У	NaN	NaN	f	f	NaN
61068	У	NaN	NaN	f	f	NaN

	habitat	season
0	d	W
1	d	u
2	d	W
3	d	W
4	d	W
•••		
61064	d	a
61065	d	a
61066	d	u
61067	d	u
61068	d	u

[61069 rows x 18 columns]

17 of them are categorical feature: ['cap-shape', 'cap-surface', 'cap-color', 'does-bruise-or-bleed', 'gill-attachment', 'gill-spacing', 'gill-color', 'stem-root', 'stem-surface', 'stem-color', 'veil-type', 'veil-color', 'has-ring', 'ring-type', 'spore-print-color', 'habitat', 'season'] I will do one-hot-encode later.

[]: df.select_dtypes(include='number')

[]:		cap-diameter	stem-height	stem-width
г ј.		cap diameter	•	Prem Miden
	0	15.26	16.95	17.09
	1	16.60	17.99	18.19
	2	14.07	17.80	17.74
	3	14.17	15.77	15.98
	4	14.64	16.53	17.20
	•••	•••	•••	•••
	61064	1.18	3.93	6.22
	61065	1.27	3.18	5.43
	61066	1.27	3.86	6.37
	61067	1.24	3.56	5.44
	61068	1.17	3.25	5.45

[61069 rows x 3 columns]

3 of them are numerical: ['cap-diameter', 'stem-height', 'stem-width'].

2.1.1 Dataset Dictionary

Target variable: class divided in edible=e and poisonous=p n: nominal, m: metrical 1. cap-diameter (m): float number in cm 2. cap-shape (n): * bell=b, * conical=c, * convex=x, * flat=f, * sunken=s, * spherical=p, * others=o 3. cap-surface (n): * fibrous=i, * grooves=g, * scaly=y, * smooth=s, * shiny=h, * leathery=l, * silky=k, * sticky=t, * wrinkled=w, * fleshy=e 4. cap-color

(n): * brown=n, * buff=b, * gray=g, * green=r, * pink=p, * purple=u, * red=e, * white=w, * yellow=y, * blue=l, * orange=o, * black=k 5. does-bruise-bleed (n): * bruises-or-bleeding=t, * no=f 6. gill-attachment (n): * adnate=a, * adnexed=x, * decurrent=d, * free=e, * sinuate=s, * pores=p, * none=f, * unknown=? 7. gill-spacing (n): * close=c, * distant=d, * none=f 8. gill-color (n): * see cap-color + * none=f 9. stem-height (m): float number in cm 10. stem-width (m): float number in mm 11. stem-root (n): * bulbous=b, * swollen=s, * club=c, * cup=u, * equal=e, * rhizomorphs=z, * rooted=r 12. stem-surface (n): see cap-surface + none=f 13. stem-color (n): see cap-color + none=f 14. veil-type (n): partial=p, universal=u 15. veil-color (n): see cap-color + none=f 16. has-ring (n): * ring=t, * none=f 17. ring-type (n): * cobwebby=c, * evanescent=e, * flaring=r, * grooved=g, * large=l, * pendant=p, * sheathing=s, * zone=z, * scaly=y, * movable=m, * none=f, * unknown=? 18. spore-print-color (n): see cap color 19. habitat (n): * grasses=g, * leaves=l, * meadows=m, * paths=p, * heaths=h, * urban=u, * waste=w, * woods=d 20. season (n): * spring=s, * summer=u, * autumn=a, * winter=w

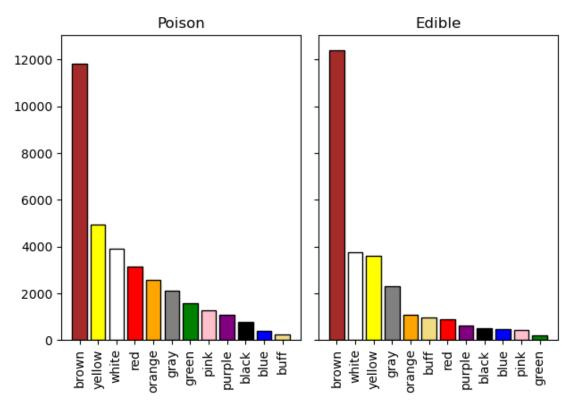
Most of the columns are categorical, so I need to do one-hot-encoding for these categorical data in order to train a model

2.2 Explain how this dataset supports your hypothesis

```
[]: | # split mushrooms df into poison and edible df.
     poison = df[df['class']=='p']
     edible = df[df['class']=='e']
[]: print(poison.shape)
     print(edible.shape)
    (33888, 21)
    (27181, 21)
[]: p color = poison['cap-color'].value counts().index
     p_counts = poison['cap-color'].value_counts().values
     e color = edible['cap-color'].value counts().index
     e counts = edible['cap-color'].value counts().values
[]: color_code = {"n":"brown",
                  "b": "#f0dc82", #buff
                  "g":"gray",
                  "r": "green",
                  "p":"pink",
                  "u": "purple",
                  "e":"red",
                  "w":"white",
                  "y":"yellow",
                  "1":"blue".
                  "o":"orange",
                  "k":"black"}
```

```
[]: import warnings
     warnings.filterwarnings('ignore')
     fig,ax = plt.subplots(1,2,sharey=True)
     ax[0].bar([color_map[c] for c in p_color],p_counts,color=[color_code[code] for_
      ⇔code in p_color],
               edgecolor='black')
     ax[0].set_title('Poison')
     ax[1].bar([color_map[c] for c in e_color],e_counts,color=[color_code[code] for_
      ⇔code in e_color],
               edgecolor='black')
     ax[1].set_title('Edible')
     # rotate xlabels
     ax[0].set_xticklabels([color_map[c] for c in p_color],rotation=90)
     ax[1].set_xticklabels([color_map[c] for c in e_color],rotation=90)
     fig.suptitle("Barchart of cap-color between Poison and Edible")
     plt.tight_layout()
     plt.show()
```





Using cap-color and other features, we are able to find some pattern to classify base on these features. For example, From the above plot, We can see most green and red mushroom are poisonous. In addition, the most common color is brown for both poison and edible mushrooms, Therefore, it might be tough to use brown to classify whether a mushroom is poisonous or edible.

3 Exploratory Data Analysis(EDA)

3.1 Descriptive Statistics

[]: df.describe()

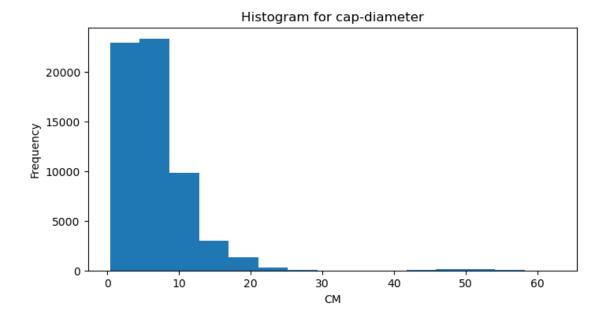
[]:		cap-diameter	stem-height	stem-width
	count	61069.000000	61069.000000	61069.000000
	mean	6.733854	6.581538	12.149410
	std	5.264845	3.370017	10.035955
	min	0.380000	0.000000	0.000000
	25%	3.480000	4.640000	5.210000
	50%	5.860000	5.950000	10.190000
	75%	8.540000	7.740000	16.570000
	max	62.340000	33.920000	103.910000

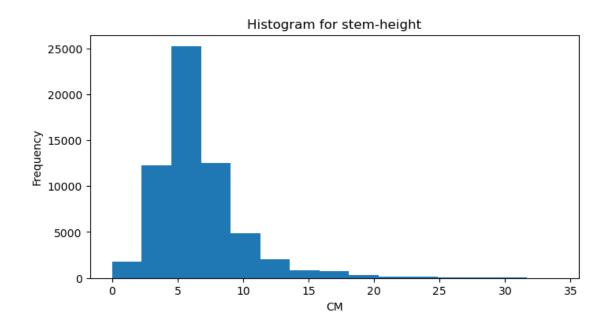
Only 3 columns are numerical. 1. cap-diameter: * min: 0.38 cm * max: 62.34 cm * mean 6.7 cm 2. stem-height: * min: 0 cm * max: 33.92 cm * mean 6.58 cm 3. stem-width: * min: 0 cm * max: 103.91 cm * mean: 10.03 cm

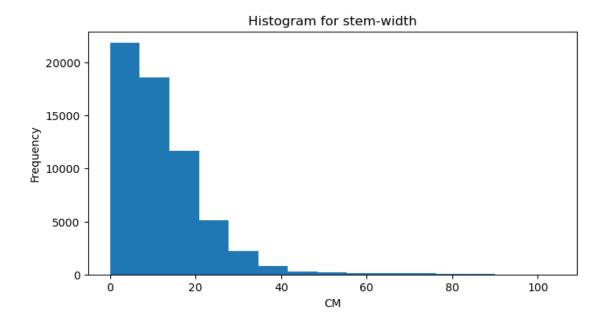
Base on the basic statistic, I can take a guess cap-diameter and stem-height are in right skewed, and stem-width is about normal by mean and 50% quartile. And they all have some outliers. I will consider whether remove them later.

```
[]: # draw histogram for all numerical columns
numerical_columns = df.select_dtypes(include='number')

for column in numerical_columns:
    fig, ax = plt.subplots(figsize=(8, 4))
    df[column].plot(kind='hist', title=f"Histogram for {column}", ax=ax,bins=15)
    ax.set_xlabel("CM")
    plt.show()
```







By the histogram plot, we can see cap-diameter and stem-width are right skewed. And only stem-height is normal distributive.

3.2 Bar chart among 2 target variables

```
[]: object_columns = df.select_dtypes(include=['object']).columns[1:]

for column in object_columns:
    # Count the occurrences of each category
    poison_C = poison[column].value_counts()
    edible_C = edible[column].value_counts()

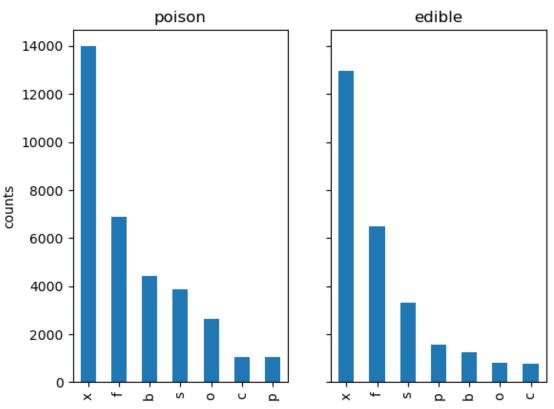
# Create a subplot chart
fig,ax = plt.subplots(1,2,sharey=True)

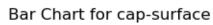
poison_C.plot(kind='bar',ax=ax[0],title='poison',ylabel='counts')
    edible_C.plot(kind='bar',ax=ax[1],title='edible')

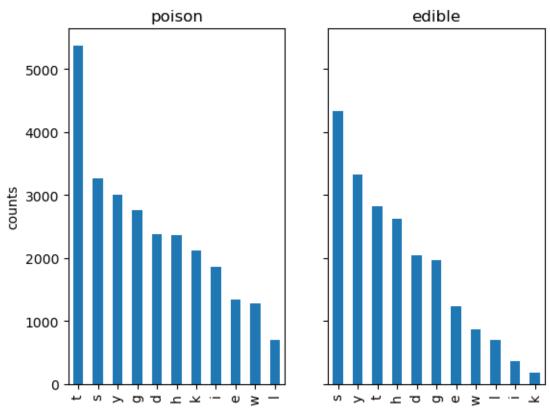
fig.suptitle(f'Bar Chart for {column}')

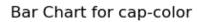
# Show the plot
plt.show()
```

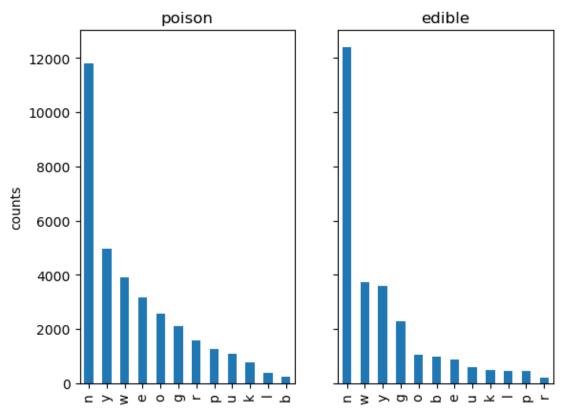
Bar Chart for cap-shape



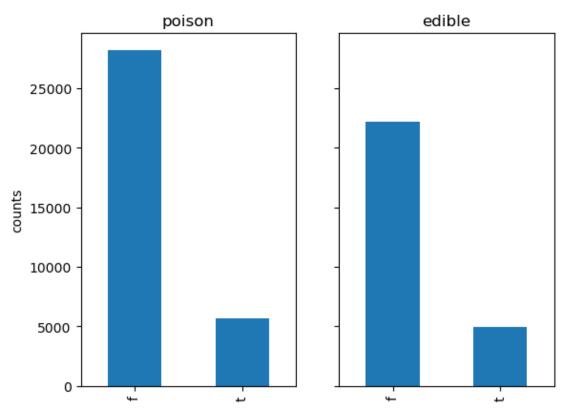




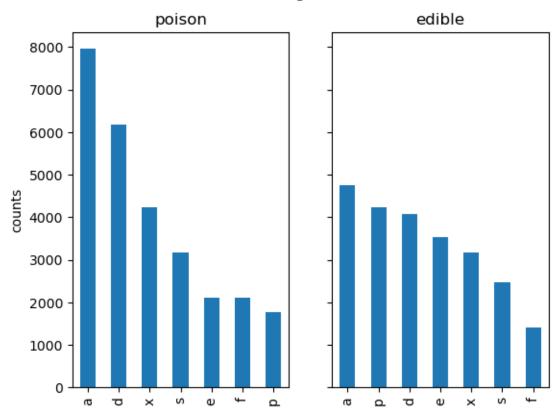


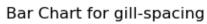


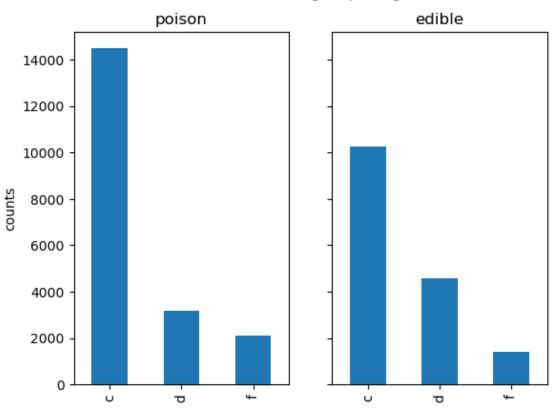
Bar Chart for does-bruise-or-bleed

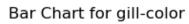


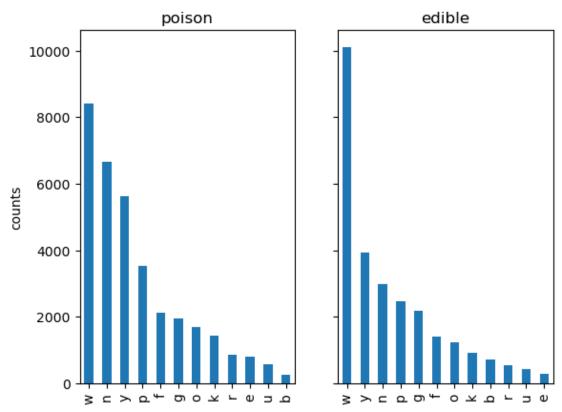
Bar Chart for gill-attachment



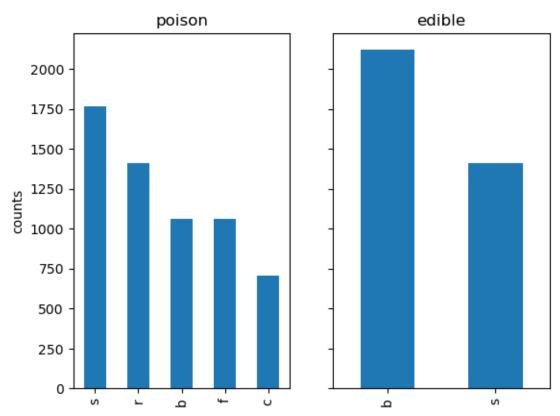




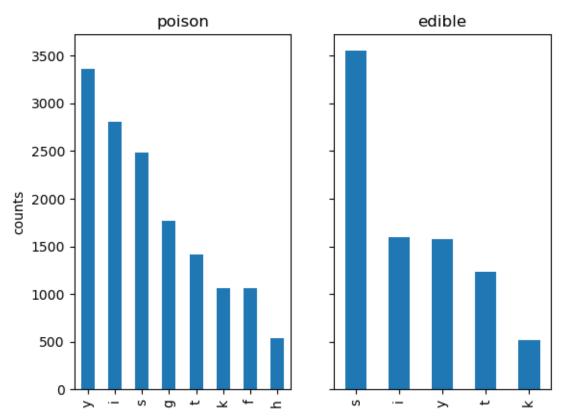




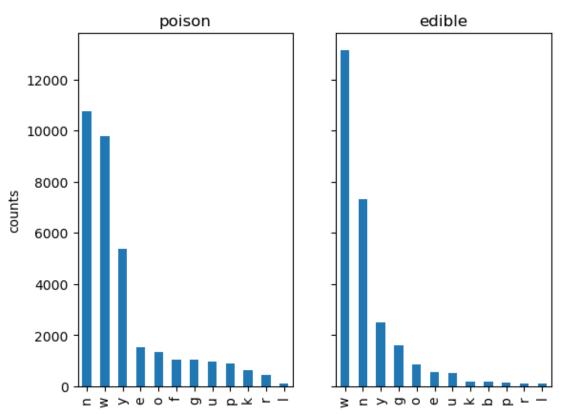
Bar Chart for stem-root

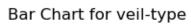


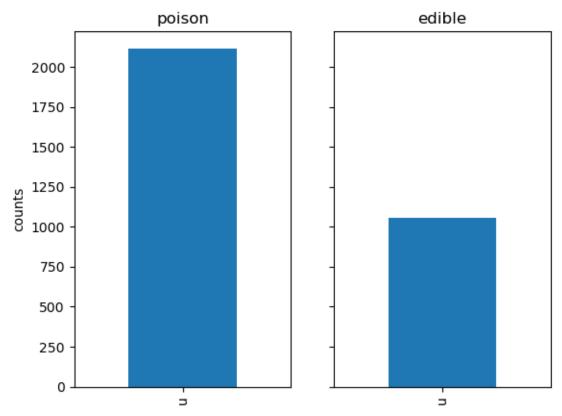
Bar Chart for stem-surface

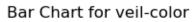


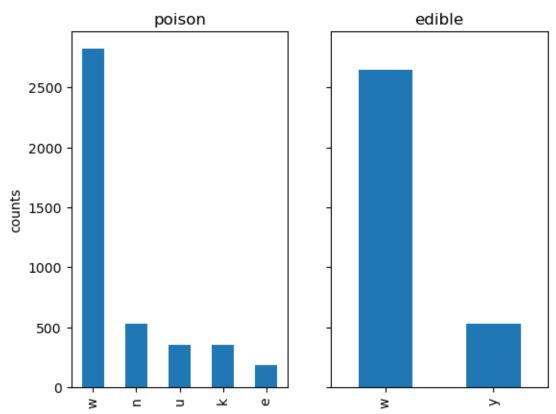
Bar Chart for stem-color

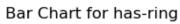


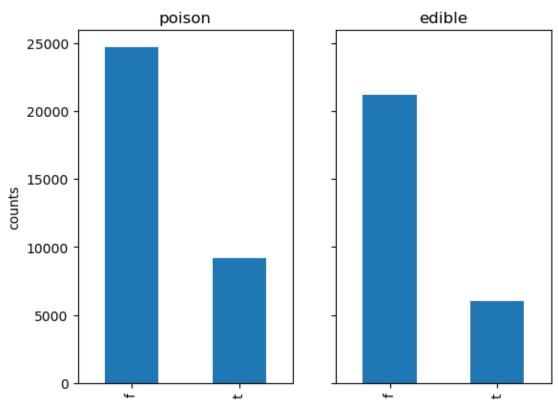


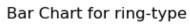


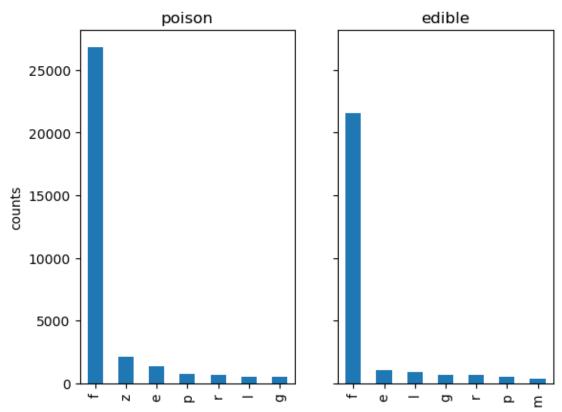




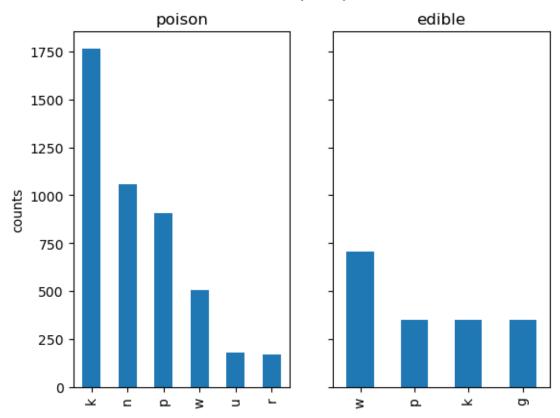




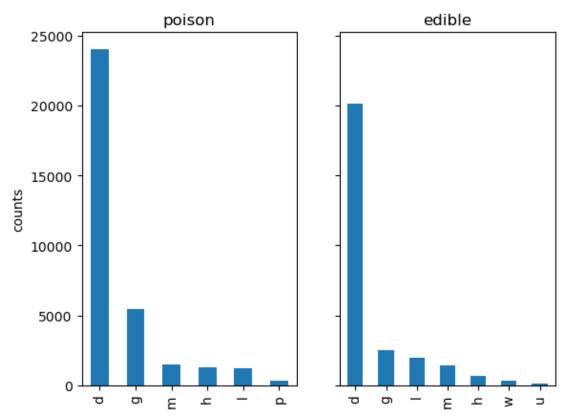




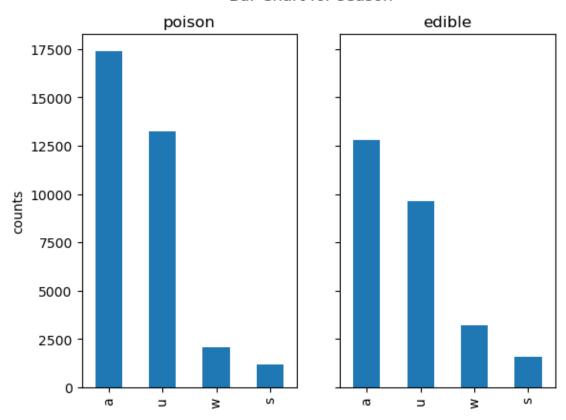
Bar Chart for spore-print-color



Bar Chart for habitat



Bar Chart for season



By the above bar charts, we can find some pattern to classify poisonous mushroom by categorical features. For example, In season bar chart, we can see most poisonous mushrooms live in autumn and summer. And we can find pattern in veil-color as well. We can easily classify the mushrooms with yellow veil is edible since there is no poisonous mushroom with yell veil. All the veil-type are universal, other than that are missing values.

3.3 Missing values

```
[]: NA_dict = dict()
for column in df.columns[1:]:
     NA_dict[column] = df[column].isna().sum()

NA_dict
```

```
'gill-spacing': 25063,
'gill-color': 0,
'stem-height': 0,
'stem-width': 0,
'stem-root': 51538,
'stem-surface': 38124,
'stem-color': 0,
'veil-type': 57892,
'veil-color': 53656,
'has-ring': 0,
'ring-type': 2471,
'spore-print-color': 54715,
'habitat': 0,
'season': 0}
```

cap-surface,gill-attachment,gill-spacing,stem-root,stem-surface,veil-type,veil-color,ring-type,spo have missing values. And all of them are categorical. All the missing values here might means unknown or not applicable. I decide to impute them using "?", since I have no idea how these missing values appear and cannot apply imputation strategy like mode or k_nearest. We can let the model to decide and find pattern when they meet the missing values "?".

```
[]: df= df.fillna('?')
```

3.4 Feature engineering

```
[]: y = df.iloc[:,0]
X = df.iloc[:,1:]
X = pd.get_dummies(X) # encode categorical data by one-hot-encoding
X = X.astype(float) # convert true and false to 1 and 0
X
```

гл.				b b	h	,
[]:	cap-diameter	stem-neight	stem-width	cap-shape_b	cap-snape_c	\
0	15.26	16.95	17.09	0.0	0.0	
1	16.60	17.99	18.19	0.0	0.0	
2	14.07	17.80	17.74	0.0	0.0	
3	14.17	15.77	15.98	0.0	0.0	
4	14.64	16.53	17.20	0.0	0.0	
•••	•••	•••	•••			
61064	1.18	3.93	6.22	0.0	0.0	
61065	1.27	3.18	5.43	0.0	0.0	
61066	1.27	3.86	6.37	0.0	0.0	
61067	1.24	3.56	5.44	0.0	0.0	
61068	1.17	3.25	5.45	0.0	0.0	
	cap-shape_f	cap-shape_o	cap-shape_p	cap-shape_s	cap-shape_x	\
0	0.0	0.0	0.0	0.0	1.0	
1	0.0	0.0	0.0	0.0	1.0	

2 3	0.0 1.0		0.0	0.0		0.0		1.0	
4	0.0		0.0	0.0		0.0		1.0	
 61064	 0.0	•••	0.0	0.0	•••	1.0		0.0	
61065	1.0		0.0	0.0		0.0		0.0	
61066	0.0		0.0	0.0		1.0		0.0	
61067	1.0		0.0	0.0		0.0		0.0	
61068	0.0		0.0	0.0		1.0		0.0	
01000	0.0		0.0	0.0		1.0		0.0	
	<pre>cap-surface_?</pre>	cap-s	-	cap-surf		cap-surf	_	\	
0	0.0		0.0		0.0		1.0		
1	0.0		0.0		0.0		1.0		
2	0.0		0.0		0.0		1.0		
3	0.0		0.0		0.0		0.0		
4	0.0		0.0		0.0		0.0		
	•••		•••	•••		•••			
61064	0.0		0.0		0.0		0.0		
61065	0.0		0.0		0.0		0.0		
61066	0.0		0.0		0.0		0.0		
61067	0.0		0.0		0.0		0.0		
61068	0.0		0.0		0.0		0.0		
01000	0.0		0.0		0.0		0.0		
	cap-surface_h	sp	ore-print-	-color_r	spore	-print-co	lor_u	\	
0	0.0		-	0.0	-	-	0.0		
1	0.0	•••		0.0			0.0		
2	0.0	•••		0.0			0.0		
3	1.0	•••		0.0			0.0		
4	1.0	•••		0.0			0.0		
-		•••					0.0		
 61064	0.0			0.0		•••	0.0		
61065	0.0	•••		0.0			0.0		
61066		•••		0.0					
	0.0	•••					0.0		
61067	0.0	•••		0.0			0.0		
61068	0.0	•••		0.0			0.0		
	spore-print-co	lor_w	habitat o	d habita	t_g h	abitat_h	habita	t_l	\
0	• •	0.0	1.0		0.0	0.0		0.0	
1		0.0	1.0)	0.0	0.0		0.0	
2		0.0	1.0		0.0	0.0		0.0	
3		0.0	1.0		0.0	0.0		0.0	
4		0.0	1.0		0.0	0.0		0.0	
								3.0	
 61064	•	0.0	1.0		0.0	0.0		0.0	
61065		0.0	1.0		0.0	0.0		0.0	
61066		0.0	1.0		0.0	0.0		0.0	
61067		0.0	1.0	,	0.0	0.0		0.0	

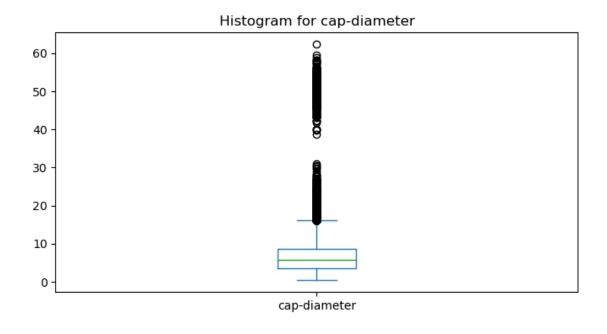
61068		0.0	1.0	0.0	0.0	0.0
	habitat_m	habitat_p	habitat_u	habitat_w	season_a	season_s \
0	0.0	0.0	0.0	0.0	0.0	0.0
1	0.0	0.0	0.0	0.0	0.0	0.0
2	0.0	0.0	0.0	0.0	0.0	0.0
3	0.0	0.0	0.0	0.0	0.0	0.0
4	0.0	0.0	0.0	0.0	0.0	0.0
•••	•••	•••			•••	
61064	0.0	0.0	0.0	0.0	1.0	0.0
61065	0.0	0.0	0.0	0.0	1.0	0.0
61066	0.0	0.0	0.0	0.0	0.0	0.0
61067	0.0	0.0	0.0	0.0	0.0	0.0
61068	0.0	0.0	0.0	0.0	0.0	0.0
	season_u	season_w				
0	0.0	1.0				
1	1.0	0.0				
2	0.0	1.0				
3	0.0	1.0				
4	0.0	1.0				
•••	•••					
61064	0.0	0.0				
61065	0.0	0.0				
61066	1.0	0.0				
61067	1.0	0.0				
61068	1.0	0.0				

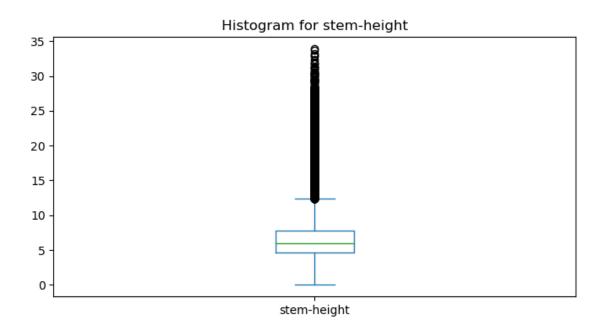
[61069 rows x 128 columns]

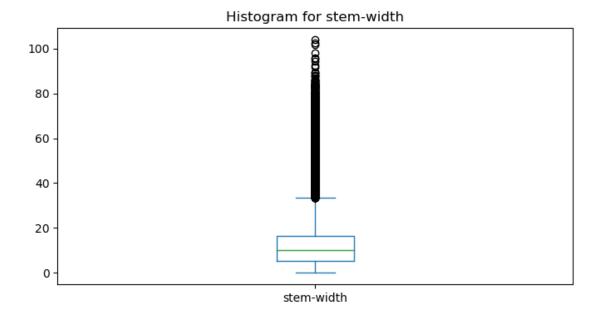
Most columns are categorical, so we need to one-hot-encode the categorical columns. Therefore the models can fit the dataset easier.

3.5 Outlier detection

```
for column in numerical_columns:
    fig, ax = plt.subplots(figsize=(8, 4))
    df[column].plot(kind='box', title=f"Histogram for {column}", ax=ax)
    plt.show()
```







By the box plot, we can see there are a lot of outlier in the positive side for all numerical columns.

```
[]: outliers_dict
```

```
[]: {'cap-diameter': 2400, 'stem-height': 3169, 'stem-width': 1967}
```

Since the amount of outliers is small compare with the total rows, I will not remove any outliers.

And I will use ensemble model. They are able to handle the outliers.

4 Model Training and Evaluation

The models I will use is logistic regression, Decision Tree, and KNN * Logistic Regression: simple model, suitable for binary classification especially when the dataset have linear pattern * Decision Tree: model that is capable for regression and classification. This model is able to find the important features easily, and capture the non-linear pattern which logistic regression cannot. * KNN: simple model that can capture nonlinear pattern, but it require to compute the distance for all data points which would are complex when the dataset is large.

All of these models are easy to understand and easy to implement, and these models are representor of Gradient_based, Tree-based and Distance-based models.

```
[]: from sklearn.model_selection import train_test_split train_X,test_X,train_y,test_y = train_test_split(X,y,test_size=0.

$\times 2$,random_state=42$)
```

4.1 Logistic Regression

```
[]: # import numpy as np
     # import pandas as pd
     # class MyLogisticRegression:
     #
           def __init__ (self, epochs = 10000, threshold=1e-3,
     #
                         regularization=None, alpha=0.01) -> None:
               # Initialize parameters and flags for the model.
     #
               self.epochs = epochs
     #
               self.threshold = threshold
               self.regularization = regularization
     #
     #
               self.alpha = alpha
     #
           def train(self, X, y, batch size=64, lr=1e-3, seed=11, verbose=False):
     #
     #
               Train the model using stochastic gradient descent.
     #
               # Set seed for reproducibility.
     #
     #
               np.random.seed(seed)
     #
               # Define the unique classes and their corresponding indices.
     #
               self.classes = np.unique(y)
               self.class labels = {c: i for i, c in enumerate(self.classes)}
     #
               # Add bias term to the features.
     #
```

```
#
          X = self.add_bias(X)
#
          # Convert labels into one-hot encoded format.
#
          y = self.one_hot(y)
          # Initialize weights matrix with zeros.
#
          self.loss = []
#
          self.weights = np.zeros(shape=(len(self.classes), X.shape[1]))
#
          # Start the training process.
          self.fit_data(X, y, batch_size, lr, verbose)
#
          return self
#
      def fit_data(self, X, y, batch_size, lr, verbose):
#
#
          Fit the data using stochastic gradient descent.
#
#
          i = 0
#
          while (not self.epochs or i < self.epochs):
              # Compute and store the cross-entropy loss.
#
#
              self.loss.append(self.cross_entropy(y, self.predict_(X)))
              # Randomly select a batch of data.
#
              idx = np.random.choice(X.shape[0], batch_size)
#
              X_batch, y_batch = X[idx], y[idx]
#
              # Calculate the error between predicted and true values.
#
              error = y_batch - self.predict_(X_batch)
#
              # Update the weights based on the error and learning rate.
              update = lr * np.dot(error.T, X_batch)
#
              # Apply regularization if specified.
#
#
              if self.regularization == 'Ridge':
                  update += self.alpha * self.weights
#
#
              elif self.regularization == 'Lasso':
#
                  update += self.alpha * np.sign(self.weights)
#
              elif self.regularization == 'Elastic Net':
                  update_w += self.alpha * (self.weights + np.sign(self.
 ⇔weights))
              self.weights += update
#
              # Stop training if updates are smaller than a threshold.
#
              if np.abs(update).max() < self.threshold:</pre>
```

```
#
                   break
              # Print training accuracy every 1000 iterations if verbose is_{\sqcup}
 \hookrightarrow True.
#
              if i \% 1000 == 0 and verbose:
                  \hookrightarrow self.evaluate_(X, y)))
              i \neq 1
      def predict(self, X):
#
          return self.predict_(self.add_bias(X))
#
      def predict (self, X):
#
          pre\_vals = np.dot(X, self.weights.T).reshape(-1,len(self.classes))
#
          return self.softmax(pre_vals)
      def softmax(self, z):
#
          return np.exp(z) / np.sum(np.exp(z), axis=1).reshape(-1,1)
      def predict_classes(self, X):
#
#
          self.probs_{\_} = self.predict(X)
          return np.vectorize(lambda c: self.classes[c])(np.argmax(self.probs_,_
 \rightarrow axis=1))
      def add bias(self, X):
          return np.insert(X, 0, 1, axis=1)
#
      def one_hot(self, y):
          return np.eye(len(self.classes))[np.vectorize(lambda c: self.
 \hookrightarrow class_labels[c])(y).reshape(-1)]
#
      def score(self, X, y):
          111
#
          Accuracy metric
          return\ round(np.mean(self.predict\_classes(X).reshape(-1,1) == y),3)
#
      def evaluate_(self, X, y):
          return \ np.mean(np.argmax(self.predict_(X), \ axis=1) == np.argmax(y, u)
 \rightarrow axis=1))
      def cross_entropy(self, y, probs):
#
          return -1 * np.mean(y * np.log(probs))
#
      def confusion_matrix(self, actual, predicted,norm=False):
#
```

```
Compute the confusion matrix for the given actual and predicted
 \hookrightarrow outputs.
          Args:
          - actual (array-like): Actual outputs (ground truth).
#
          - predicted (array-like): Predicted outputs from the model.
          Returns:
#
          - matrix (np.ndarray): N x N confusion matrix, where N is the number
 ⇔of unique classes.
          11 11 11
          # Create an empty matrix
          matrix = np.zeros((len(self.classes), len(self.classes)), dtype=float)
          # Fill the matrix
#
          for i, true class in enumerate(self.classes):
#
              for j, pred_class in enumerate(self.classes):
                  matrix[i, j] = np.sum((actual == true_class) & (predicted ==_
 →pred_class))
          if norm:
              for i in range(len(self.classes)):
                  total = np.sum(matrix[i])
                  for j in range(len(self.classes)):
                      matrix[i,j] = round(matrix[i,j] / total,2)
          matrix\_df = pd.DataFrame(matrix, index=self.classes, columns=self.
 ⇔classes)
          return matrix_df
```

```
[]: from LogisticRegression import MyLogisticRegression
lm = MyLogisticRegression()
lm.train(train_X.values,train_y.values)
```

[]: <LogisticRegression.MyLogisticRegression at 0x133453650>

4.1.1 Training set evaluation

```
[]: pred_train_y = lm.predict_classes(train_X.values)
lm.confusion_matrix(train_y,pred_train_y)
```

```
e 20183.0 1624.0
p 7378.0 19670.0
```

logistic regression can classify e much better than p, but the overall performance is great. We treat e as positive and p as negative, then if we misclassify e to p, we call it type 1 error(false positive). And misclassify p as e, we call it type 2 error(false negative). Type 2 is bad for this case, because people eat the poisonous mushrooms which were classified as edible. Therefore, we should increase the overall accuracy or decrease the type 2 error.

```
[]: print(f'accuracy: {np.mean(pred_train_y == train_y)}')
```

accuracy: 0.8157404564527684

Logistic regression perform great in training set, acquire 81.5% accuracy.

4.1.2 Test set evaluation

```
[ ]: pred_test_y = lm.predict_classes(test_X)
lm.confusion_matrix(test_y,pred_test_y)
```

```
e p
e 4934.0 440.0
p 1865.0 4975.0
```

```
[]: print(f'accuracy: {np.mean(pred_test_y == test_y)}')
```

accuracy: 0.8112821352546259

The logistic model seems perform about the same in test dataset. It acquires 81.1% accuracy which is just 0.4% lower than the accuracy in test dataset.

4.1.3 Important features

```
[]: pd.set_option('display.max_columns', None)
     pd.DataFrame(lm.weights,index=[0,1],columns=X.columns.insert(0,'bias'))
                                              stem-width cap-shape_b
[]:
            bias
                  cap-diameter stem-height
                                                                        cap-shape c \
                                                             -1.341672
                                                                          -0.191079
     0 - 1.465632
                      0.165223
                                      0.0796
                                                0.092808
     1 1.465632
                     -0.165223
                                     -0.0796
                                               -0.092808
                                                              1.341672
                                                                           0.191079
                                                cap-shape_s
                                                              cap-shape_x
        cap-shape_f
                     cap-shape_o
                                   cap-shape_p
     0
           0.104998
                       -0.926183
                                      0.137476
                                                   0.482723
                                                                 0.268105
                                                  -0.482723
                                                                -0.268105
     1
          -0.104998
                        0.926183
                                     -0.137476
        cap-surface_?
                       cap-surface_d
                                       cap-surface_e
                                                      cap-surface_g
                                                                      cap-surface_h \
     0
             0.229483
                            0.241876
                                            -0.66786
                                                            0.811854
                                                                           0.623892
     1
            -0.229483
                            -0.241876
                                             0.66786
                                                           -0.811854
                                                                          -0.623892
        cap-surface i
                       cap-surface_k cap-surface_l
                                                      cap-surface_s
                                                                      cap-surface_t
     0
            -1.622021
                            -3.011944
                                            1.189832
                                                            0.574294
                                                                          -0.686148
     1
             1.622021
                            3.011944
                                           -1.189832
                                                           -0.574294
                                                                           0.686148
```

```
cap-surface w cap-surface_y cap-color_b cap-color_e cap-color_g \
0
       0.068476
                      0.782634
                                  1.066569
                                              -1.446941
                                                              0.35109
      -0.068476
                     -0.782634
                                  -1.066569
                                                1.446941
                                                             -0.35109
1
   cap-color_k cap-color_l cap-color_n cap-color_o cap-color_p
    -0.618687
                  1.166477
                              0.849755
                                         -0.035764
                                                        -0.606174
0
     0.618687
1
                 -1.166477
                             -0.849755
                                          0.035764
                                                        0.606174
   cap-color_r cap-color_u cap-color_w cap-color_y does-bruise-or-bleed_f \
0
    -2.086839
                 -0.480523
                              0.090853
                                           0.284552
                                                                   -0.827921
1
     2.086839
                  0.480523
                             -0.090853
                                                                    0.827921
                                           -0.284552
   does-bruise-or-bleed_t gill-attachment_? gill-attachment_a \
                                  -0.801695
0
               -0.637711
                                                     -0.893634
1
                0.637711
                                   0.801695
                                                      0.893634
   gill-attachment_d gill-attachment_e gill-attachment_f gill-attachment_p \
0
          -2.158874
                              1.088682
                                                -0.047301
                                                                    2.981584
1
           2.158874
                             -1.088682
                                                 0.047301
                                                                   -2.981584
   gill-attachment_s gill-attachment_x gill-spacing_? gill-spacing_c
0
          -0.306381
                             -1.328013
                                             -1.006156
                                                             -0.38982
1
           0.306381
                             1.328013
                                             1.006156
                                                              0.38982
   gill-spacing_d gill-spacing_f gill-color_b gill-color_e gill-color_f \
0
        -0.022356
                       -0.047301
                                      1.247254
                                                  -0.938048
                                                                -0.047301
        0.022356
                        0.047301
                                     -1.247254
                                                    0.938048
                                                                  0.047301
   gill-color_g gill-color_k gill-color_n gill-color_o gill-color_p
0
       0.14095
                   -0.856506
                                -0.848034
                                               0.174075
                                                              0.24918
1
      -0.14095
                    0.856506
                                  0.848034
                                               -0.174075
                                                              -0.24918
   gill-color_r gill-color_u gill-color_w gill-color_y stem-root_?
0
      0.215163
                    0.012258
                                   0.46076
                                              -1.275383
                                                             2.445175
     -0.215163
                   -0.012258
                                  -0.46076
                                               1.275383
                                                            -2.445175
   stem-root_b stem-root_c stem-root_f stem-root_r stem-root_s \
0
     2.774851
                 -3.522699
                              -0.151454
                                          -2.844927
                                                       -0.166577
    -2.774851
                  3.522699
                               0.151454
1
                                            2.844927
                                                        0.166577
   stem-surface ? stem-surface f stem-surface g stem-surface h \
                                                       -1.049839
         1.212086
                       -0.151454
                                       -3.119523
0
       -1.212086
                        0.151454
                                        3.119523
                                                        1.049839
   stem-surface_i stem-surface_k stem-surface_s stem-surface_t \
        -0.006051
0
                       -0.816623
                                        1.511848
                                                        2.198119
1
        0.006051
                        0.816623
                                       -1.511848
                                                       -2.198119
```

```
stem-surface v stem-color b stem-color e stem-color f stem-color g \
                      0.460045
0
       -1.244194
                                  -1.315729
                                                 -0.151454
                                                               1.145737
1
        1.244194
                     -0.460045
                                    1.315729
                                                 0.151454
                                                              -1.145737
   stem-color_k stem-color_l stem-color_n stem-color_o stem-color_p \
0
     -0.975333
                    0.560633
                                 -0.059773
                                               0.901764
                                                             -1.59306
1
      0.975333
                   -0.560633
                                  0.059773
                                              -0.901764
                                                              1.59306
   stem-color_r stem-color_u stem-color_w stem-color_y veil-type_? \
0
      -0.379744
                   -0.795127
                                               -0.578656
                                                            1.196461
                                  1.315065
1
      0.379744
                    0.795127
                                 -1.315065
                                               0.578656
                                                           -1.196461
   veil-type_u veil-color_? veil-color_e veil-color_k veil-color_n
0
    -2.662093
                  -0.688759
                                -0.540438
                                             -0.184157
                                                           -1.108955
1
                                                            1.108955
     2.662093
                   0.688759
                                 0.540438
                                              0.184157
   veil-color_w veil-color_y has-ring_f has-ring_t \
0
     -1.147823
                    0.706619
                                 1.497881
                                                        -1.743884
                                             0.278252
                   -0.706619
                                            -0.278252
      1.147823
                                 -1.497881
                                                         1.743884
   ring-type_? ring-type_e ring-type_f ring-type_g ring-type_l \
0
      0.88374
                 -1.047384
                             -1.160445
                                           0.750416
                                                        0.387372
      -0.88374
                  1.047384
                              1.160445
                                          -0.750416
                                                       -0.387372
1
   ring-type_m ring-type_p ring-type_r ring-type_z spore-print-color_? \
                                          -3.020089
     1.548294
                 -0.894767
                                1.08723
                                                               -0.029702
0
    -1.548294
                  0.894767
                               -1.08723
                                           3.020089
                                                                0.029702
   spore-print-color_g spore-print-color_k spore-print-color_n
                                                     -1.083105
0
             1.168732
                                 -1.231195
1
            -1.168732
                                  1.231195
                                                      1.083105
                                            spore-print-color_u
   spore-print-color_p
                       spore-print-color_r
                                  -0.08949
0
            -1.023482
                                                     -0.186547
1
             1.023482
                                   0.08949
                                                       0.186547
   spore-print-color_w habitat_d habitat_g habitat_h habitat_l habitat_m \
             1.009156 -0.585411 -0.645654 -0.198758 -0.226998
0
                                                                    0.24723
1
            -1.009156
                       0.585411 0.645654
                                             0.198758
                                                        0.226998
                                                                   -0.24723
   habitat_p habitat_u habitat_w season_a season_s season_u season_w
                        0.504056 -0.962381 0.229055 -1.014429 0.282124
0 -0.693242
              0.133146
   0.693242 -0.133146 -0.504056 0.962381 -0.229055 1.014429 -0.282124
```

- cap-surface_g: 0.81
- cap-color l: 1.16

• ring-type_m: 1.54

These are the important features. The important feature in logistic regression would have higher weight

4.2 Decision Tree

```
[]: # import numpy as np
     # import pandas as pd
    # class Node():
          def __init__(self, feature_index=None, threshold=None, left=None,_
      →right=None, var_red=None, info_gain=None, value=None):
              ''' constructor '''
              # for decision node
              self.feature_index = feature_index
              self.threshold = threshold # threshold for splitting data
     #
              self.left = left
              self.right = right
              self.var_red = var_red # index for regressor, choose the highest
      ⇔feature to split
              self.info\_gain = info\_gain \# index for classifier, choose the highest \sqcup
     ⇔to split
              # for leaf node
              self.value = value
    # class DecisionTreeRegressor():
          def __init__(self, min_samples_split=2, max_depth=2):
              ''' constructor '''
    #
              # initialize the root of the tree
    #
              self.root = None
     #
              # stopping conditions
              self.min_samples_split = min_samples_split # min elements need to_
     \hookrightarrowsplit
              self.max_depth = max_depth # max depth for tree
          def build_tree(self, dataset, curr_depth=0):
              ''' recursive function to build the tree '''
    #
              X, Y = dataset[:,:-1], dataset[:,-1]
    #
     #
              num_samples, num_features = np.shape(X)
```

```
best\_split = \{\}
          # split until stopping conditions are met
#
          if num samples>=self.min samples split and curr depth<=self.max depth:
#
               # find the best split
              best\_split = self.get\_best\_split(dataset, num\_samples, 
 →num_features)
#
               # check if information gain is positive
              if best split["var red"]>0:
#
                   # recur left
                   left_subtree = self.build_tree(best_split["dataset_left"],__
 \hookrightarrow curr_depth+1)
#
                   # recur right
                   right_subtree = self.build_tree(best_split["dataset_right"],__
 \hookrightarrow curr_depth+1)
                   # return decision node
                   return Node(best_split["feature_index"],_
 ⇔best_split["threshold"],
                               left subtree, right subtree,
 ⇒best_split["var_red"])
          # compute leaf node
#
          leaf_value = self.calculate_leaf_value(Y)
          # return leaf node
#
          return Node(value=leaf_value)
      def get_best_split(self, dataset, num_samples, num_features):
#
#
#
          function to find the best split
#
          Returns:
#
              Dict: A dict that contains information for best split
#
          # dictionary to store the best split
#
          best_split = {}
#
          max\_var\_red = -float("inf")
#
          # loop over all the features
#
          for feature_index in range(num_features):
#
              feature_values = dataset[:, feature_index]
#
              possible_thresholds = np.unique(feature_values)
#
              # loop over all the feature values present in the data
              for threshold in possible_thresholds:
#
                   # get current split
                   dataset_left, dataset_right = self.split(dataset,__
 → feature_index, threshold)
#
                   # check if childs are not null
```

```
#
                  if len(dataset_left)>0 and len(dataset_right)>0:
                      y, left_y, right_y = dataset[:, -1], dataset_left[:, -1],
 ⇔dataset_right[:, -1]
#
                      # compute information gain
                      curr_var_red = self.variance_reduction(y, left_y, right_y)
#
#
                      # update the best split if needed
                      # if curr is higher, update everything
#
#
                      if curr var red>max var red:
#
                           best_split["feature_index"] = feature_index
                           best_split["threshold"] = threshold
#
#
                           best_split["dataset_left"] = dataset_left
#
                           best_split["dataset_right"] = dataset_right
#
                           best_split["var_red"] = curr_var_red
#
                          max\_var\_red = curr\_var\_red
          # return best split
#
          return best split
#
      def split(self, dataset, feature index, threshold):
#
          ''' function to split the data using a feature and threshold
          Returns:
              Tuple: dataset_left, dataset_right
          dataset_left = np.array([row for row in dataset if_
 →row[feature_index]<=threshold])
          dataset_right = np.array([row for row in dataset if_
 →row[feature_index]>threshold])
          return dataset_left, dataset_right
      def variance_reduction(self, parent, l_child, r_child):
#
          ''' function to compute variance reduction
#
          Returns:
              float: measure the information gain for a certain split
          111
#
          weight_l = len(l_child) / len(parent)
          weight_r = len(r_child) / len(parent)
          reduction = np.var(parent) - (weight_l * np.var(l_child) + weight_r *_u
 \hookrightarrow np.var(r child))
          return reduction
      def calculate_leaf_value(self, Y):
#
          ''' function to compute leaf node using mean
```

```
#
          Returns:
              float: mean value for given node value
#
          val = np.mean(Y)
#
          return val
#
      def print_tree(self, tree=None, indent=" "):
          ''' function to print the tree '''
#
#
          if not tree:
               tree = self.root
#
          if tree.value is not None:
#
              print(tree.value)
#
          else:
              print("X_"+str(tree.feature_index), "<=", tree.threshold, "?", "</pre>
 \hookrightarrow tree.var_red)
              print("%sleft:" % (indent), end="")
#
#
              self.print tree(tree.left, indent + indent)
#
              print("%sright:" % (indent), end="")
#
              self.print_tree(tree.right, indent + indent)
      def pretty_print(self,node=None,depth=0,prefix="Root: "):
#
#
          '''Print tree in pretty format using post order traversal'''
          if node is None:
              node = self.root
#
          indent = " " * depth * 4
          if node.value is not None:
              print(f"{indent}{prefix}Leaf => Value: {round(node.value,2)} /__
 \rightarrow depth: {depth}")
#
          else:
#
              self.pretty_print(node.right, depth + 1, prefix="R--> ")
              print(f"{indent}{prefix} X {node.feature index} <= {round(node.</pre>
 →threshold,2)} / depth: {depth}")
              self.pretty_print(node.left, depth + 1, prefix="L--> ")
#
      def fit(self, X, Y):
#
          ''' function to train the tree '''
          dataset = np.concatenate((X, Y), axis=1)
#
#
          self.root = self.build_tree(dataset)
```

```
#
     def make_prediction(self, x, tree):
#
         ''' function to predict new dataset '''
#
         if tree.value!=None: return tree.value
         feature_val = x[tree.feature_index]
#
         if feature_val<=tree.threshold:</pre>
             return self.make_prediction(x, tree.left)
#
         else:
             return self.make prediction(x, tree.right)
#
     def predict(self, X):
#
         ''' function to predict a single data point '''
#
         preditions = [self.make_prediction(x, self.root) for x in X]
#
         return preditions
     def mse(self,y_true, y_pred,verbose=False):
#
#
         ''' Calculate Mean Square Error'''
         res = np.mean((y_true - y_pred)**2)
#
         if verbose:
#
             print(f"MSE: {res}")
         return res
     def r2(self,y_true,y_pred,verbose=False):
         ''' Calculate r2 score for regressor, range form -inf to 1, higher
 \hookrightarrowmeans better
        111
         y_{mean} = np.mean(y_true)
         # Calculate total sum of squares
#
         SST = np.sum((y_true - y_mean) ** 2)
         # Calculate residual sum of squares
#
         SSR = np.sum((y_true - y_pred) ** 2)
#
         # Calculate R^2 score
         r2 = 1 - (SSR / SST)
         return r2
```

```
# #__
 # class DecisionTreeClassifier():
      def __init__(self, min_samples_split=2, max_depth=2):
          ''' constructor '''
#
         # initialize the root of the tree
         self.root = None
         # stopping conditions
#
#
         self.min\_samples\_split = min\_samples\_split
         self.max\_depth = max\_depth
#
      def build_tree(self, dataset, curr_depth=0):
#
          ''' recursive function to build the tree '''
#
         X, Y = dataset[:,:-1], dataset[:,-1]
#
#
         num_samples, num_features = np.shape(X)
#
         # split until stopping conditions are met
#
         if num samples >= self.min samples split and curr depth <= self.max depth:
#
              # find the best split
             best_split = self.get_best_split(dataset, num_samples,_
 ⇔num_features)
#
              # check if information gain is positive
#
              if best_split["info_gain"]>0:
#
                 # recur left
                 left_subtree = self.build_tree(best_split["dataset_left"],__
 \hookrightarrow curr_depth+1)
#
                  # recur right
                 right_subtree = self.build_tree(best_split["dataset_right"],___
 \hookrightarrow curr_depth+1)
                  # return decision node
                  return Node(best_split["feature_index"],_
 ⇔best split["threshold"],
                              left_subtree,_
 →right_subtree, info_gain=best_split["info_gain"])
         # compute leaf node
         leaf_value = self.calculate_leaf_value(Y)
         # return leaf node
#
#
         return Node(value=leaf_value)
      def get_best_split(self, dataset, num_samples, num_features):
#
#
          ''' function to find the best split '''
```

```
#
          # dictionary to store the best split
#
          best\_split = \{\}
#
          max\_info\_qain = -float("inf")
          # loop over all the features
#
#
          for feature_index in range(num_features):
              feature values = dataset[:, feature index]
#
#
              possible_thresholds = np.unique(feature_values)
              # loop over all the feature values present in the data
#
              for threshold in possible_thresholds:
#
                  # get current split
                  dataset_left, dataset_right = self.split(dataset,__
 → feature_index, threshold)
#
                  # check if childs are not null
#
                  if len(dataset left)>0 and len(dataset right)>0:
                      y, left_y, right_y = dataset[:, -1], dataset_left[:, -1],
 → dataset_right[:, -1]
#
                      # compute information gain
#
                      curr_info_gain = self.information_gain(y, left_y,_
 →right_y, "gini")
#
                      # update the best split if needed
#
                      if curr_info_gain>max_info_gain:
#
                           best split["feature index"] = feature index
                           best split["threshold"] = threshold
#
                           best split["dataset left"] = dataset left
#
#
                          best_split["dataset_right"] = dataset_right
                          best_split["info_gain"] = curr_info_gain
#
#
                          max\_info\_gain = curr\_info\_gain
#
          # return best split
#
          return best_split
      def split(self, dataset, feature_index, threshold):
#
#
          ''' function to split the data '''
          dataset_left = np.array([row for row in dataset if_
 →row[feature_index]<=threshold])
          dataset_right = np.array([row for row in dataset if_
 →row[feature_index]>threshold])
          return dataset_left, dataset_right
      def information_gain(self, parent, l_child, r_child, mode="entropy"):
#
          ''' function to compute information gain '''
#
          weight_l = len(l_child) / len(parent)
          weight r = len(r child) / len(parent)
#
          if mode=="gini":
```

```
qain = self.qini_index(parent) - (weight_l*self.
 ⇒qini_index(l_child) + weight_r*self.qini_index(r_child))
          else:
              gain = self.entropy(parent) - (weight_l*self.entropy(l_child) +__
 \neg weight\_r*self.entropy(r\_child))
          return gain
      def entropy(self, y):
#
#
          ''' function to compute entropy '''
#
          class\_labels = np.unique(y)
          entropy = 0
#
          for cls in class_labels:
#
              p_cls = len(y[y == cls]) / len(y)
              entropy += -p_cls * np.log2(p_cls)
#
          return entropy
      def gini_index(self, y):
#
          ''' function to compute gini index '''
#
#
          class_labels = np.unique(y)
#
          gini = 0
#
          for cls in class_labels:
#
              p_{cls} = len(y[y == cls]) / len(y)
#
             qini += p_cls**2
#
          return 1 - qini
#
      def calculate_leaf_value(self, Y):
#
          ''' function to compute leaf node '''
          Y = list(Y)
#
          # return the maximum count element
#
          return max(Y, key=Y.count)
#
      def print_tree(self, tree=None, indent=" "):
#
          ''' function to print the tree '''
#
          if not tree:
#
              tree = self.root
#
          if tree.value is not None:
              print(tree.value)
          else:
#
              print("X_"+str(tree.feature_index), "<=", tree.threshold, "?",__</pre>
 \rightarrow tree. info_gain)
              print("%sleft:" % (indent), end="")
```

```
#
                                         self.print_tree(tree.left, indent + indent)
#
                                         print("%sright:" % (indent), end="")
#
                                         self.print_tree(tree.right, indent + indent)
                 def pretty_print(self,node=None,depth=0,prefix="Root: "):
#
#
                              '''Print tree in pretty format using post order traversal'''
#
                             if node is None:
                                         node = self.root
                             indent = " " * depth * 4
#
                             # print leaves node
#
                             if node.value is not None:
                                         print(f"{indent}{prefix}Leaf => Value: {node.value} / depth:
   \hookrightarrow \{depth\}")
                             # print feature node
#
                             else:
                                         self.pretty_print(node.right, depth + 1, prefix="R--> ")
                                       print(f''\{indent\}\{prefix\} X_{node.feature\_index\} \le \{node.feature\_index\} \le \{node.feature\_i
   ⇔threshold} / depth: {depth}")
                                         self.pretty_print(node.left, depth + 1, prefix="L--> ")
#
                 def fit(self, X, Y):
#
                              ''' function to train the tree '''
                             dataset = np.concatenate((X, Y), axis=1)
#
                             self.root = self.build_tree(dataset)
#
                 def predict(self, X):
#
                              ''' function to predict new dataset
                             Returns:
                                         List: list contains predictions
                             preditions = [self.make_prediction(x, self.root) for x in X]
#
                             return preditions
#
                 def make_prediction(self, x, tree):
#
                              ''' function to predict a single data point '''
                             # if the node is leaf node, return the value
                             if tree.value!=None: return tree.value
#
                             # otherwise travel to left or right node by the threshold
#
#
                             feature_val = x[tree.feature_index]
#
                             if feature val<=tree.threshold:</pre>
```

```
#
                 return self.make_prediction(x, tree.left)
    #
             else:
    #
                return self.make_prediction(x, tree.right)
    #
         def accuracy(self, y_true, y_pred):
             ''' Calculate the accuracy '''
    #
             count = 0
    #
             N = len(y true)
             for i in range(N):
                 if y_true[i] == y_pred[i]:
                    count += 1
    #
             return count / N
    # #__
     	ilde{ullet}
[]: from DST import DecisionTreeClassifier
[]: dt = DecisionTreeClassifier(min_samples_split=5,max_depth=10)
    dt.fit(train_X.values,train_y.values.reshape(-1,1))
```

4.2.1 Training set

```
[]: pred_y = dt.predict(train_X.values)
```

```
[ ]: np.mean(pred_y == train_y)
```

[]: 0.8666052604646403

4.2.2 test set

```
[]: pred_y = dt.predict(test_X.values)
np.mean(pred_y == test_y)
```

[]: 0.8648272474209923

4.2.3 important features

```
[]: dt.pretty_print()
```

```
R--> Leaf => Value: e | depth: 4
                         L--> X_72 \le 0.0 \mid depth: 3
                                         R--> Leaf => Value: p | depth: 5
                                 L--> X_64 \le 0.0 \mid depth: 4
                                                                  R--> Leaf =>
Value: e | depth: 8
                                                          R--> X_37 <= 0.0
depth: 7
R--> Leaf => Value: p | depth: 10
                                                                           R-->
X_90 \le 0.0 \mid depth: 9
L--> Leaf => Value: e | depth: 10
                                                                  L--> X_0 <=
10.06 | depth: 8
        R--> Leaf => Value: p | depth: 11
R--> X_33 \le 0.0 \mid depth: 10
        L--> Leaf => Value: e | depth: 11
                                                                           L-->
X_2 \le 9.26 \mid depth: 9
L--> Leaf => Value: p | depth: 10
                                                  R--> X_59 <= 0.0 \mid depth: 6
                                                                  R--> Leaf =>
Value: p | depth: 8
                                                          L--> X_87 <= 0.0
depth: 7
                                                                           R-->
Leaf => Value: p | depth: 9
                                                                  L--> X_24 <=
0.0 | depth: 8
        R--> Leaf => Value: p | depth: 11
R--> X_74 \le 0.0 \mid depth: 10
        L--> Leaf => Value: e | depth: 11
                                                                           L-->
X_0 \le 3.46 \mid depth: 9
        R--> Leaf => Value: p | depth: 11
L--> X_32 <= 0.0 \mid depth: 10
        L--> Leaf => Value: e | depth: 11
                                         L--> X_18 \le 0.0 \mid depth: 5
                                                                  R--> Leaf =>
Value: e | depth: 8
                                                          R--> X_56 <= 0.0
depth: 7
                                                                  L--> Leaf =>
Value: p | depth: 8
                                                 L--> X_30 <= 0.0 \mid depth: 6
R--> Leaf => Value: e | depth: 10
                                                                           R-->
X_{11} \le 0.0 \mid depth: 9
```

```
L--> Leaf => Value: p | depth: 10
                                                                  R--> X_108 <=
0.0 | depth: 8
                                                                           L-->
Leaf => Value: e | depth: 9
                                                          L--> X_46 <= 0.0
depth: 7
                                                                           R-->
Leaf => Value: p | depth: 9
                                                                  L--> X_83 <=
0.0 | depth: 8
R--> Leaf => Value: p | depth: 10
                                                                           L-->
X_{62} \le 0.0 \mid depth: 9
        R--> Leaf => Value: e | depth: 11
L--> X_41 <= 0.0 \mid depth: 10
        L--> Leaf => Value: e | depth: 11
Root: X_2 \le 8.22 \mid depth: 0
                                 R--> Leaf => Value: p | depth: 4
                         R--> X_17 <= 0.0 \mid depth: 3
                                         R--> Leaf => Value: p | depth: 5
                                 L--> X_11 <= 0.0 \mid depth: 4
                                                 R--> Leaf => Value: e | depth: 6
                                         L--> X_2 <= 1.79 \mid depth: 5
                                                 L--> Leaf => Value: p | depth: 6
                R--> X_1 <= 3.69 \mid depth: 2
                                 R--> Leaf => Value: e | depth: 4
                         L--> X_2 \le 3.01 \mid depth: 3
                                         R--> Leaf => Value: e | depth: 5
                                 L--> X_3 <= 0.0 \mid depth: 4
                                                  R--> Leaf => Value: e | depth: 6
                                         L--> X_5 <= 0.0 \mid depth: 5
                                                          R--> Leaf => Value: e |
depth: 7
                                                  L--> X_21 <= 0.0 \mid depth: 6
                                                          L--> Leaf => Value: p |
depth: 7
        L--> X_46 <= 0.0 \mid depth: 1
                                         R--> Leaf => Value: e | depth: 5
                                 R--> X_17 \le 0.0 \mid depth: 4
                                                  R--> Leaf => Value: e | depth: 6
                                         L--> X_10 <= 0.0 \mid depth: 5
                                                  L--> Leaf => Value: p | depth: 6
                         R--> X_117 <= 0.0 \mid depth: 3
                                                  R--> Leaf => Value: p | depth: 6
                                         R--> X_4 <= 0.0 \mid depth: 5
                                                          R--> Leaf => Value: p |
depth: 7
```

```
L--> X_65 <= 0.0 \mid depth: 6
R--> Leaf => Value: e | depth: 10
                                                                           R-->
X_21 \le 0.0 \mid depth: 9
        R--> Leaf => Value: e | depth: 11
L--> X_17 \le 0.0 \mid depth: 10
        L--> Leaf => Value: p | depth: 11
                                                                  R--> X_44 <=
0.0 | depth: 8
R--> Leaf => Value: p | depth: 10
                                                                           L-->
X_37 \le 0.0 \mid depth: 9
L--> Leaf => Value: e | depth: 10
                                                          L--> X_58 <= 0.0
depth: 7
        R--> Leaf => Value: e | depth: 11
R--> X_0 \le 2.84 \mid depth: 10
        L--> Leaf => Value: p | depth: 11
                                                                           R-->
X_55 \le 0.0 \mid depth: 9
        R--> Leaf => Value: p | depth: 11
L--> X_70 \le 0.0 \mid depth: 10
        L--> Leaf => Value: e | depth: 11
                                                                  L--> X_2 <=
0.93 | depth: 8
                                                                          L-->
Leaf => Value: p | depth: 9
                                 L--> X_97 <= 0.0 \mid depth: 4
                                                 R--> Leaf => Value: e | depth: 6
                                         L--> X_1 <= 7.23 \mid depth: 5
                                                 L--> Leaf => Value: p | depth: 6
                L--> X_86 <= 0.0 \mid depth: 2
                                         R--> Leaf => Value: e | depth: 5
                                 R--> X_13 \le 0.0 \mid depth: 4
                                         L--> Leaf => Value: p | depth: 5
                        L--> X_4 <= 0.0 \mid depth: 3
                                                 R--> Leaf => Value: e | depth: 6
                                         R--> X_106 \le 0.0 \mid depth: 5
                                                                  R--> Leaf =>
Value: e | depth: 8
                                                          R--> X_11 <= 0.0
depth: 7
                                                                           R-->
Leaf => Value: e | depth: 9
                                                                  L--> X_9 <= 0.0
| depth: 8
                                                                           L-->
Leaf => Value: p | depth: 9
```

```
L--> X_55 <= 0.0 \mid depth: 6
R--> Leaf => Value: e | depth: 10
                                                                           R-->
X_43 \le 0.0 \mid depth: 9
L--> Leaf => Value: p | depth: 10
                                                                   R--> X_58 <=
0.0 | depth: 8
R--> Leaf => Value: e | depth: 10
                                                                           L-->
X_34 \le 0.0 \mid depth: 9
L--> Leaf => Value: p | depth: 10
                                                          L--> X_10 <= 0.0
depth: 7
        R--> Leaf => Value: p | depth: 11
R--> X_8 <= 0.0 \mid depth: 10
        L--> Leaf => Value: e | depth: 11
                                                                           R-->
X_1 \le 4.08 \mid depth: 9
        R--> Leaf => Value: e | depth: 11
L--> X_{117} \le 0.0 \mid depth: 10
        L--> Leaf => Value: p | depth: 11
                                                                  L--> X 54 <=
0.0 | depth: 8
        R--> Leaf => Value: p | depth: 11
R--> X_15 \le 0.0 \mid depth: 10
        L--> Leaf => Value: e | depth: 11
                                                                           L-->
X_{49} \le 0.0 \mid depth: 9
        R--> Leaf => Value: e | depth: 11
L--> X_{22} \le 0.0 \mid depth: 10
        L--> Leaf => Value: p | depth: 11
                                 L--> X_2 <= 5.24 \mid depth: 4
                                                                  R--> Leaf =>
Value: e | depth: 8
                                                          R--> X_9 <= 0.0
depth: 7
                                                                  L--> Leaf =>
Value: p | depth: 8
                                                  R--> X_73 <= 0.0 \mid depth: 6
                                                                           R-->
Leaf => Value: e | depth: 9
                                                                   R--> X_43 <=
0.0 | depth: 8
R--> Leaf => Value: p | depth: 10
                                                                           L-->
X_1 \le 2.95 \mid depth: 9
L--> Leaf => Value: e | depth: 10
                                                          L--> X_14 <= 0.0
```

```
depth: 7
R--> Leaf => Value: e | depth: 10
                                                                           R-->
X_{10} \le 0.0 \mid depth: 9
L--> Leaf => Value: p | depth: 10
                                                                   L--> X_8 <= 0.0
| depth: 8
                                                                           L-->
Leaf => Value: p | depth: 9
                                         L--> X_34 <= 0.0 \mid depth: 5
                                                                   R--> Leaf =>
Value: e | depth: 8
                                                          R--> X_11 <= 0.0
depth: 7
                                                                   L--> Leaf =>
Value: p | depth: 8
                                                  L--> X_0 <= 1.4 \mid depth: 6
                                                          L--> Leaf => Value: e |
depth: 7
```

The first 3 split is 2,46,68. Since decision tree will find the use best split in each decision node, the most important features will be in the first coupes splits.

```
[]: X.iloc[:,[2,26,68]].columns
```

[]: Index(['stem-width', 'cap-color_l', 'stem-surface_g'], dtype='object')

'stem-width', 'cap-color_l', 'stem-surface_g' are the important features.

4.3 KNN

```
# distances = np.linalg.norm(self.X_train - x, axis=1)

# Find the indices of the k nearest neighbors

# k_index = np.argpartition(distances, self.k)[:self.k]

# Retrieve the labels of the k nearest neighbors

# k_nearest_labels = self.y_train[k_index]

# return list(Counter(k_nearest_labels).keys())[0]
```

```
from pca import PCA
from KNN import KNN
pca = PCA(n_components=2)
pca.fit(X)

X_transform = pca.transform(X)
X_transform = pd.DataFrame(X_transform,columns=['PC1',"PC2"])
train_X,test_X,train_y,test_y = train_test_split(X_transform,y,test_size=0.
$\infty 2$, random_state=42)
```

4.3.1 Training set

```
[]: knn = KNN(k=247)
knn.fit(train_X.values,train_y.values)
pred_y = knn.predict(train_X.values)
```

```
[]: print(f'accuracy: {np.mean(train_y == pred_y)}')
```

accuracy: 0.6252379490328523

4.3.2 Test set

```
[]: pred_y = knn.predict(test_X.values)
print(f'accuracy: {np.mean(test_y == pred_y)}')
```

accuracy: 0.6240379891927297

KNN perform worse than logistic regression and Decision Tree. I choose coupe k values: 7,11,21,41 they perform similarly as k=247. The best accuracy i got from KNN is 0.71 which is still worse than other models.

4.3.3 important feature

KNN cannot find the most important features

5 Model selection

• Logistic Regression:

run time: 1minTrain: 0.815Test: 0.811

• Decision Tree:

run time: 3hoursTrain: 0.866Test: 0.864

• KNN:

run time: 1minTrain: 0.625Test: 0.624

The model I will choose is Decision Tree, It has the best performance compare with other two models, but it takes long time to train the model. In addition, Decision Tree can find the important features easily. I think I can optimize the run time using some data structure and using vectorization by numpy.