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Haberman's survival data set:
            The data set, contains cases from a study, that was conducted between 1958 and 1970, at the
            University of Chicago Billings Hospital, on the survival of patients, who had undergone surgery for
            breast cancer.
            Objective:
            To classify whether a patient survived 5 year or longer or the patient died within 5 years
            No. of instances:
            306
            No. of attribute:
            4(including class atribute)
            Attribute information:
                    a.Age of patient at the time of operation(numerical)
                    b.patient's year of operation(year-1900, numerical)
                    c.no. of positive axillary nodes detected(numerical)
                    d.survival status (class attribute)1=the patient survived 5 years o
                r longer 2= the patient died within 5 years
   In [2]: import pandas as pd
            import seaborn as sns
            import numpy as np
            import matplotlib.pyplot as plt
            #loading haberman.csv into panda dataframe
            haber= pd.read_csv("haberman.csv")
   In [3]: import os
            os.getcwd()
   Out[3]: '/Users/user/Downloads'
   In [4]: os.chdir("/Users/user/Downloads")
   In [5]: import warnings
            warnings.filterwarnings('ignore')
   In [5]: #finding no. of datapoints and features
            print(haber.shape)
            (306, 4)
   In [6]: #finding the column names in our dataset
            print(haber.columns)
            Index([u'age', u'year', u'nodes', u'status'], dtype='object')
   In [7]: #to find the survival status(class attribute)
            haber["status"].value_counts()
                 225
            2
                  81
            Name: status, dtype: int64
            this shows 1) 225 patients survived 5 years or longer 2)81 patients died within 5 years
            Bi-variate analysis
            2-D Scatter plot:
            hard to classify with only one colour.
   In [3]: haber.plot(kind='scatter', x='age', y='nodes');
            plt.title("2-D scatter plot")
            plt.show()
                                  2-D scatter plot
               50
               40
               30
               20
               10
                                                 70
            2-D scatter with differnt colors for different survival
            status:
   In [6]: sns.FacetGrid(haber, hue="status", size=3).map(plt.scatter, "age", "nodes").add_le
            plt.title("2-D scatter with different colors for different survival status:")
            plt.show()
             2-D scatter with differnt colors for different survival status:
                        50
                        40
                      30 20
                                                        status
                                                        1
from the above figure it is hard to classify status 1 and status2, as they overlap at many places.
            2-D pair plot:
                1) suppose if there are more than 2 features , then pair plot is used , \ensuremath{\mathbf{w}}
                here different pairs of features are tried.
                2)It is used to visualize the data in 2-D and it cannot be used for 3-D
                or 4-D etc.
  In [11]: sns.set style("whitegrid");a= sns.pairplot(haber, hue="status", vars=["age", "y
            ear", "nodes"], height=3);
            a.fig.suptitle("2 D pair plot")
  Out[11]: Text(0.5,0.98,'2 D pair plot')
               70
               60
               30
               68
                                                                    (0) (0)
               66
                                                                                       status
               60
                     . . . . . . . . . . . . .
               50
            Observation from bi-variate analysis:
                1) In plots of ,age and year data overlaps with each other.
                2) Nodes is a better attribute to classify.
            Univariate analysis:
            Histogram:
  In [13]: sns.FacetGrid(haber, hue='status', height=4).map(sns.distplot, "year").add legend
            plt.title("histogram for year")
  Out[13]: Text(0.5,1,'histogram')
                             histogram
             0.12
             0.10
             0.08
                                                  status
             0.06
             0.04
             0.02
             0.00
            Observation:
            the data is overlapping so it is hard to classify
  In [14]: sns.FacetGrid(haber, hue='status', height=4).map(sns.distplot, "age").add legend
            plt.title("histogram for age");
                           histogram for age
             0.040
             0.035
             0.030
             0.025
                                                  status
             0.020
                                                  2
             0.015
             0.010
             0.005
             0.000
                                 60
            Observation:
                1) Patients within age 30-34 have survived more than five years after op
                2) Patients within age 77-84 have not survived more than 5 years after o
                3) The rest of the data is overlapping.
  In [15]: sns.FacetGrid(haber, hue='status', height=8).map(sns.distplot, "nodes").add legen
            plt.title("histogram for nodes");
                                            histogram for nodes
             0.5
             0.4
                                                                                       status
             0.2
             0.1
            Observations:
                1) As the no. of nodes increases, survival status decreases.
                2) The rest of the data is overlapping.
            Box plot and whiskers:
  In [16]: sns.boxplot(x='status',y='year',data=haber)
            plt.title("box plot and whiskers for year")
  Out[16]: Text(0.5,1,'box plot and whiskers for year')
                             box plot and whiskers for year
               68
               66
                            1
                                      status
  In [17]: sns.boxplot(x='status', y='age', data=haber)
            plt.title("box plot and whiskers for age")
  Out[17]: Text(0.5,1,'box plot and whiskers for age')
                             box plot and whiskers for age
               80
               70
               60
               50
               40
            Observation:
                1) Patients within age 30-34 have survived more than five years after op
                2) Patients within age 77-83 have not survived more than five years afte
                r operation.
  In [20]: sns.boxplot(x='status', y='nodes', data=haber)
            plt.title("box plot and whiskers for nodes")
  Out[20]: Text(0.5,1,'box plot and whiskers for nodes')
                            box plot and whiskers for nodes
               50
                            ٠
               40
               30
               20
                                      status
            Observation:
                1) Patients with 0 nodes have higher chance of survival
            Violin plots:
  In [21]: sns.violinplot(x='status', y='year', data=haber)
            plt.title("violin plots for year")
  Out[21]: Text(0.5,1,'violin plots for year')
                                  violin plots for year
               72.5
               70.0
               67.5
               65.0
               62.5
               60.0
               55.0
            Observation:
                1) Hard to classify as the data is overlapping.
  In [22]: sns.violinplot(x='status', y='age', data=haber)
            plt.title("violin plot for age")
  Out[22]: Text(0.5,1,'violin plot for age')
```

20 1 2 status

In [23]: sns.violinplot(x='status',y='nodes',data=haber)
 plt.title("violin plot for nodes")

1) Hard to classify as the data is overlapping.

violin plot for age

90

80

70

50

**Observation:** 

PDF:
In [24]: counts, bin\_edges=np.histogram(haber['nodes'], bins=10, density=True)
pdf=counts/(sum(counts))
cdf=np.cumsum(pdf)
plt.plot(bin\_edges[1:], pdf)
plt.plot(bin\_edges[1:], cdf)
counts, bin\_edges=np.histogram(haber['nodes'], bins=10, density=True)

plt.plot(bin\_edges[1:],pdf,color="dodgerblue")
plt.plot(bin\_edges[1:],cdf,color="orange")

PDF and CDF

Pdf for the patients who survive more than 5 years

Cdf for the patients who survive more than 5 years

plt.legend(['Pdf for the patients who survive more than 5 years',

'Cdf for the patients who survive more than 5 years'])

pdf=counts/(sum(counts))

plt.title("PDF and CDF ")

not survive is 54

plt.xlabel("nodes")

plt.show()

cdf=np.cumsum(pdf)

## 0.6

1.0

0.2 0.0 50 20 nodes Mean In [20]: less\_than\_five=haber[haber['status']==2] more\_than\_five=haber[haber['status']==1] In [20]: print(np.mean(more\_than\_five)) 52.017778 age 62.862222 year 2.791111 status 1.000000 dtype: float64 In [21]: print(np.mean(less\_than\_five)) 53.679012 age year 62.827160 nodes 7.456790 2.000000 status dtype: float64 **Observation from univariate analysis:** 1) Mean age of patients who survived more than 5 years is 52 and who did

2) Patients having nodes less than 3 nodes have survived more than 5 yea