

```
In [1]: import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np
import warnings
warnings.filterwarnings("ignore")
```

```
In [2]: data =pd.read_csv('haberman.csv')
```

Assignment to be solved

- 1) High level statistics of the dataset: number of points, number of features, number of classes, data-points per class.
- 2) Explain objective.
- 3) Perform Univariate analysis(PDF, CDF, Boxplot, Violin plots) to understand which features are useful towards classification.
- 4) Perform Bi-variate analysis (scatter plots, pair-plots) to see if combinations of features are useful in classification.
- 5) Write your observations in english as crisply and unambiguously as possible. Always quantify your results.

```
In [3]: data['Status']=np.where(data['status']==2, 'Not survived', 'survived')
```

```
In [4]: data.drop('status',axis=1,inplace=True)
```

```
In [5]: data.rename(columns = {'Status':'status'},inplace=True)
```

```
In [6]: print('Number of points : ',data.shape[0]),
print('Number of features is ',data.shape[1], ' and they are ',data.columns[0],',
print('Number of classes',len(data['status'].unique()),' they are ',data['status']
print('data-points per class','\n',data['status'].value_counts())
```

```
Number of points : 306
Number of features is 4 and they are age , year , nodes , status
Number of classes 2 they are ['survived' 'Not survived']
data-points per class
survived      225
Not survived   81
Name: status, dtype: int64
```

```
In [7]: #Data imbalance

Status_1=round(225/(225+81),2)
Status_2=round(81/(225+81),2)

print('Dataset is imbalance with the ratio',Status_1,':',Status_2)
```

Dataset is imbalance with the ratio 0.74 : 0.26

```
In [8]: #Checking for Null values
data.isnull().sum()
```

```
Out[8]: age      0
       year      0
       nodes     0
       status     0
       dtype: int64
```

Observation:

- 1) Dataset is imbalance with the ratio 0.74 : 0.26
- 2) There is zero data point with null values

Note: Survival status

- 1 = the patient survived 5 years or longer(survived)
- 2 = the patient died within 5 year (Not survived)

Objective: To predict/classify survival of patients who had undergone surgery for breast cancer. i.e whether patient survived more than 5 years or survived less than 5 years

Univarait analysis(PDF, CDF, Boxplot, Voilin plots)

```
In [9]: #Making two sub dataset

status_1=data[data['status']=='survived']
status_2=data[data['status']=='Not survived']
```

```
In [10]: status_1.columns
```

```
Out[10]: Index(['age', 'year', 'nodes', 'status'], dtype='object')
```

```
In [11]: counts_1age, bin_edges_1age = np.histogram(status_1['age'], bins=10,
            density = True)
counts_1year, bin_edges_1year = np.histogram(status_1['year'], bins=10,
            density = True)
counts_1nodes, bin_edges_1nodes = np.histogram(status_1['nodes'], bins=10,
            density = True)
counts_2age, bin_edges_2age = np.histogram(status_2['age'], bins=10,
            density = True)
counts_2year, bin_edges_2year = np.histogram(status_2['year'], bins=10,
            density = True)
counts_2nodes, bin_edges_2nodes = np.histogram(status_2['nodes'], bins=10,
            density = True)
```

```
In [38]: # User defined function for creating PDF and CDF plots
def pdf_cdf_plot(counts,bin_edges):
    pdf = counts/(sum(counts))
    print('pdf values :',pdf);
    #print(bin_edges);
    cdf = np.cumsum(pdf)
    plt.title('PDF and CDF Plot')
    plt.plot(bin_edges[1:],pdf,)
    plt.plot(bin_edges[1:], cdf)
    plt.legend(['pdf','cdf'])
```

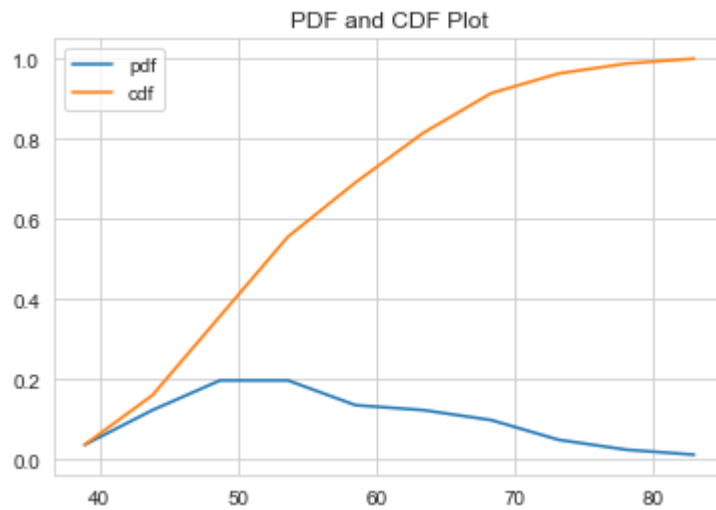
```
In [39]: pdf_cdf_plot(counts_1age, bin_edges_1age)
```

```
pdf values : [0.05333333 0.10666667 0.12444444 0.09333333 0.16444444 0.16444444
0.09333333 0.11111111 0.06222222 0.02666667]
```



```
In [40]: pdf_cdf_plot(counts_2age, bin_edges_2age)
```

```
pdf values : [0.03703704 0.12345679 0.19753086 0.19753086 0.13580247 0.12345679  
0.09876543 0.04938272 0.02469136 0.01234568]
```

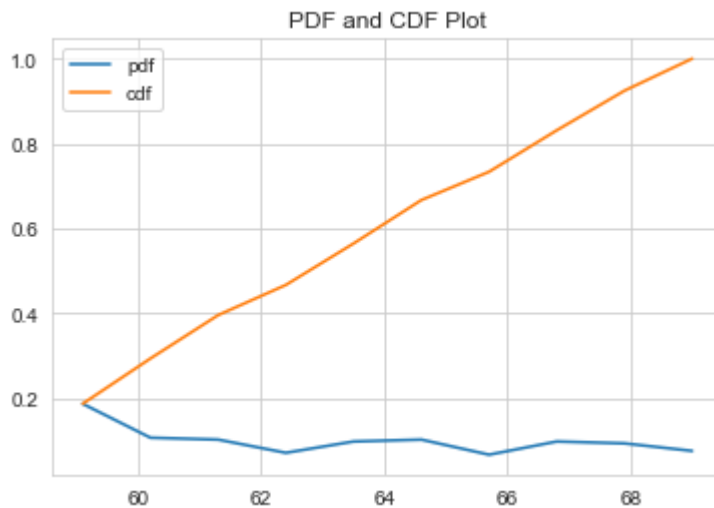


Observation:

From the above to plot , its evident that the people having age less than or equal to 40 have very high chance of survival.

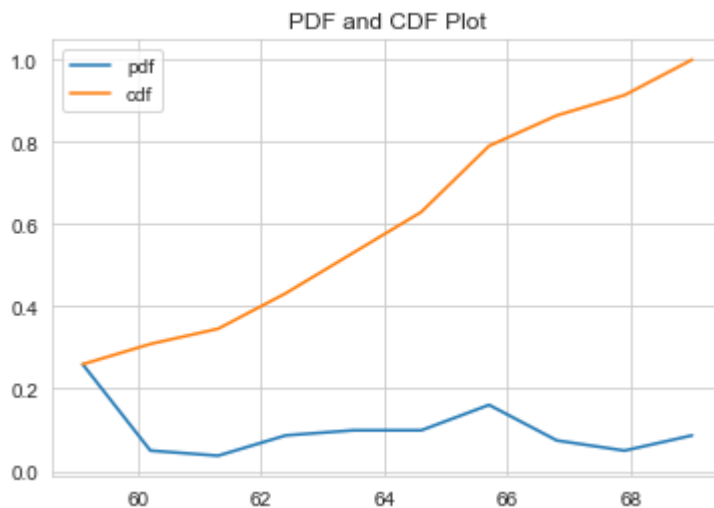
```
In [41]: pdf_cdf_plot(counts_1year, bin_edges_1year)
```

```
pdf values : [0.18666667 0.10666667 0.10222222 0.07111111 0.09777778 0.10222222  
0.06666667 0.09777778 0.09333333 0.07555556]
```



```
In [42]: pdf_cdf_plot(counts_2year, bin_edges_2year)
```

```
pdf values : [0.25925926 0.04938272 0.03703704 0.08641975 0.09876543 0.09876543  
0.16049383 0.07407407 0.04938272 0.08641975]
```

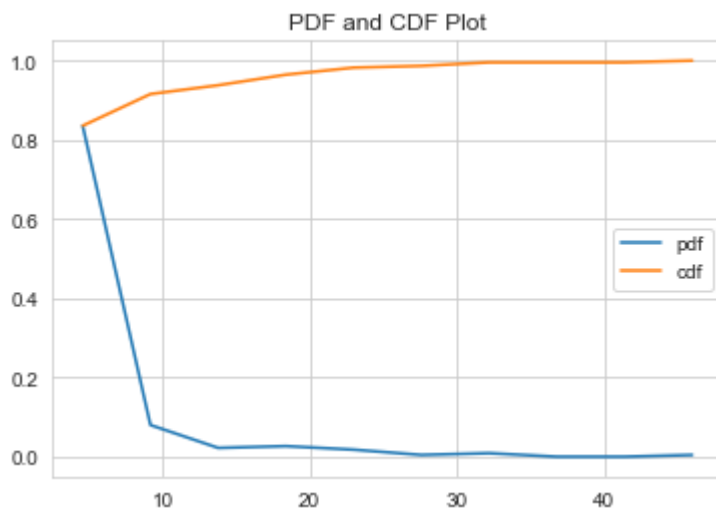


Observation:

Year feature resembles the same as the age feature.

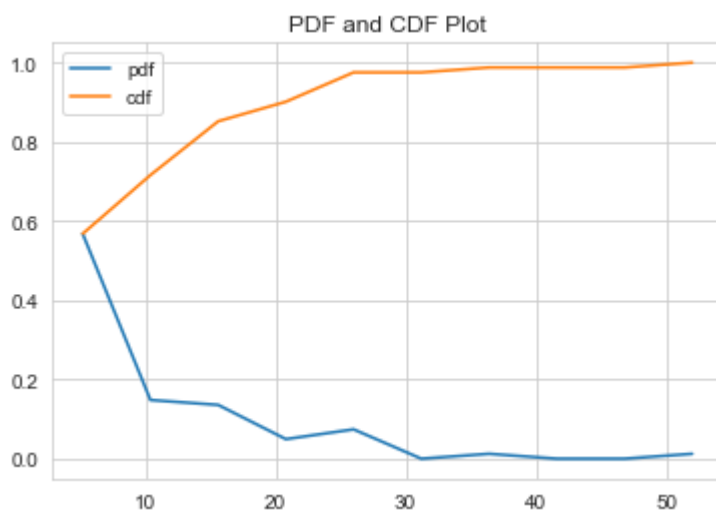
```
In [43]: pdf_cdf_plot(counts_1nodes, bin_edges_1nodes)
```

```
pdf values : [0.83555556 0.08      0.02222222 0.02666667 0.01777778 0.00444444
0.00888889 0.      0.      0.00444444]
```



```
In [44]: pdf_cdf_plot(counts_2nodes, bin_edges_2nodes)
```

```
pdf values : [0.56790123 0.14814815 0.13580247 0.04938272 0.07407407 0.
0.01234568 0.      0.      0.01234568]
```



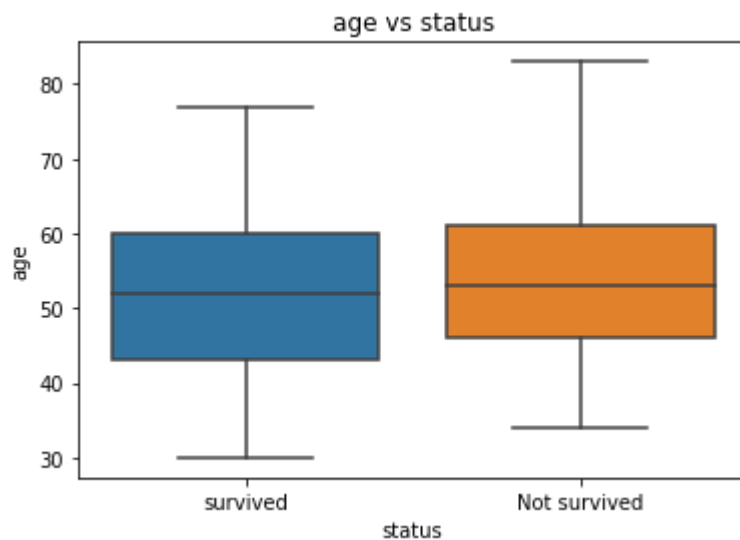
Observation:

The patient having 50+ auxillary_lymph_node never survived.

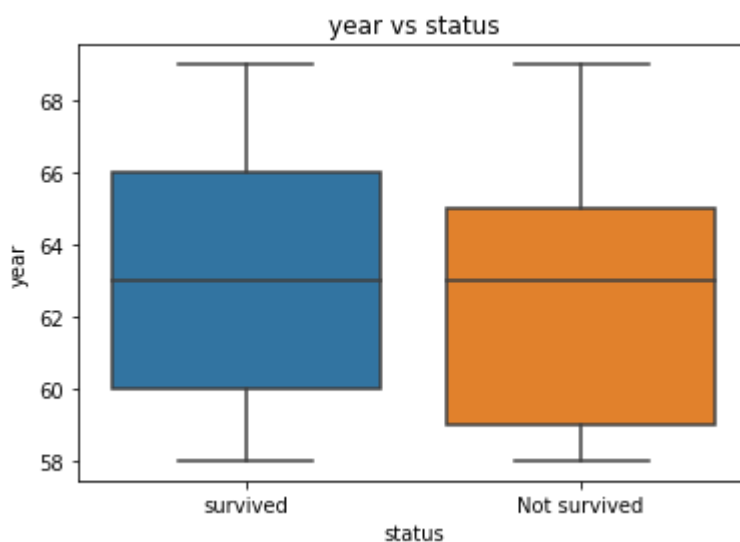
The patient having less than 5 auxillary_lymph_node has neverly 83% of survival chance.

Box Plot

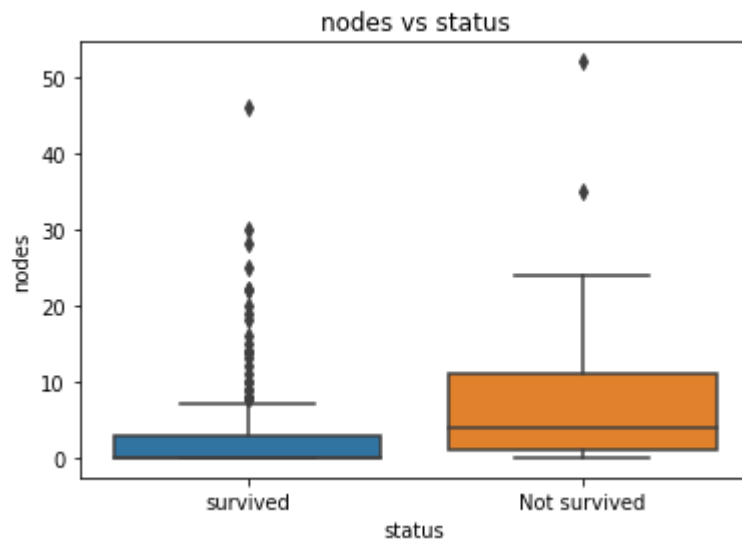
```
In [19]: sns.boxplot(x='status',y='age', data=data,).set_title("age vs status")  
plt.show()
```



```
In [20]: sns.boxplot(x='status',y='year', data=data).set_title("year vs status")  
plt.show()
```



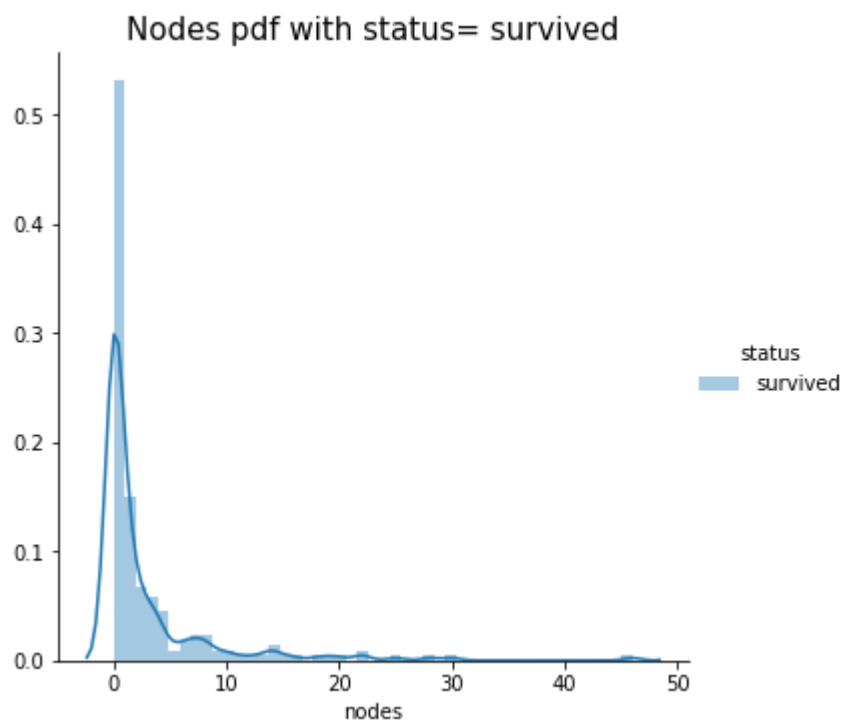
```
In [21]: sns.boxplot(x='status',y='nodes', data=data).set_title("nodes vs status")  
plt.show()
```



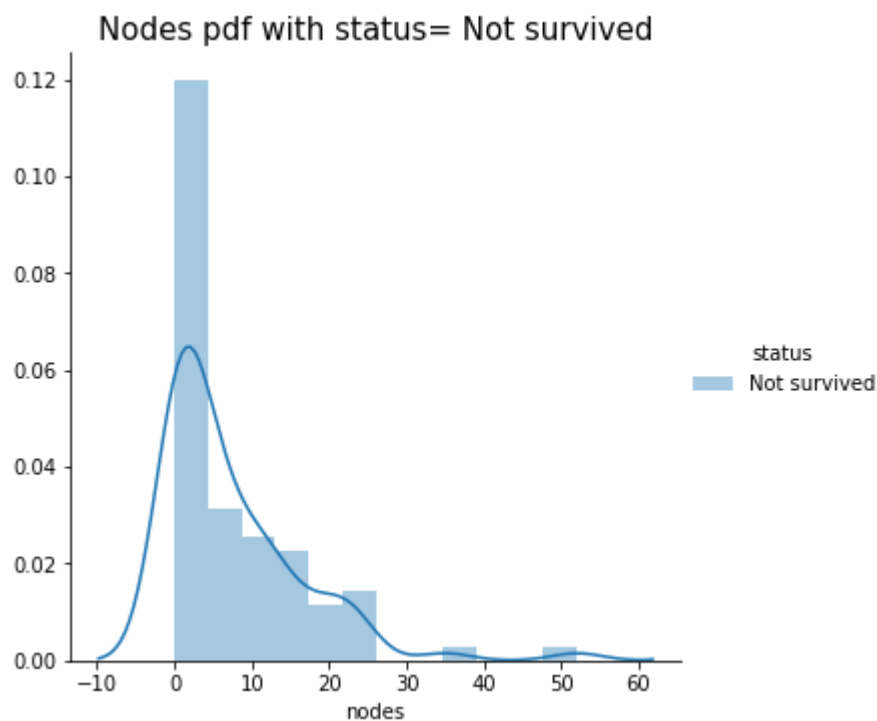
Observation:

One can easily see lot of outlier in the survival class.

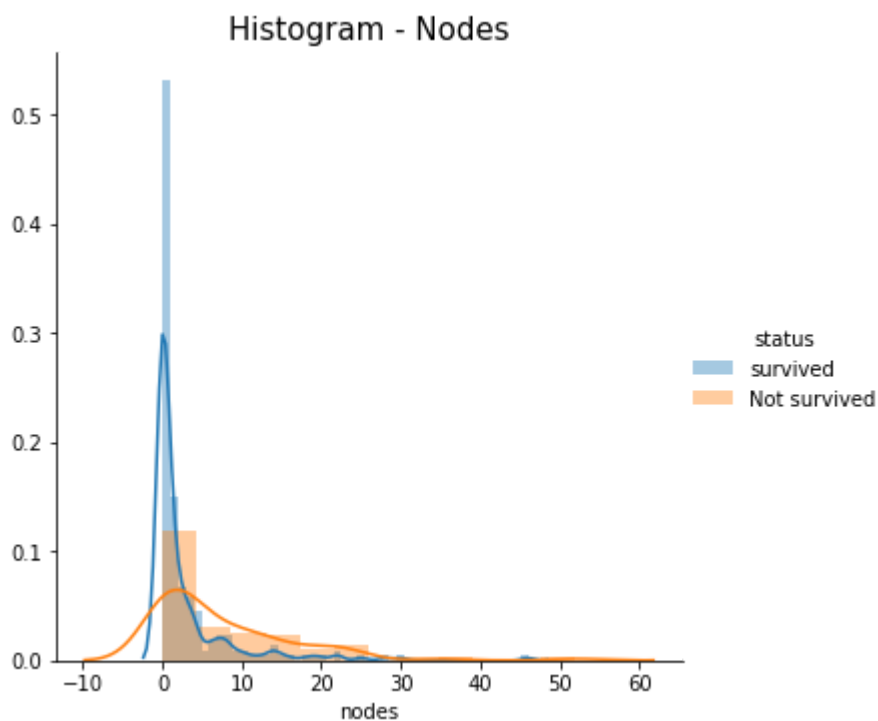

```
In [22]: sns.FacetGrid(data[data['status']=='survived'], hue="status", size=5) \
        .map(sns.distplot, "nodes") \
        .add_legend();
plt.title('Nodes pdf with status= survived',fontsize=15)
plt.show();
```



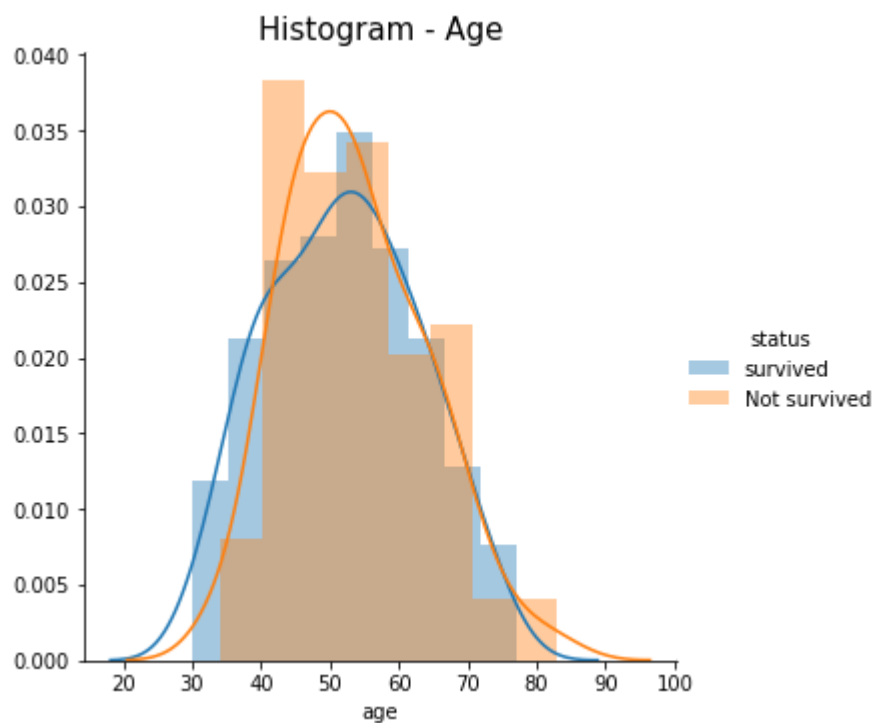
```
In [23]: sns.FacetGrid(data[data['status']=='Not survived'], hue="status", size=5) \
        .map(sns.distplot, "nodes") \
        .add_legend();
plt.title('Nodes pdf with status= Not survived', fontsize=15)
plt.show();
```



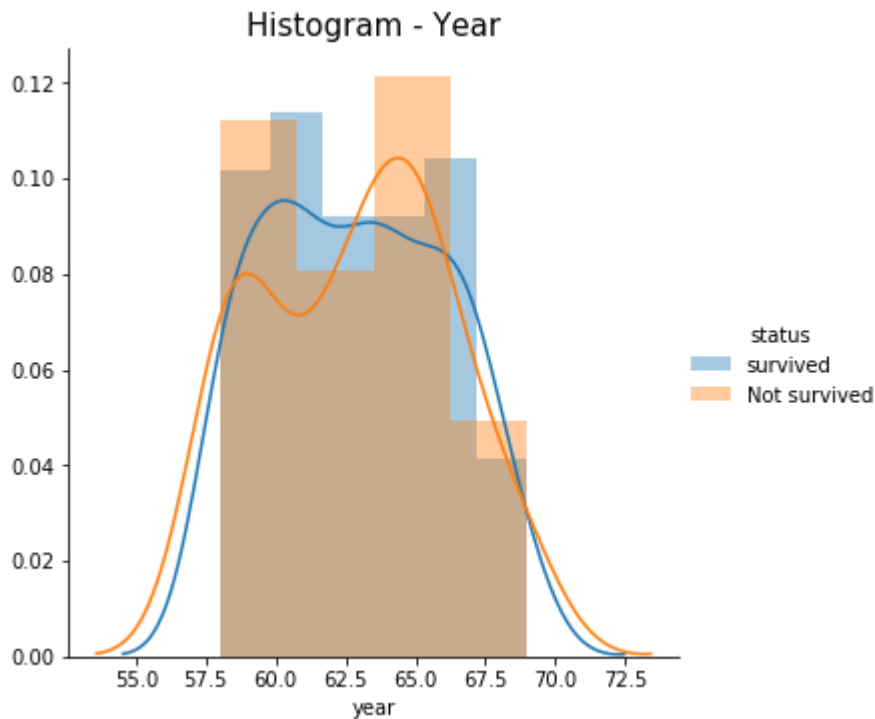
```
In [24]: sns.FacetGrid(data, hue="status", size=5) \
        .map(sns.distplot, "nodes") \
        .add_legend();
plt.title('Histogram - Nodes',fontsize=15)
plt.show();
```



```
In [25]: sns.FacetGrid(data, hue="status", size=5) \
        .map(sns.distplot, "age") \
        .add_legend();
plt.title('Histogram - Age',fontsize=15)
plt.show();
```



```
In [26]: sns.FacetGrid(data, hue="status", size=5) \
        .map(sns.distplot, "year") \
        .add_legend();
plt.title('Histogram - Year',fontsize=15)
plt.show();
```



Observation:

There is no information that can be gather from the above as they are overlapped.

Bi-varaite analysis

```
In [27]: plt.close();
sns.set_style("whitegrid");
sns.pairplot(data, hue="status", size=4);
plt.title('Pair-plot', fontsize=15)
plt.show()
```



Conclusion:

- 1) Age and nodes are useful features which talks clearly about the survival of patients who had undergone surgery for breast cancer.
- 2) People with age less than or equal 40 and having nodes less 30 have high probability of survival(more than 5 years) than others.
- 3) People having higher age have higher risk on the survival.
- 4) Around 90% of the people having node less than 10 have survived more than 5 year after a surgery.

In []:

