# **Haberman Dataset Exploratory Data Analysis**

The dataset contains cases from a study that was conducted between 1958 and 1970 at the University of Chicago's Billings Hospital on the survival of patients who had undergone surgery for breast cancer.

Attribute Information: Age of patient at time of operation (numerical) Patient's year of operation (year - 1900, numerical) Number of positive axillary nodes detected (numerical) Survival status (class attribute) 1 = the patient survived 5 years or longer 2 = the patient died within 5 year

Source: https://www.kaggle.com/gilsousa/habermans-survival-data-set

```
In [2]:
```

```
#importing the necessary libraries:
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings('ignore')
```

### In [3]:

```
#loading the dataset into dataframe:
data = pd.read_csv('haberman.csv')
```

# Initial analysis

```
In [4]:
```

```
data.head()
```

### Out[4]:

	age	year	nodes	status
0	30	64	1	1
1	30	62	3	1
2	30	65	0	1
3	31	59	2	1
4	31	65	4	1

```
In [5]:

print(data.shape)

(306, 4)

In [6]:

print(data.columns)

Index(['age', 'year', 'nodes', 'status'], dtype='object')

In [7]:

print(data['status'].value_counts())
```

```
2 81
Name: status, dtype: int64
```

#### In [8]:

```
class_1 = data[data['status']==1]
class_2 = data[data['status']==2]
print('='*18+'Class 1 Stats'+'='*18)
print(class_1.describe())
print('='*49)
print('='*18+'Class 2 Stats'+'='*18)
print(class_2.describe())
```

```
==========Class 1 Stats==========
          age
                   vear
                           nodes status
count 225.000000 225.000000 225.000000 225.0
mean 52.017778 62.862222 2.791111 1.0
std
      11.012154
                3.222915
                          5.870318
                                    0.0
     30.000000 58.000000
                        0.000000
min
                                    1.0
     43.000000 60.000000 0.000000
25%
                                   1.0
50%
     52.000000 63.000000 0.000000
                                   1.0
                                   1.0
75%
     60.000000 66.000000
                         3.000000
      77.000000 69.000000 46.000000
max
                                    1.0
_____
==========Class 2 Stats==========
                         nodes status
          age
                vear
count 81.000000 81.000000 81.000000 81.0
mean 53.679012 62.827160 7.456790
                               2.0
std
     10.167137
              3.342118
                       9.185654
                                  0.0
     34.000000 58.000000 0.000000
min
                                  2.0
   46.000000 59.000000 1.000000
25%
                                 2.0
50% 53.000000 63.000000 4.000000
                                 2.0
75% 61.000000 65.000000 11.000000
                                 2.0
max 83.000000 69.000000 52.000000
                                 2.0
```

# **Initial Observations:**

- The means of the attributes 'age' and 'year' for class 1 and class 2 are overlapping. The quartiles also are very close.
- The 'nodes' attribute has fairly seperate mean for class 1 and class 2 and may act as the deciding attribute in case of classification.
- The mean for the 'nodes' attribute suggests that patients having less nodes had higher chances of surviving 5 years or longer.

# **Univariate Analysis**

#### In [9]:

```
sns.set_style('darkgrid')
sns.FacetGrid(data, hue='status').map(sns.distplot, 'age').add_legend()
plt.title('Distribution of survival status wrt to age.')
plt.show()
```

0.035 0.030 0.025 0.020 0.015 0.010 0.005

60

age

100

80

Distribution of survival status wrt to age.

## **Observations:**

40

0.000

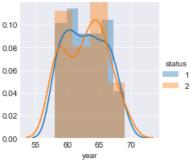
As was suggested by initial analysis, age is not a very strong deciding factor for survival in this case.

- Patients in the age group ~25-40 had slightly higher chances of surviving 5 years or more.
- Patients in the age group ~40-60 had lesser chances of surviving 5 years or more.

#### In [10]:

```
sns.set_style('darkgrid')
sns.FacetGrid(data, hue='status').map(sns.distplot, 'year').add_legend()
plt.title('Distribution of survival status wrt to year.')
plt.show()
```

Distribution of survival status wrt to year.

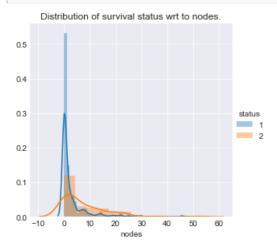


#### **Observations:**

- Patients who were operated upon between ~1958-1962 had comparatively more chances of surviving 5 years or more.
- Patients who were operated upon between ~1962-1966 had comparatively lesser chances of surviving 5 years or more.
- Like the age attribute, the year also does not play a very huge role in deciding the chances of survival.

### In [11]:

```
sns.set_style('darkgrid')
sns.FacetGrid(data, hue='status', size=4).map(sns.distplot, 'nodes').add_legend()
plt.title('Distribution of survival status wrt to nodes.')
plt.show()
```



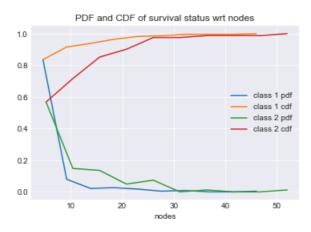
### Obsevations:

- There is a significant higher chance of survival for patients having 1 or no nodes.
- For patients having more than 3 nodes, the chances of survival 5 years or more dramatically decreases.
- The number of nodes looks like the strongest deciding factor in case of surviving 5 years or more.

#### In [12]:

```
#Since 'nodes' is the most important attribute, let's explore the cdf:
counts_1, edges_1 = np.histogram(class_1['nodes'], bins=10, density=True)
pdf_1 = counts_1/sum(counts_1)
cdf_1 = np.cumsum(pdf_1)
print('class 1')
print(pdf_1)
print(edges_1)
plt.plot(edges_1[1:], pdf_1)
plt.plot(edges_1[1:], cdf_1)
```

```
plt.xlabel('nodes')
print('=======
                    -----<sup>1</sup>)
counts 2, edges 2 = np.histogram(class 2['nodes'], bins=10, density=True)
pdf 2 = counts \frac{2}{sum(counts 2)}
cdf 2 = np.cumsum(pdf 2)
print('class 2')
print(pdf 2)
print(edges 2)
plt.plot(edges_2[1:], pdf_2)
plt.plot(edges 2[1:], cdf 2)
plt.xlabel('nodes')
plt.title('PDF and CDF of survival status wrt nodes')
plt.legend(['class 1 pdf','class 1 cdf','class 2 pdf','class 2 cdf'])
plt.show()
class 1
[0.83555556 0.08
                     0.02222222 0.02666667 0.01777778 0.00444444
 0.00888889 0. 0.
                               0.00444444]
[ 0. 4.6 9.2 13.8 18.4 23. 27.6 32.2 36.8 41.4 46. ]
```



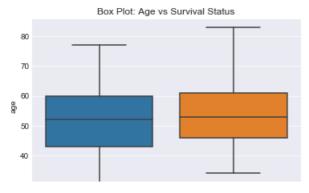
# Observations:

• More than 83% of the patients who survived had less than 4.6 nodes.

### Analysis with box and violin plots

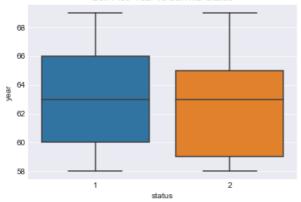
```
In [13]:
```

```
sns.boxplot(x='status',y='age',data=data)
plt.title('Box Plot: Age vs Survival Status')
plt.show()
sns.boxplot(x='status',y='year',data=data)
plt.title('Box Plot: Year vs Survival Status')
plt.show()
sns.boxplot(x='status',y='nodes',data=data)
plt.title('Box Plot: Nodes vs Survival Status')
plt.show()
```

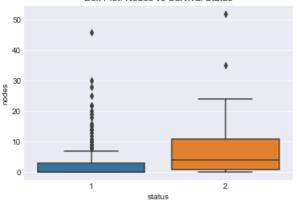








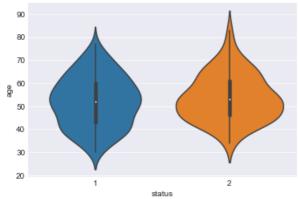
Box Plot: Nodes vs Survival Status



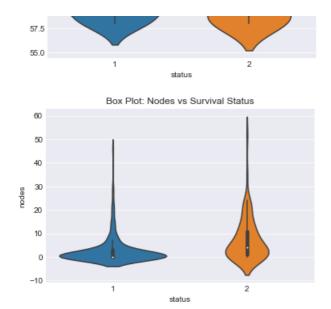
# In [17]:

```
sns.violinplot(x='status',y='age',data=data)
plt.title('Box Plot: Age vs Survival Status')
plt.show()
sns.violinplot(x='status',y='year',data=data)
plt.title('Box Plot: Year vs Survival Status')
plt.show()
sns.violinplot(x='status',y='nodes',data=data)
plt.title('Box Plot: Nodes vs Survival Status')
plt.show()
```

Box Plot: Age vs Survival Status







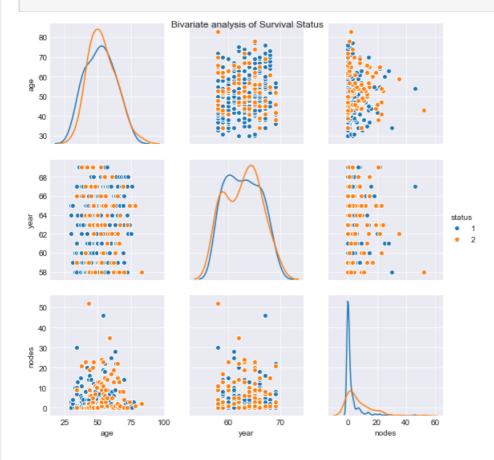
#### **Observations:**

• The violin plot combined with the box plot for 'nodes' strengthens our earlier observation that lesser the number of nodes, more the chances of surviving 5 years or more.

# **Bivariate Analysis**

### In [22]:

```
sns.pairplot(data, hue='status', x_vars = ['age','year','nodes'], y_vars = ['age','year','nodes'],
diag_kind = 'kde').fig.\
suptitle('Bivariate analysis of Survival Status')
plt.show()
```



# Observations:

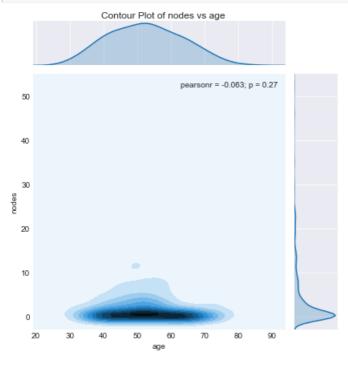
Though not much significant, the pairplots do provide some additional insights:

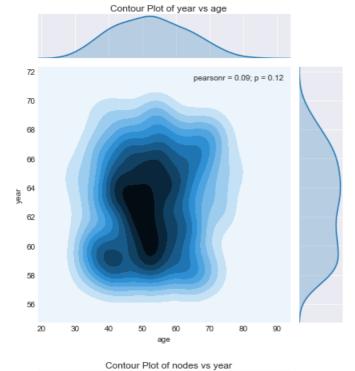
- Looking at pairplot of nodes and age, we can deduce that patients less than 40 years of age had more survivals.
- The patients operated upon in 1960 had more survival rate.
- Patients of age more than 60 who were operated upon in 1968 had more survivals.

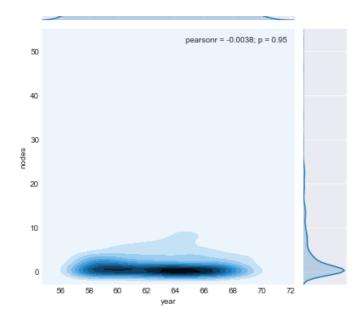
# **Analysis with Contour Plot**

### In [36]:

```
sns.jointplot(x='age',y='nodes',kind='kde',data=data)
plt.title('Contour Plot of nodes vs age',y=1.2,x=-3)
plt.show()
sns.jointplot(x='age',y='year',kind='kde',data=data)
plt.title('Contour Plot of year vs age',y=1.2,x=-3)
plt.show()
sns.jointplot(x='year',y='nodes',kind='kde',data=data)
plt.title('Contour Plot of nodes vs year',y=1.2,x=-3)
plt.show()
```







# Observations:

• From the first and third contour plot we can loosely see that, irrespective of the age and year of operation, the number of nodes decides the survival category.

# **Conclusions:**

# From the above data analysis we conclude that:

- Number of nodes are most important in deciding the survival category.
- More than 83% of the patients who survived had less than 4.6 nodes.
- Age and Year play a very minor role in deciding the survival category.