## **Data Visualization Lab 2**

Eda Haberman dataset

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# **Exploratory Data Analysis (EDA)**

load the data and perform some initial analysis. Assuming you have the dataset in a CSV file:

```
In [2]:
```

```
# Import necessary libraries
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
# Load the dataset
data = pd.read csv('haberman.csv')
# Display the first few rows of the dataset
print(data.head())
# Check for missing values
print(data.isnull().sum())
# Summary statistics
print(data.describe())
# Data types and counts
print(data.info())
   age year nodes status
0
   30 64 1 1
   30
           62
                     3
1
                               1
                              1
   30 65 0
2
3 31 59
                   2
                              1
4 31 65
                    4
       0
age
          0
year
nodes
status
          Ω
dtype: int64
                age
                             year
                                           nodes status
count 305.000000 305.000000 305.000000 305.000000
                        3.242783 7.200501
58.000
mean 52.357377 62.868852
                                                    1.262295
        10.678010
                                                     0.440605
std

      10.6/8010
      3.242783
      7.200528
      0.440605

      30.000000
      58.000000
      0.000000
      1.000000

      44.000000
      60.000000
      0.000000
      1.000000

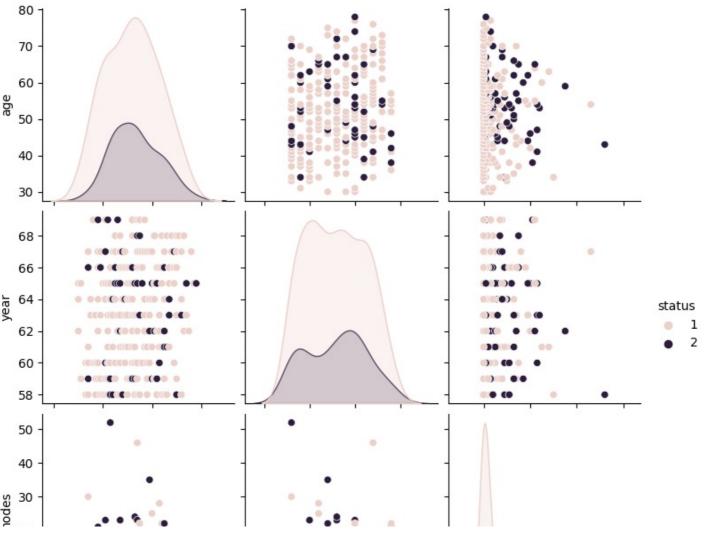
      52.000000
      66.000000
      4.000000
      2.000000

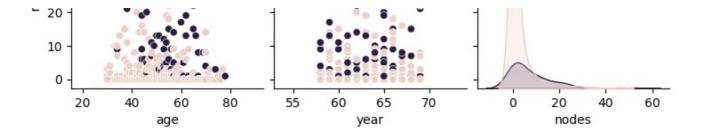
      78.000000
      69.00000
      52.000000
      2.000000

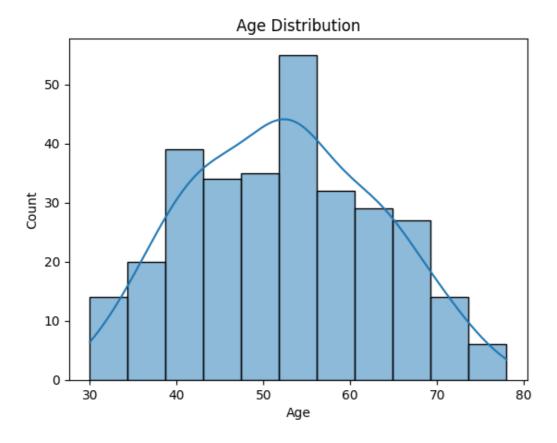
        30.000000 58.000000
min
25%
50%
75%
max
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 305 entries, 0 to 304
Data columns (total 4 columns):
 # Column Non-Null Count Dtype
--- ----- -----
 0 age 305 non-null int64
1 year 305 non-null int64
 2 nodes 305 non-null int64
3 status 305 non-null int64
dtypes: int64(4)
memory usage: 9.7 KB
None
```

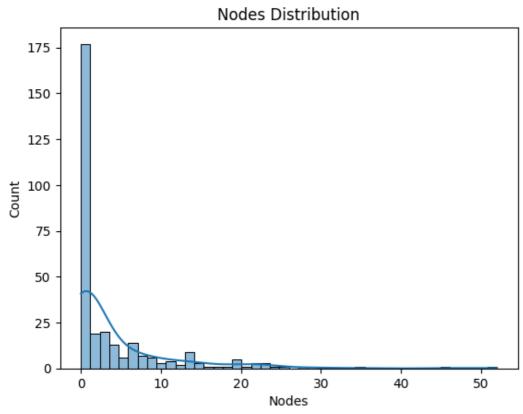
### **EDA**

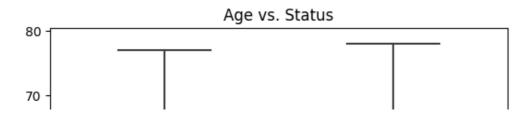
```
In [3]:
# Pairplot to visualize relationships between numerical attributes
sns.pairplot(data, hue='status')
plt.show()
# Histograms to visualize the distribution of individual features
sns.histplot(data['age'], kde=True)
plt.xlabel('Age')
plt.title('Age Distribution')
plt.show()
sns.histplot(data['nodes'], kde=True)
plt.xlabel('Nodes')
plt.title('Nodes Distribution')
plt.show()
# Box plots to identify outliers
sns.boxplot(x='status', y='age', data=data)
plt.xlabel('Status')
plt.ylabel('Age')
plt.title('Age vs. Status')
plt.show()
sns.boxplot(x='status', y='nodes', data=data)
plt.xlabel('Status')
plt.ylabel('Nodes')
plt.title('Nodes vs. Status')
plt.show()
C:\Users\utkar\AppData\Local\Packages\PythonSoftwareFoundation.Python.3.11 qbz5n2kfra8p0\
LocalCache\local-packages\Python311\site-packages\seaborn\axisgrid.py:118: UserWarning: T
he figure layout has changed to tight
  self. figure.tight_layout(*args, **kwargs)
   80
   70
   60
```

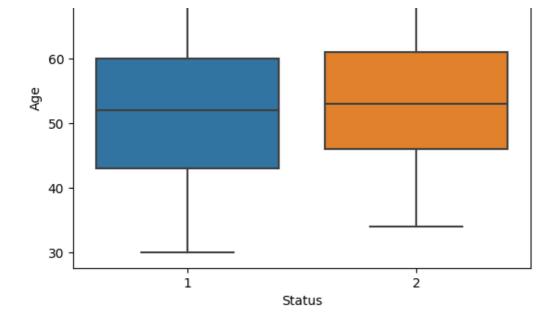


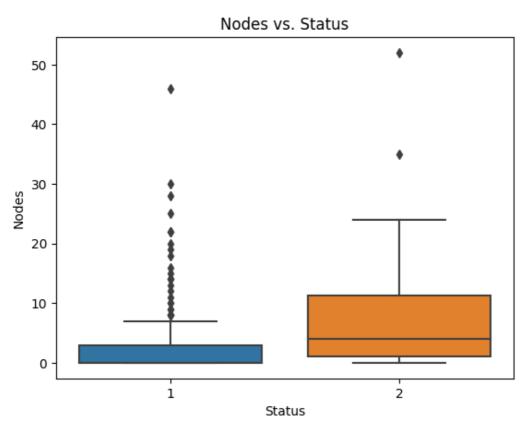












# **Basic Statistical Analysis**

#### In [4]:

```
# Calculate mean, median, and standard deviation for age and nodes
mean_age = data['age'].mean()
median_age = data['age'].std()

mean_nodes = data['nodes'].mean()
median_nodes = data['nodes'].median()
std_nodes = data['nodes'].std()

print(f"Age: Mean={mean_age}, Median={median_age}, Standard Deviation={std_age}")
print(f"Nodes: Mean={mean_nodes}, Median={median_nodes}, Standard Deviation={std_nodes}")
```

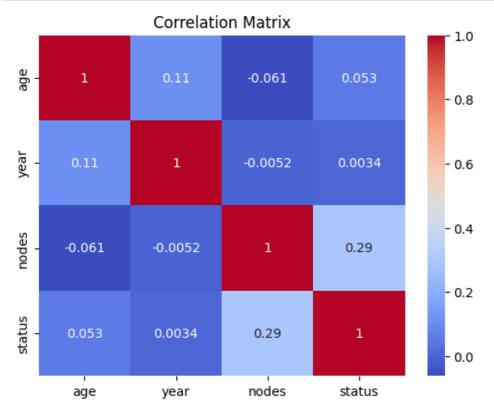
Age: Mean=52.35737704918033, Median=52.0, Standard Deviation=10.678009544589289 Nodes: Mean=4.032786885245901, Median=1.0, Standard Deviation=7.200528153862346

## **Multivariate Analysis**

#### **Correlation Analysis**

#### In [5]:

```
# Correlation matrix
correlation_matrix = data.corr()
sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm')
plt.title('Correlation Matrix')
plt.show()
```



### **Logistic Regression (Predictive Analysis):**

#### In [8]:

```
from sklearn.model selection import train test split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score, classification_report
# Define your features (X) and target variable (y)
X = data[['age', 'year', 'nodes']]
y = data['status']
# Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y, test size=0.2, random state=42
# Create a logistic regression model
model = LogisticRegression()
model.fit(X_train, y_train)
# Predict on the test set
y pred = model.predict(X test)
# Calculate accuracy and print classification report
accuracy = accuracy score(y test, y pred)
print(f"Accuracy: {accuracy}")
print(classification_report(y_test, y_pred))
# Assuming you have already trained a logistic regression model
# Model parameters (intercept and coefficients)
intercept = model.intercept
coefficients = model.coef_[0]
```

```
# Create a scatter plot of the data points
plt.scatter(X_test['age'], X_test['nodes'], c=y_test, cmap='RdYlBu', marker='o', label='
Data Points')

# Add labels and legend
plt.xlabel('Age')
plt.ylabel('Nodes')
plt.title('Logistic Regression Decision Boundary')
plt.legend(loc='best')

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

Accuracy: 0.6721311475409836

	precision	recall	f1-score	support
1	0.68	0.95	0.80	41
2	0.50	0.10	0.17	20
accuracy			0.67	61
macro avg	0.59	0.53	0.48	61
weighted avg	0.62	0.67	0.59	61

# Logistic Regression Decision Boundary

