

```
In [5]: cd "D:\AppliedAI\"
```

D:\AppliedAI

Importing packages

```
In [6]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import warnings

warnings.filterwarnings('ignore')
```

loading the dataset and peeking into it

Haberman's Survival : Exploratory Data Analysis Haberman's Cancer survival :EDA

Haberman's survival 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 the 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 years

Goal is to find the chances of survival for more than 5 years after treatment

Age group, gender,etc. which are likely to die early after treatment

In [7]: `data = pd.read_csv("haberman.csv")`
`data.head()`

Uploading .csv file

Out[7]:

	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 [8]: `data.info()`

checking for number of

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 306 entries, 0 to 305
Data columns (total 4 columns):
age          306 non-null int64
year         306 non-null int64
nodes        306 non-null int64
status       306 non-null int64
dtypes: int64(4)
memory usage: 9.7 KB
```

In [9]: `data.columns`

Out[9]: `Index(['age', 'year', 'nodes', 'status'], dtype='object')`

In [10]: `data.values`

Out[10]: `array([[30, 64, 1, 1],`
 `[30, 62, 3, 1],`
 `[30, 65, 0, 1],`
 `...,`
 `[77, 65, 3, 1],`
 `[78, 65, 1, 2],`
 `[83, 58, 2, 2]], dtype=int64)`

In [11]: data.describe()

basic stats

Out[11]:

	age	year	nodes	status
count	306.000000	306.000000	306.000000	306.000000
mean	52.457516	62.852941	4.026144	1.264706
std	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
max	83.000000	69.000000	52.000000	2.000000

Observations

median age of patients is 52 years

75% of the patients have nodes <=4.

peeking into number of years survived

```
In [12]: x = data["status"]                                # total number of survivors
y = data[data["status"]==2]["status"]                    # total number of survivors w

print("total number of survivors are {}".format(len(x)))
print("total number of survivors died within 5 years of operation are {}".format(len(y)))
print("the percentage of people who live for more than 5 years is {}".format((306-len(y))/306))
```

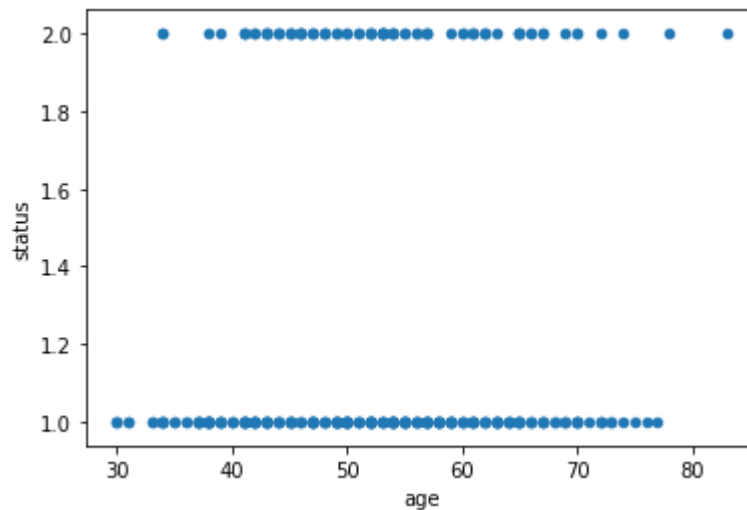
total number of survivors are 306

total number of survivors died within 5 years of operation are 81

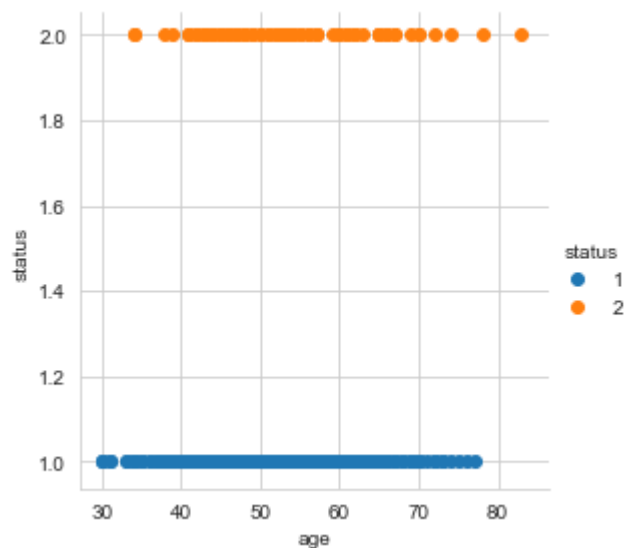
the percentage of people who live for more than 5 years is 0.7352941176470589

2D plots

```
In [13]: data.plot(kind='scatter', x='age', y='status') ;
plt.show()
```



```
In [14]: sns.set_style("whitegrid");
sns.FacetGrid(data, hue="status", size=4) \
    .map(plt.scatter, "age", "status") \
    .add_legend();
plt.show();
```



```
<p style = "background : pink">
Observations : <br>
1. On a positive note, we can say, survivors whose age is below 40 are likely
to survive for more than 5 years <br>
<br>
let us calculate the percentage of patients who can survive for more than 5
years having age<40
```

```
In [27]: a=data[(data["status"]==1) & (data["age"]<=40)]["status"].sum()      # finding
b=data[(data["status"]==2) & (data["age"]<=40)]["status"].sum()      # finding
print(a)
print(b)
print(a/(a+b))      # calculate
```

39

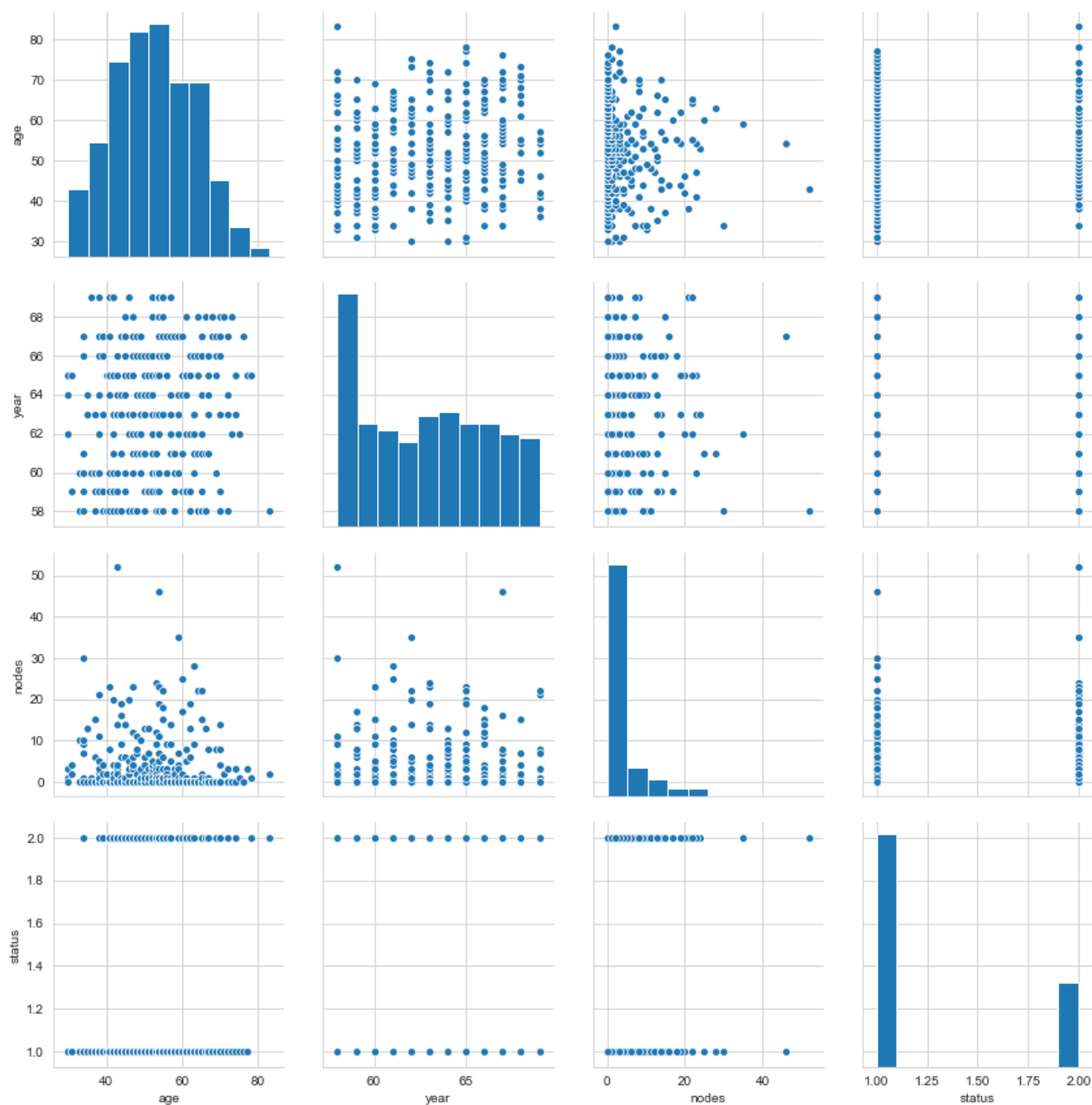
8

0.8297872340425532

Pair Plots

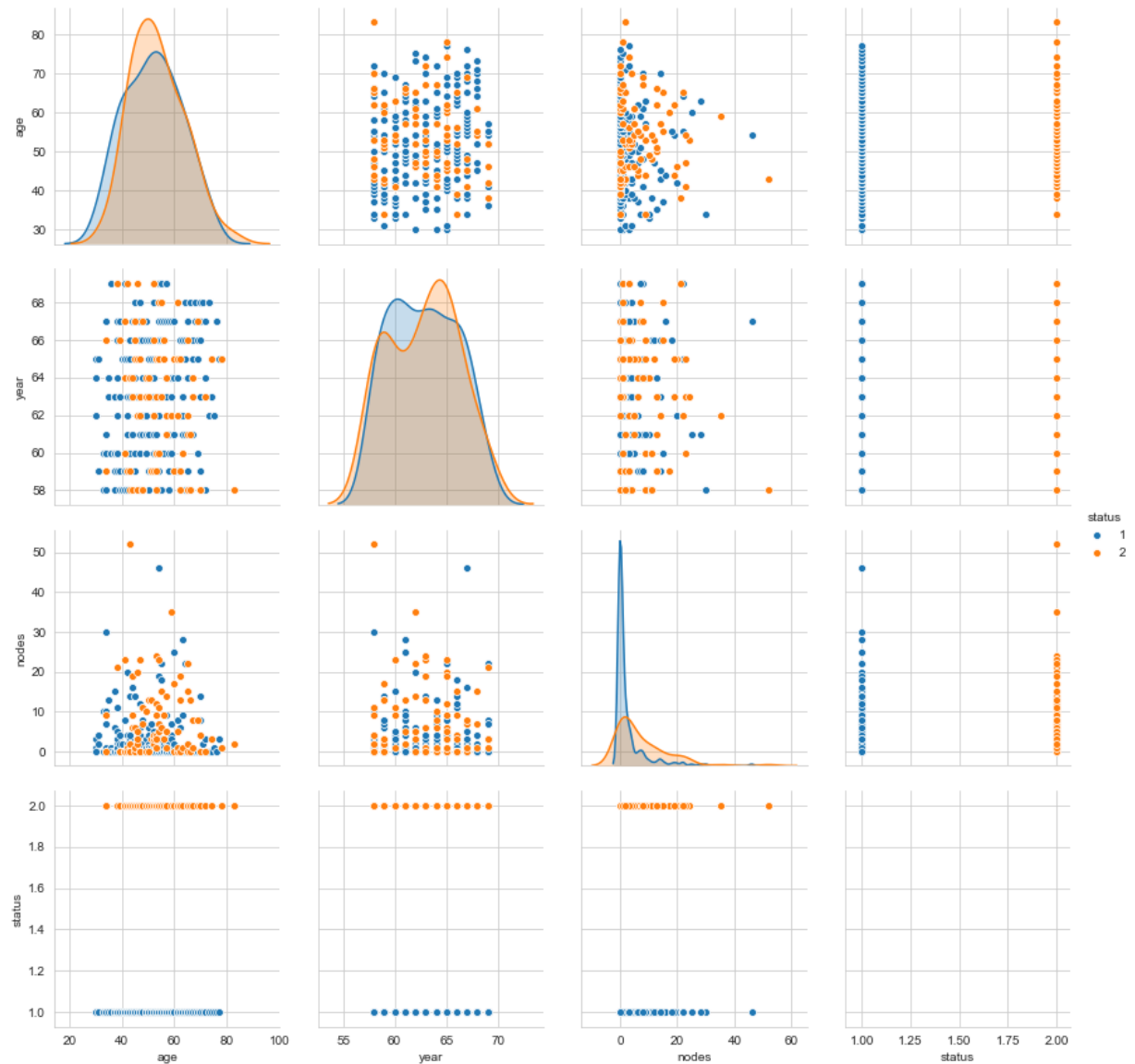
```
In [28]: sns.set_style("whitegrid");
sns.pairplot(data, size=3);
plt.show()
```

alltogether without 'hue'



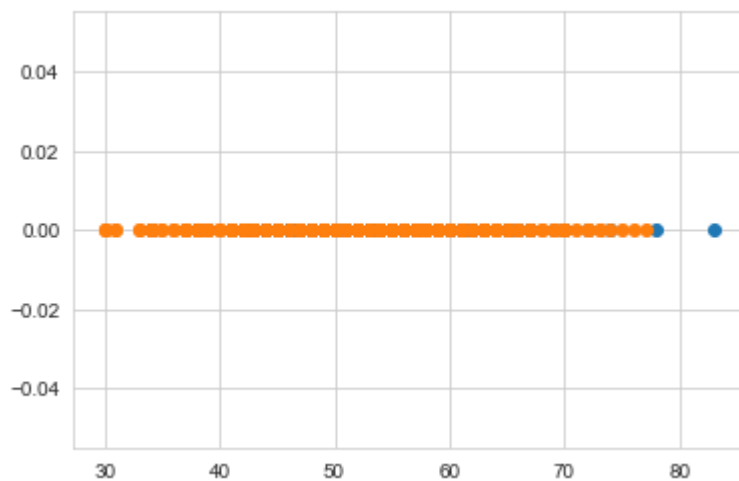
```
In [29]: sns.set_style("whitegrid");
sns.pairplot(data, hue = "status", size=3);
plt.show()
```

with 'hue on sta

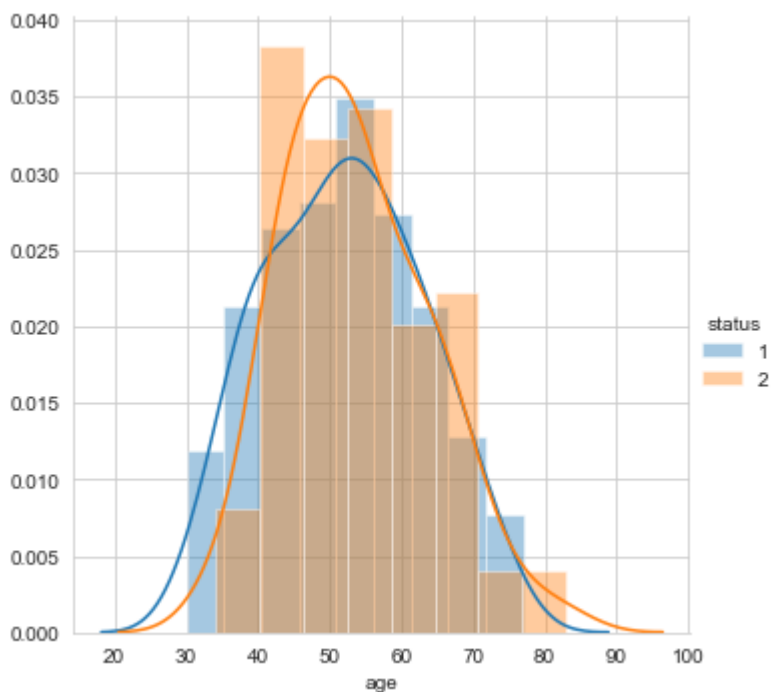


Histogram

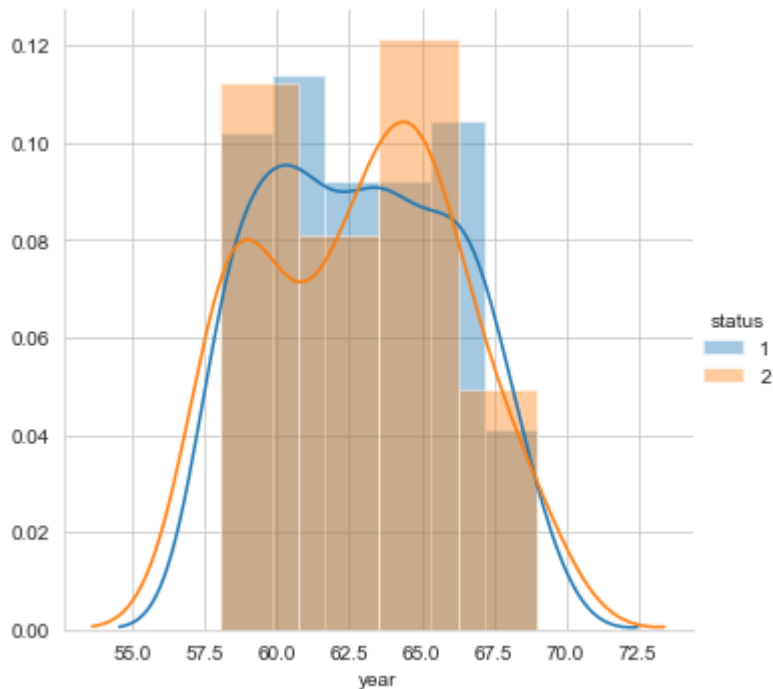
```
In [30]: data_below5years = data.loc[data["status"] == 2];
data_above5years = data.loc[data["status"] == 1];
#print(iris_setosa["petal_length"])
plt.plot(data_below5years["age"], np.zeros_like(data_below5years["age"]), 'o')
plt.plot(data_above5years["age"], np.zeros_like(data_above5years["age"]), 'o')
plt.show()
```



```
In [31]: sns.FacetGrid(data, hue="status", size=5) \
    .map(sns.distplot, "age") \
    .add_legend();
plt.show();
```

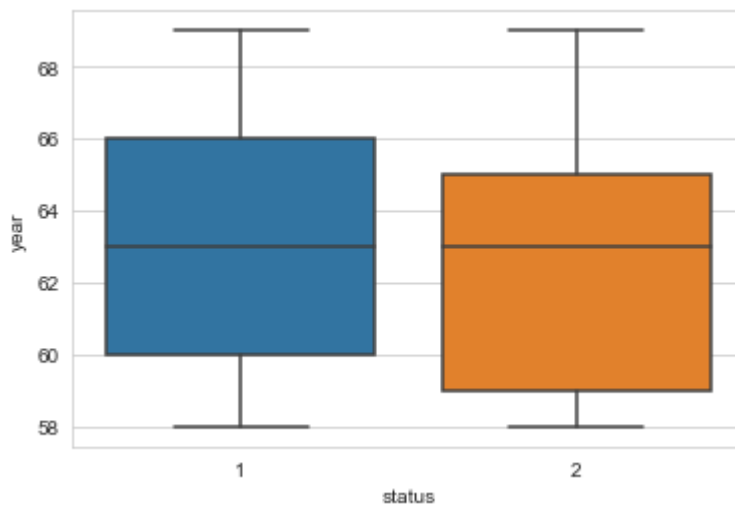



```
In [32]: sns.FacetGrid(data, hue="status", size=5) \
        .map(sns.distplot, "year") \
        .add_legend();
plt.show();
```

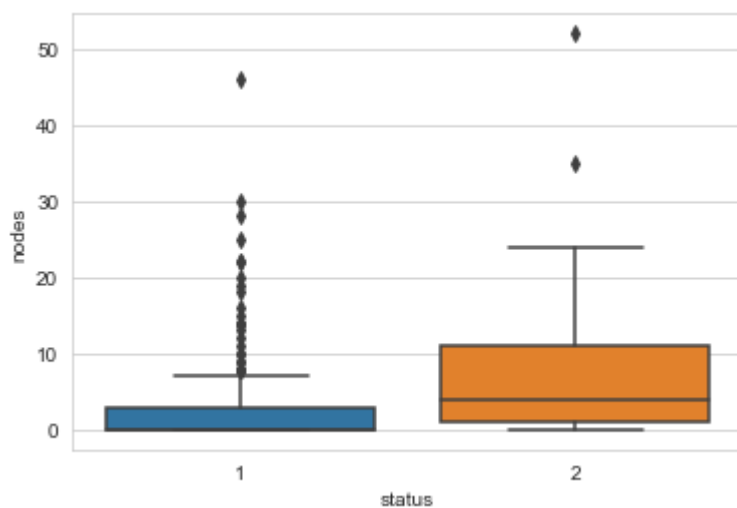


Box plot and Whiskers

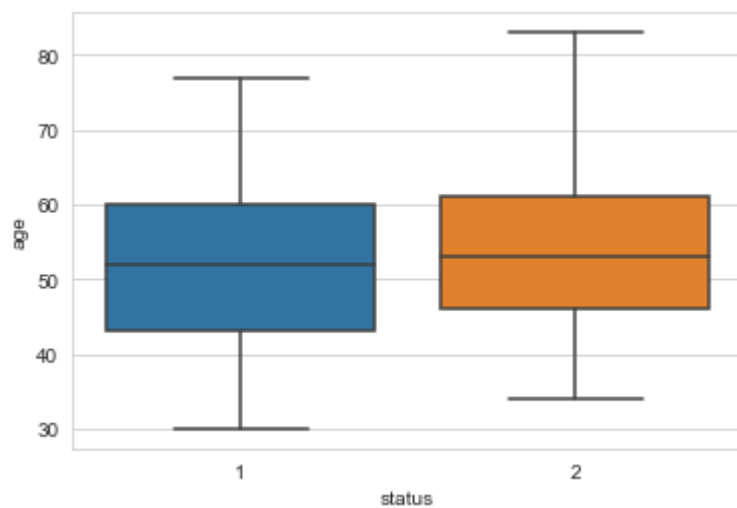
```
In [33]: sns.boxplot(x='status', y='year', data=data)
plt.show()
```



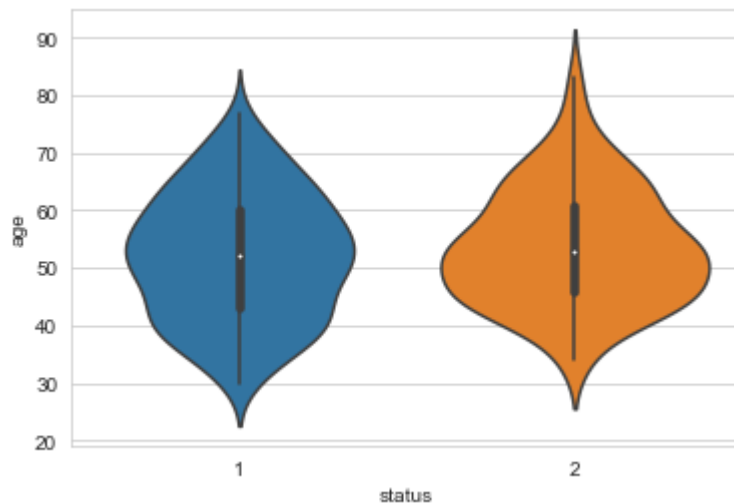
```
In [34]: sns.boxplot(x='status',y='nodes', data=data)  
plt.show()
```



```
In [35]: sns.boxplot(x='status',y='age', data=data)  
plt.show()
```



```
In [36]: sns.violinplot(x="status", y="age", data=data, size=8)  
plt.show()
```



Take away points

median age of patients is 52 years

- 75% of the patients have nodes ≤ 4 .
- total number of survivors are 306
- total number of survivors died within 5 years of operation are 81
- the percentage of people who live for more than 5 years is 0.7352941176470589
- On a positive note, we can say, survivors whose age is below 40 are likely to survive for more than 5 years