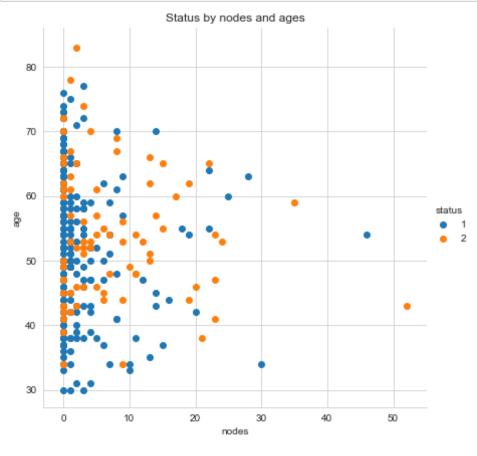
```
In [1]: import pandas as pd
         import numpy as np
         import seaborn as sns
         import matplotlib.pyplot as plt
         import warnings
         warnings.filterwarnings("ignore")
In [2]: df=pd.read csv("haberman.csv")
In [3]: df.shape
         # Haberman Dataset has 306 Observations and 4 Columns
Out[3]: (306, 4)
In [4]: df.info()
         df.dtypes
         <class 'pandas.core.frame.DataFrame'>
         RangeIndex: 306 entries, 0 to 305
         Data columns (total 4 columns):
                   306 non-null int64
         age
         year
                   306 non-null int64
                   306 non-null int64
         nodes
                   306 non-null int64
         status
         dtypes: int64(4)
         memory usage: 9.6 KB
Out[4]: age
                   int64
                   int64
         year
                   int64
         nodes
         status
                   int64
         dtype: object
In [11]: #we can use above statement to store all the coloumns in a list format
         col=df.columns.tolist()
```

```
df.columns
Out[11]: Index(['age', 'year', 'nodes', 'status'], dtype='object')
In [12]: #Here when we see that status 73% BELONGS to 1 and 26% belongs to 2, so
           this implies it is not a balanced dataset
          df['status'].value counts()
Out[12]: 1
               225
                81
          Name: status, dtype: int64
In [13]: #We dont have any missing values in the records
          #we also see from the above statistical table mean and median(50%) are
           almost equailavent to each other except for nodes
          df.describe()
Out[13]:
                      age
                               year
                                        nodes
                                                  status
           count 306.000000 306.000000 306.000000 306.000000
                 52.457516
                           62.852941
                                      4.026144
                                                1.264706
           mean
                 10.803452
                            3.249405
                                      7.189654
                                                0.441899
            min
                 30.000000
                           58.000000
                                      0.000000
                                               1.000000
            25% 44.000000
                           60.000000
                                      0.000000
                                               1.000000
            50% 52.000000
                           63.000000
                                     1.000000
                                               1.000000
            75%
                 60.750000
                           65.750000
                                      4.000000
                                                2.000000
                 83.000000 69.000000
                                     52.000000
                                                2.000000
 In [6]: # 2d scatter plot
          #2d Scatter plot with respect to ages and nodes.
          sns.set style("whitegrid");
          sns.FacetGrid(df,hue="status",height=6)\
              .map(plt.scatter, "nodes", "age")\
```

```
.add_legend()
plt.title('Status by nodes and ages')

plt.show()
#0bservations
#1,we have used scatter plot belwo using seaborn but it is really confusing to understand
# as all the data scattered randomly'
#2,But from my obervations i see that most of the dots are visible at
0 nodes and i also see that most of them belong tp 'status'=1'''
```



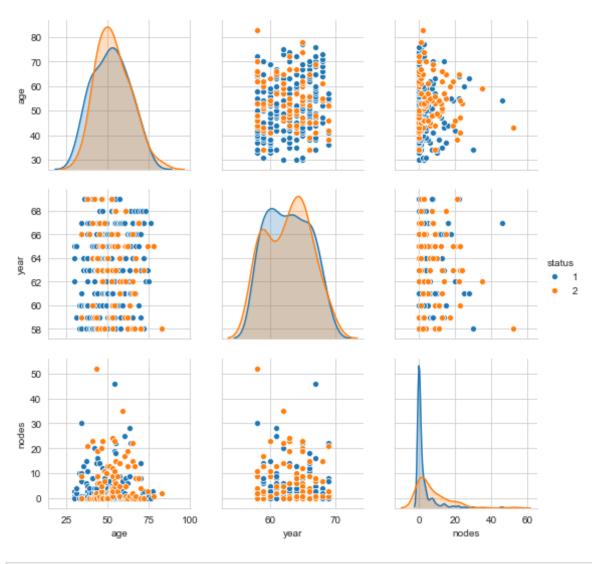
In [15]: # from this below line code we see that 38 % of node belonging to 0 and 10 % of nodes belonging are in 1(Status) and all the other points are

```
dfagg2=pd.crosstab(df['status'], df['nodes'],normalize=True)
# when we do the same thing for ages we see that all the points are ran
domly distibuted here and there
dfagg3=pd.crosstab(df['status'], df['age'],normalize=True)

In [10]: # Pair Plots

sns.set_style("whitegrid");
sns.pairplot(df,hue='status',vars=['age','year','nodes'])
plt.show()

#Observations
# 1,Accoring to me i am not able to make out any obervations using pair
plots
#2, I can see only one thing in the above observations that some of the
m whose age is above 65 and belonging to year 1968 are in status 1 with
linearly related
```



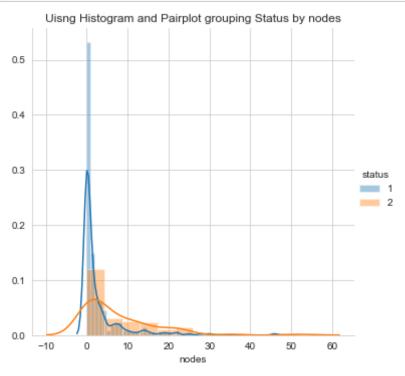
```
.add_legend();
plt.title('Uisng Histogram and Pairplot grouping Status by nodes')

plt.show()

#Observations
#1,By using pairplot chart i can make out one thing that nodes belongin
g to -10 to -0.2 are of surival status 2

#2,we can also see that we have clear representation of axil nodes belo
nging from range 48 to 62 are of status 2

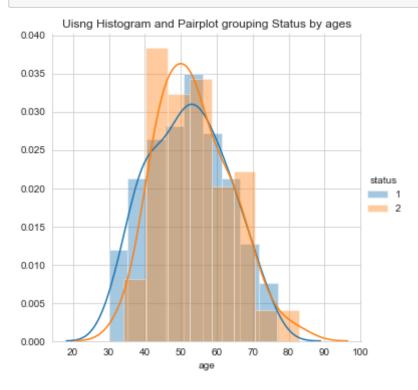
#3,Both the status 1 and 2 corresponding to nodes are overlapping each
other from -0.25
```



```
In [12]: #Histogram and Pairplot for ages
sns.FacetGrid(df,hue='status',height=5)\
    .map(sns.distplot,"age")\
```

```
.add_legend();
plt.title('Uisng Histogram and Pairplot grouping Status by ages')
plt.show()

#Observations
#1,we came across 1 obervation in this pair plot is who age is below 1
9 are in status of 1 and whose age is greater the 88 are in status 2, i
cannot find any differentiate
```

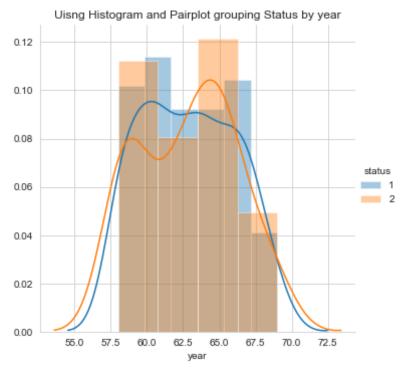


```
In [13]: #Histogram and Pairplot for year

sns.FacetGrid(df,hue='status',height=5)\
    .map(sns.distplot,"year")\
    .add_legend();
plt.title('Uisng Histogram and Pairplot grouping Status by year')
```

```
plt.show()

# Observations
#1,we can see that the year in range from 58 to 69 have both status 1 a
nd 2
#2,we can see that there are less % who are in both status 1 and status
after year 72 and in year 55
```

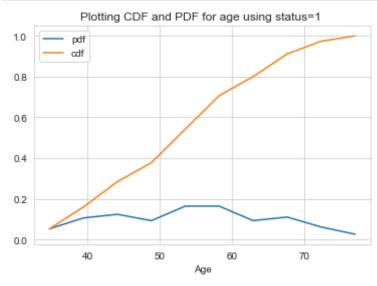


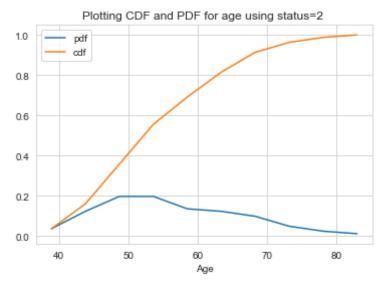
In [16]: #CDF and PDF for age
 import numpy as np
 dfstatus1=df.loc[df["status"]==1]
 ddfstatus2=df.loc[df['status']==2]

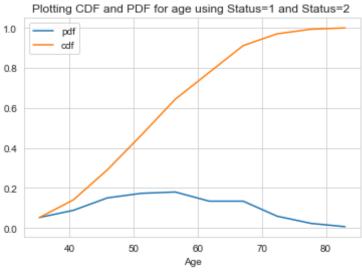
In [20]: #Computing CDF and PDF in comparision to age.
#Plotting cdf and pdf for status=1 in comparision to age

```
counts, bins edges=np.histogram(dfstatus1['age'], bins=10, density=True)
pdf=counts/(sum(counts))
pdf
bins_edges
#compute cdf for status=1
cdf=np.cumsum(pdf)
plt.plot(bins edges[1:],pdf,label='pdf')
plt.plot(bins edges[1:],cdf,label='cdf')
plt.legend()
plt.title('Plotting CDF and PDF for age using status=1 ')
plt.xlabel('Age')
plt.show()
#Plotting cdf and pdf for status=2 in comparision to age
counts,bins edges=np.histogram(ddfstatus2['age'],bins=10,density=True)
pdf=counts/(sum(counts))
pdf
bins edges
#compute cdf for status=2
cdf=np.cumsum(pdf)
plt.plot(bins edges[1:],pdf,label='pdf')
plt.plot(bins edges[1:],cdf,label='cdf')
plt.legend()
plt.title('Plotting CDF and PDF for age using status=2 ')
plt.xlabel('Age')
plt.show()
#Plotting cdf and pdf for both status=1 and status=2 in comparision to
age.
counts, bins edges=np.histogram(df['age'], bins=10, density=True)
pdf=counts/(sum(counts))
pdf
bins edges
#compute cdf for both the status
cdf=np.cumsum(pdf)
```

```
plt.plot(bins edges[1:],pdf,label='pdf')
plt.plot(bins edges[1:],cdf,label='cdf')
plt.legend()
plt.title('Plotting CDF and PDF for age using Status=1 and Status=2 ')
plt.xlabel('Age')
plt.show()
#Observation
#1,70%(159) of them belonging to age group are below 58 and their statu
s is 1
#2,57%(45) of them belonging to age group are below 53 and their status
#3, when we check for both the status 64 % of them belonging to age grou
p less then 56
#4, According to me this classification will only work when want to com
pare both status diiffferently and the results are invalid with resepec
t to ages
# as to see for two status at once, this type of analysis will not wo
rk and will give us irrelvant anaswers
```

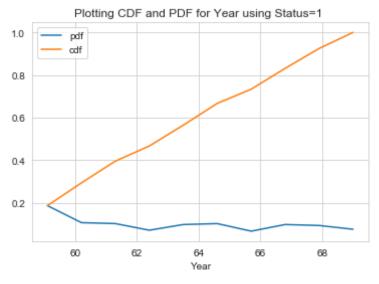


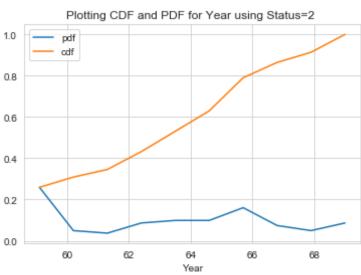


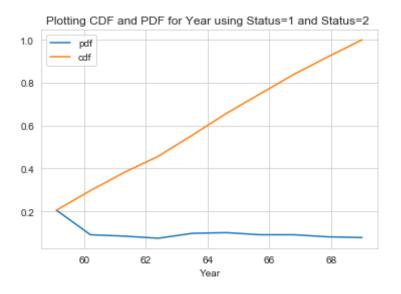


```
In [21]: #Computing CDF and pdf for Year
         #Computing pdf for status=1 with respect to Year
         counts,bins edges=np.histogram(dfstatus1['year'],bins=10,density=True)
         pdf=counts/(sum(counts))
         pdf
         bins edges
         #compute cdf for status=1
         cdf=np.cumsum(pdf)
         plt.plot(bins edges[1:],pdf,label='pdf')
         plt.plot(bins edges[1:],cdf,label='cdf')
         plt.legend()
         plt.title('Plotting CDF and PDF for Year using Status=1 ')
         plt.xlabel('Year')
         plt.show()
         #Plotting cdf and pdf for status=2 in comparision to year
         counts,bins edges=np.histogram(ddfstatus2['year'],bins=10,density=True)
         pdf=counts/(sum(counts))
         pdf
         bins_edges
```

```
#compute cdf for status=2
cdf=np.cumsum(pdf)
plt.plot(bins edges[1:],pdf,label='pdf')
plt.plot(bins edges[1:],cdf,label='cdf')
plt.legend()
plt.title('Plotting CDF and PDF for Year using Status=2 ')
plt.xlabel('Year')
plt.show()
#Plotting cdf and pdf for status=1 and status=2 in comparision to year.
counts,bins edges=np.histogram(df['year'],bins=10,density=True)
pdf=counts/(sum(counts))
pdf
bins edges
#compute cdf for both the status
cdf=np.cumsum(pdf)
plt.plot(bins edges[1:],pdf,label='pdf')
plt.plot(bins edges[1:],cdf,label='cdf')
plt.legend()
plt.title('Plotting CDF and PDF for Year using Status=1 and Status=2 ')
plt.xlabel('Year')
plt.show()
#0bservations
#1, we cannot see a correct distribution of cdf and pair plot in year i
n dependent to 1 (status) as in the year
#2, when we see the cdf and pdf for status=2 , i see that 39 of the val
ues are in distribution of year 60-65 with respect to status=2
#3, we see that we have totally 66%(204/306) of values belonging to stat
us =1 and status=2 who are in between year of 60 and 68
```





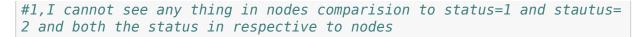


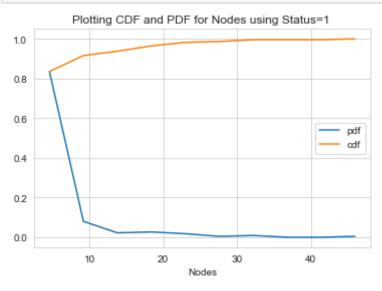
```
In [23]: #compute cdf and pdf for nodes in relation to status
#computing cdf and pdf in respect to status=1 with respect to nodes

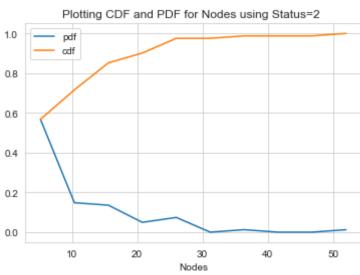
#compute pdf for status=1 in respect to nodes
counts,bins_edges=np.histogram(dfstatus1['nodes'],bins=10,density=True)
pdf=counts/(sum(counts))
pdf
bins_edges

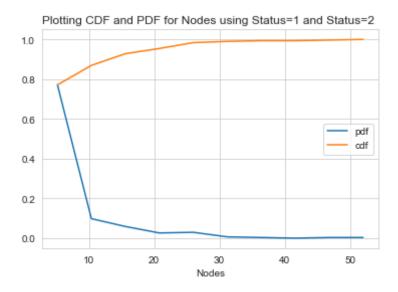
#compute cdf for status=1 in respect to nodes
cdf=np.cumsum(pdf)
plt.plot(bins_edges[1:],pdf,label='pdf')
plt.plot(bins_edges[1:],cdf,label='cdf')
plt.legend()
plt.title('Plotting CDF and PDF for Nodes using Status=1 ')
```

```
plt.xlabel('Nodes')
plt.show()
#computing cdf and pdf in respect to status=2 with respect to nodes
counts, bins edges=np.histogram(ddfstatus2['nodes'], bins=10, density=True
pdf=counts/(sum(counts))
pdf
bins edges
#compute cdf for status=2 in respect to nodes
cdf=np.cumsum(pdf)
plt.plot(bins edges[1:],pdf,label='pdf')
plt.plot(bins edges[1:],cdf,label='cdf')
plt.legend()
plt.title('Plotting CDF and PDF for Nodes using Status=2 ')
plt.xlabel('Nodes')
plt.show()
#computing cdf and pdf in respective to both status with respect to no
des
#compute pdf for nodes with respect to status=1 and status=2
counts,bins edges=np.histogram(df['nodes'],bins=10,density=True)
pdf=counts/(sum(counts))
pdf
bins edges
#compute cdf for nodes with respect to status=1 and status=2
cdf=np.cumsum(pdf)
plt.plot(bins edges[1:],pdf,label='pdf')
plt.plot(bins edges[1:],cdf,label='cdf')
plt.legend()
plt.title('Plotting CDF and PDF for Nodes using Status=1 and Status=2 '
plt.xlabel('Nodes')
plt.show()
#Observations
```









```
In [32]: #Calculating the mean and std of the haberman dataset
         print('The mean of the nodes of haberman dataset which has status 1 and
         status 2 are given below for nodes')
         print(np.mean(dfstatus1['nodes']))
                                            #calculating the mean of the
         nodes with status 1
         print(np.mean(ddfstatus2['nodes'])) #calculating the mean of th
         e nodes with status 2
         print('The standard deviation of the nodes of haberman dataset which ha
         s status 1 and status 2 are given below for nodes')
         print(np.std(dfstatus1['nodes']))
                                                   #calculating the standard de
         viation with status 1
         print(np.std(ddfstatus2['nodes']))
         #Calculating the mean and std of the haberman dataset with respect to a
         ges
         print('The mean of the nodes of haberman dataset which has status 1 and
         status 2 are given below for ages')
         print(np.mean(dfstatus1['age']))
                                                 #calculating the mean of the n
         odes with status 1
         print(np.mean(ddfstatus2['age']))
                                                 #calculating the mean of the
          nodes with status 2
```

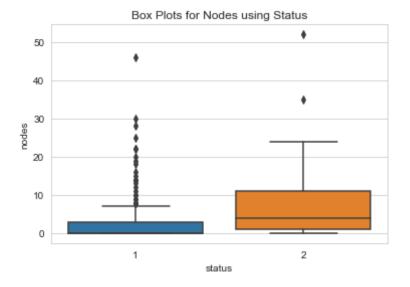
```
print('The standard deviation of the nodes of haberman dataset which ha
s status 1 and status 2 are given below for ages')
print(np.std(dfstatus1['age']))
                                       #calculating the standard devi
ation with status 1
print(np.std(ddfstatus2['age'])) #calculating the standard devi
ation with status 2
#Calculating the mean and std of the haberman dataset with respect to y
print('The mean of the nodes of haberman dataset which has status 1 and
status 2 are given below for year')
print(np.mean(dfstatus1['year'])) #calculating the mean of the
nodes with status 1
print(np.mean(ddfstatus2['year'])) #calculating the mean of the
nodes with status 2
print('The standard deviation of the nodes of haberman dataset which ha
s status 1 and status 2 are given below for year')
print(np.std(dfstatus1['year'])) #calculating the standard dev
iation with status 1
print(np.std(ddfstatus2['year']))
#conclusion:By using visual charts we are unable to find any kind of pa
tterns with respect to status 1 and status 2
#we cannot use any of the columns and say correctly that buy taking th
is column we can clearly see the difference in status
#But using mean and std we can see that only for nodes column the mean
and std proves us worthy with repspect to status 1 and 2
'''The mean of the nodes of haberman dataset which has status 1 and sta
tus 2 are given below
2.7911111111111113
7.45679012345679
The standard deviation of the nodes of haberman dataset which has statu
s 1 and status 2 are given below
5.857258449412131
9.128776076761632'''
```

The mean of the nodes of haberman dataset which has status 1 and status 2 are given below for nodes 2.791111111111113

7.45679012345679 The standard deviation of the nodes of haberman dataset which has statu s 1 and status 2 are given below for nodes 5.857258449412131 9.128776076761632 The mean of the nodes of haberman dataset which has status 1 and status 2 are given below for ages 52.017777777778 53.67901234567901 The standard deviation of the nodes of haberman dataset which has statu s 1 and status 2 are given below for ages 10.98765547510051 10.10418219303131 The mean of the nodes of haberman dataset which has status 1 and status 2 are given below for year 62.862222222222 62.82716049382716 The standard deviation of the nodes of haberman dataset which has statu s 1 and status 2 are given below for year 3.2157452144021956 3.3214236255207883 Out[32]: 'The mean of the nodes of haberman dataset which has status 1 and statu deviation of the nodes of haberman dataset which has status 1 and statu s 2 are given below\n5.857258449412131\n9.128776076761632' In [24]: # Lets only use nodes and take a look at boxplot if we are able to find anything as we are able to see a lot of #variation of mean and std for nodes for both the status #I see from below box plot that the distribution of data is not equally spread btw 25-100 for both the sttaus which can be clearly represented for both the nodes # Box Plot sns.boxplot(x='status',y='nodes',data=df)

plt.title("Box Plots for Nodes using Status")

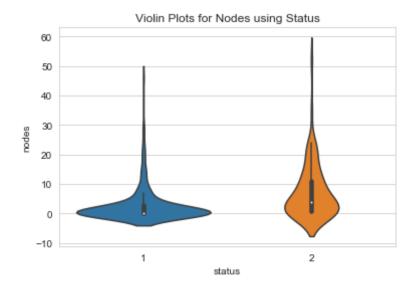
Out[24]: Text(0.5, 1.0, 'Box Plots for Nodes using Status')



```
In [25]: #violin plot for nodes in comparison to status
    sns.violinplot(x='status',y='nodes',data=df)
    plt.title("Violin Plots for Nodes using Status")

#Observations:
#1,I see no observations from below point.
```

Out[25]: Text(0.5, 1.0, 'Violin Plots for Nodes using Status')



#Conclusion #I think after doing total visual analysis of all the obervations we can say that we can use nodes column for significant observations in status as we see that we got a mean and std for both the status of nodes with a very large variance