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In [ ]: # Dataset: Differentiated Thyroid Cancer Recurrence
```

This data set contains 13 clinicopathologic features aiming to predict recur

Gathering Data

```
In [1]: # Importing necessary libraries
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```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

#import dataset
patient_data = pd.read_csv('https://raw.githubusercontent.com/karanzijm/MLEx
patient_data
```

```
Out[1]:
```

	Age	Gender	Smoking	Hx Smoking	Hx Radiotherapy	Thyroid Function	Physical Examination	A
0	27	F	No	No	No	Euthyroid	Single nodular goiter-left	
1	34	F	No	Yes	No	Euthyroid	Multinodular goiter	
2	30	F	No	No	No	Euthyroid	Single nodular goiter-right	
3	62	F	No	No	No	Euthyroid	Single nodular goiter-right	
4	62	F	No	No	No	Euthyroid	Multinodular goiter	
...	...	...	...	...	...	...	...	
378	72	M	Yes	Yes	Yes	Euthyroid	Single nodular goiter-right	
379	81	M	Yes	No	Yes	Euthyroid	Multinodular goiter	
380	72	M	Yes	Yes	No	Euthyroid	Multinodular goiter	
381	61	M	Yes	Yes	Yes	Clinical Hyperthyroidism	Multinodular goiter	
382	67	M	Yes	No	No	Euthyroid	Multinodular goiter	

383 rows × 17 columns

```
In [ ]: # Patient columns
-       T: Tumor size and extent
-       - T1a and T1b: Indicates a small tumor size, typically less than 2cm i
-       N: Lymph node involvement.
-       - N0: No regional lymph node metastasis (cancer has not spread to nearby
-       - N1b: Cancer has spread to certain lymph nodes (such as cervical or upp
-       M: Distant metastasis.
-       - M0: No distant metastasis (cancer has not spread to other parts of the
-       Hx Smoking: History of smoking
-       Hx Radiotherapy: History of radiotherapy.
-       Thyroid Function: The functional state of the thyroid.
-       Euthyroid: This means that the thyroid is functioning normally. The
-       Clinical Hyperthyroidism: This indicates that the patient has overactive
-       Clinical Hypothyroidism: This indicates that the patient has underac
-       Physical Examination: Results of a physical examination of the thyro
-       Single nodular goiter-left: A single nodule (enlarged portion of the
-       Multinodular goiter: Multiple nodules are present in the thyroid gla
-       Stages:
-       - Stages I & II are typically early-stage cancers, with Stage II sometimes
-       - Stage III often involves larger tumors or some lymph node involvement bu
-       - Stage IV is advanced, with the cancer either spreading to nearby tissues
-       Adenopathy: Swelling or disease of lymph nodes.
-       "No" indicates no adenopathy, meaning there is no lymph node involve
-       Pathology: The study of the disease, especially cancer.
-       Focality: The number of distinct tumor sites.
-       Uni-focal: The cancer is localized to a single focus or site within
-       Risk: The level of cancer risk or recurrence.
-       "Low" means the patient is considered at low risk for recurrence or
-       Response: The clinical assessment of how well the patient's condition resp
```

```
In [ ]: All columns seem to have all their data consistent at first glance. There se

Check is the data has any null and duplicate values and remove them.
```

```
In [3]: print(patient_data.isnull().sum())
```

```
Age                0
Gender             0
Smoking            0
Hx Smoking         0
Hx Radiothreapy    0
Thyroid Function   0
Physical Examination 0
Adenopathy         0
Pathology          0
Focality           0
Risk               0
T                  0
N                  0
M                  0
Stage              0
Response           0
Recurred           0
dtype: int64
```

```
In [5]: print(patient_data.duplicated().sum())
patient_data = patient_data.drop_duplicates()
```

19

```
In [7]: patient_data.sample(5)
```

```
Out[7]:
```

	Age	Gender	Smoking	Hx Smoking	Hx Radiotherapy	Thyroid Function	Physical Examination	Adenopathy
366	64	F	No	Yes	No	Euthyroid	Multinodular goiter	
105	42	F	No	No	No	Euthyroid	Single nodular goiter-right	
89	31	M	Yes	No	No	Euthyroid	Multinodular goiter	R
238	29	F	Yes	No	No	Euthyroid	Single nodular goiter-left	
280	37	F	No	No	No	Euthyroid	Single nodular goiter-right	

```
In [9]: print(patient_data.dtypes)
```

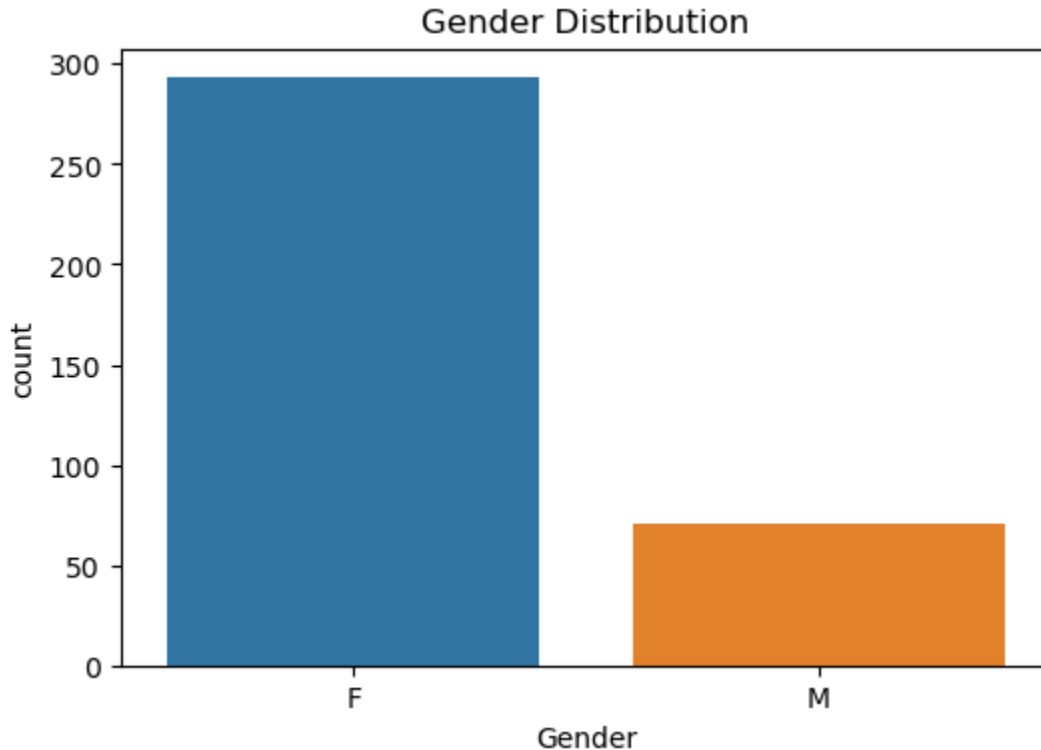
```
Age                int64
Gender              object
Smoking             object
Hx Smoking          object
Hx Radiotherapy     object
Thyroid Function    object
Physical Examination object
Adenopathy          object
Pathology           object
Focality            object
Risk                object
T                   object
N                   object
M                   object
Stage               object
Response            object
Recurred            object
dtype: object
```

```
In [11]: patient_data.Gender.value_counts()
```

```
Out[11]: Gender
F      293
M       71
Name: count, dtype: int64
```

```
In [13]: plt.figure(figsize=(6, 4))
sns.countplot(x='Gender', data=patient_data)
```

```
plt.title('Gender Distribution')  
plt.show()
```



```
In [ ]: Females are more than males.
```

```
In [15]: from scipy.stats import chi2_contingency  
import numpy as np  
  
def cramers_v(x, y):  
    confusion_matrix = pd.crosstab(x, y)  
    chi2 = chi2_contingency(confusion_matrix)[0]  
    n = confusion_matrix.sum().sum()  
    r, k = confusion_matrix.shape  
    return np.sqrt(chi2 / (n * (min(k, r) - 1)))  
  
print(cramers_v(patient_data['Hx Smoking'], patient_data['Recurred']))  
print(cramers_v(patient_data['Pathology'], patient_data['Recurred']))  
print(cramers_v(patient_data['Gender'], patient_data['Recurred']))  
print(cramers_v(patient_data['Smoking'], patient_data['Recurred']))  
print(cramers_v(patient_data['Focality'], patient_data['Recurred']))  
print(cramers_v(patient_data['M'], patient_data['Recurred']))  
print(cramers_v(patient_data['Stage'], patient_data['Recurred']))  
print(cramers_v(patient_data['Adenopathy'], patient_data['Recurred']))  
print(cramers_v(patient_data['T'], patient_data['Recurred']))  
print(cramers_v(patient_data['N'], patient_data['Recurred']))
```

```

0.11718749999999997
0.25107586767432133
0.3101425326756299
0.3164898153242812
0.36236533566053664
0.3372913838647582
0.4993900250890623
0.6331196116773835
0.5996441476424428
0.624612084273687

```

In [ ]: Above figures show the association of the individual features and recurrence. From the figures, Tumor size(T), Lymph node involvement(N) and Adenopathy ha

In [ ]: We can also use the Chi-square & p-value as another way to determine the extent of the association.

Chi-square: This value measures the difference between the observed data and the expected data.

P-value: This value indicates significance of the correlation between observed and expected data.

```

In [17]: ct_gender = pd.crosstab(patient_data['Gender'], patient_data['Recurred'])
print(ct_gender)

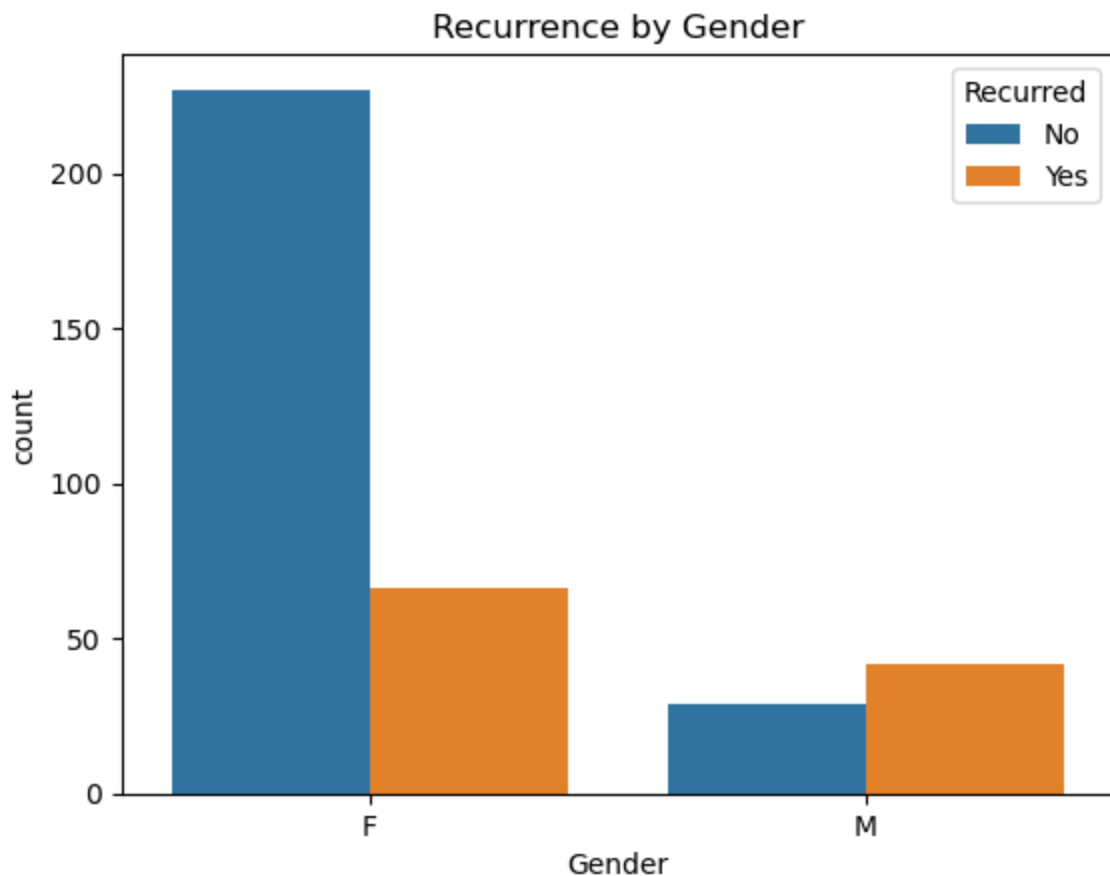
chi2, p, dof, expected = chi2_contingency(ct_gender)
print(f'Chi-square: {chi2}, p-value: {p}')

# Graph To Analyzing the effect of Gender on Recurred
sns.countplot(data=patient_data, x='Gender', hue='Recurred')
plt.title('Recurrence by Gender')
plt.show()

```

	No	Yes
Gender		
F	227	66
M	29	42

Chi-square: 35.01257416910131, p-value: 3.2758306763157053e-09



In [ ]: There **is** a statistical significant relationship between gender **and** recurrence. Males seem to have a higher recurrence rate compared to females. But it may have been better to have more male specimen so **as** to draw a better

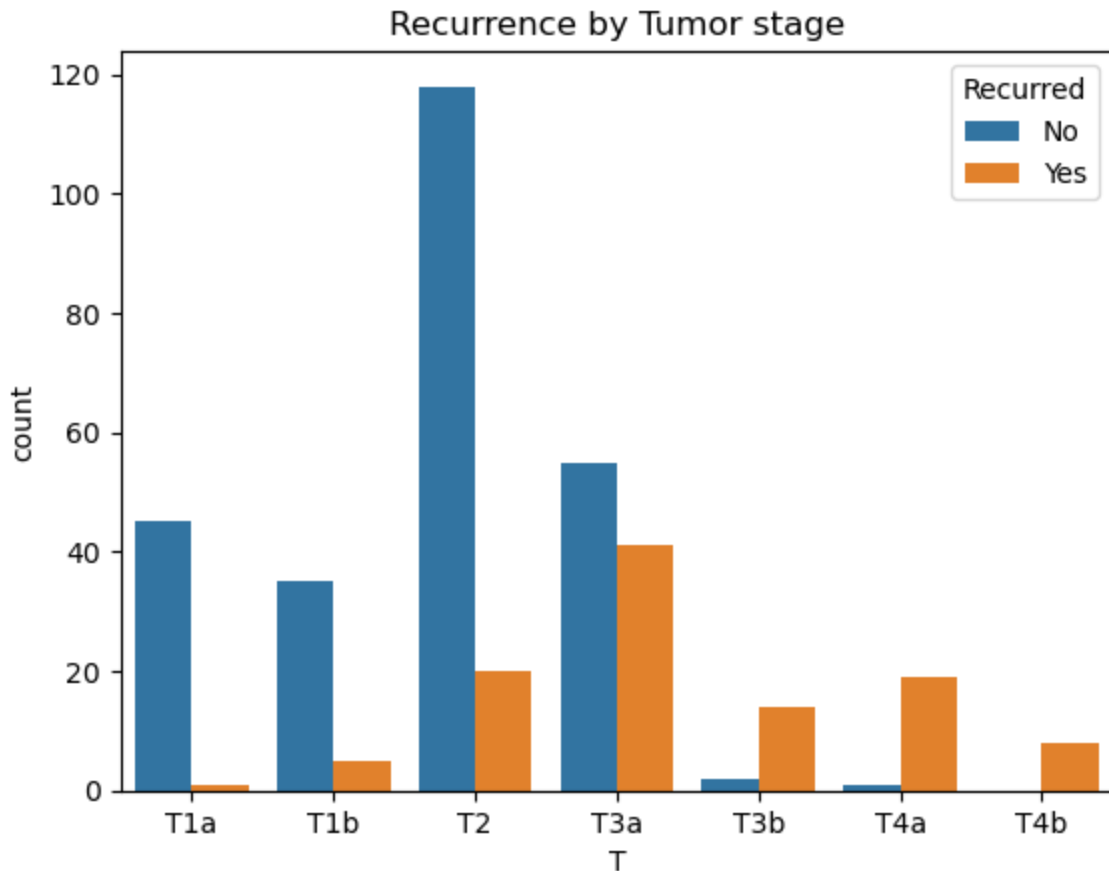
```
In [19]: ct_t = pd.crosstab(patient_data['T'], patient_data['Recurred'])
print(ct_t)

chi2, p, dof, expected = chi2_contingency(ct_t)
print(f'Chi-square: {chi2}, p-value: {p}')

# Graph To Analyzing the effect of Tumor stage on Recurred
sns.countplot(data=patient_data, x='T', hue='Recurred')
plt.title('Recurrence by Tumor stage')
plt.show()
```

Recurred	No	Yes
T		
T1a	45	1
T1b	35	5
T2	118	20
T3a	55	41
T3b	2	14
T4a	1	19
T4b	0	8

Chi-square: 130.88460978386675, p-value: 8.370007602185988e-26



In [ ]: There **is** a strong association between tumor stage (T) **and** recurrence (Recurred).

- Early-stage tumors like T1a **and** T1b show fewer cases of recurrence.
- Later stages like T3a, T3b, T4a, **and** T4b show more frequent recurrence, in

This result **is** highly statistically significant, meaning that the stage of tumor **is** associated with recurrence. However, this test only shows association, **not** causation. Tumor size/stage **is**

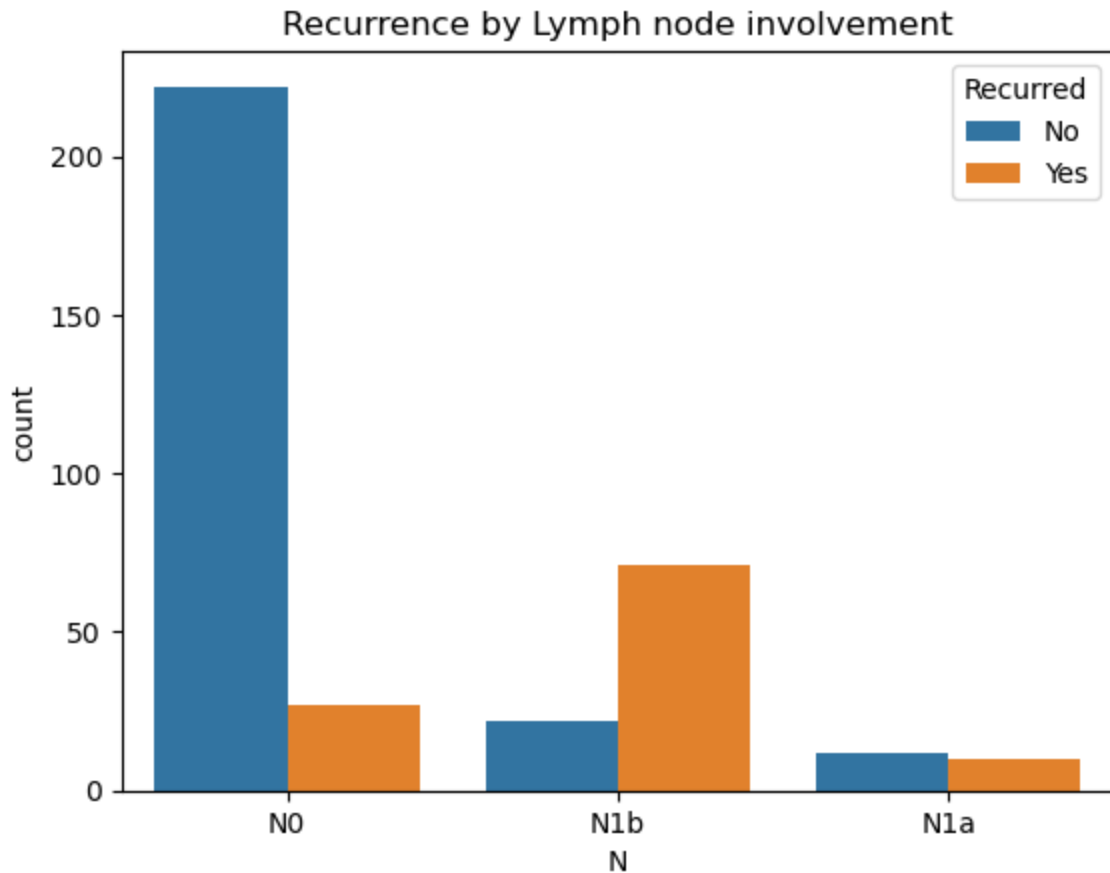
```
In [21]: ct_n = pd.crosstab(patient_data['N'], patient_data['Recurred'])
print(ct_n)

chi2, p, dof, expected = chi2_contingency(ct_n)
print(f'Chi-square: {chi2}, p-value: {p}')

# Graph To Analyzing the effect of Lymph node involvement on Recurred
sns.countplot(data=patient_data, x='N', hue='Recurred')
plt.title('Recurrence by Lymph node involvement')
plt.show()
```

Recurred	No	Yes
N		
N0	222	27
N1a	12	10
N1b	22	71

Chi-square: 142.0110531187419, p-value: 1.4544260034549868e-31



In [ ]: There **is** a very strong **and** significant association between the N (lymph node involvement) and Recurred. Patients **with** higher lymph node involvement (N1a, N1b) are much more likely to recur.

```
In [23]: ct_m = pd.crosstab(patient_data['M'], patient_data['Recurred'])
print(ct_m)

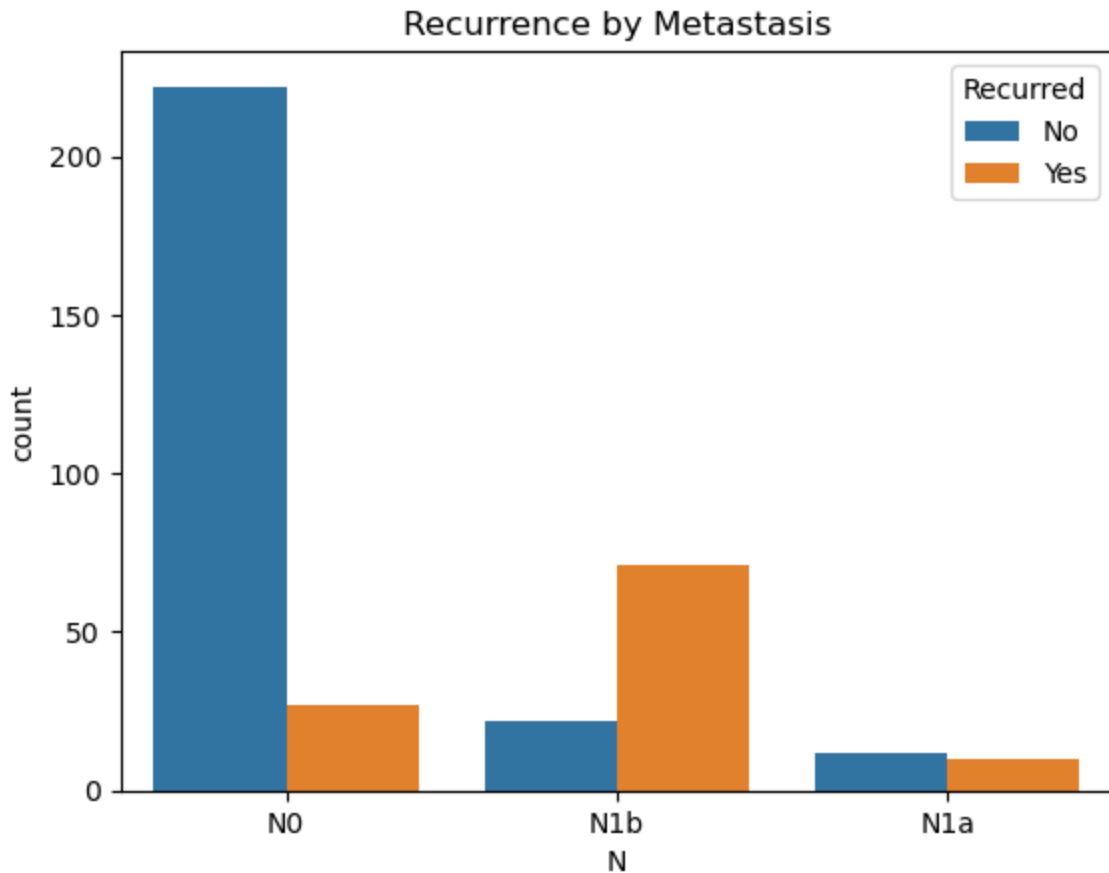
chi2, p, dof, expected = chi2_contingency(ct_m)
print(f'Chi-square: {chi2}, p-value: {p}')

# Graph To Analyzing the effect of Metastasis
sns.countplot(data=patient_data, x='N', hue='Recurred')
plt.title('Recurrence by Metastasis')
plt.show()
```

Recurred	No	Yes
M0	256	90
M1	0	18

Chi-square: 41.41063385710294, p-value: 1.2338437472012865e-10





In [ ]: There **is** a statistically significant relationship between distant metastasis

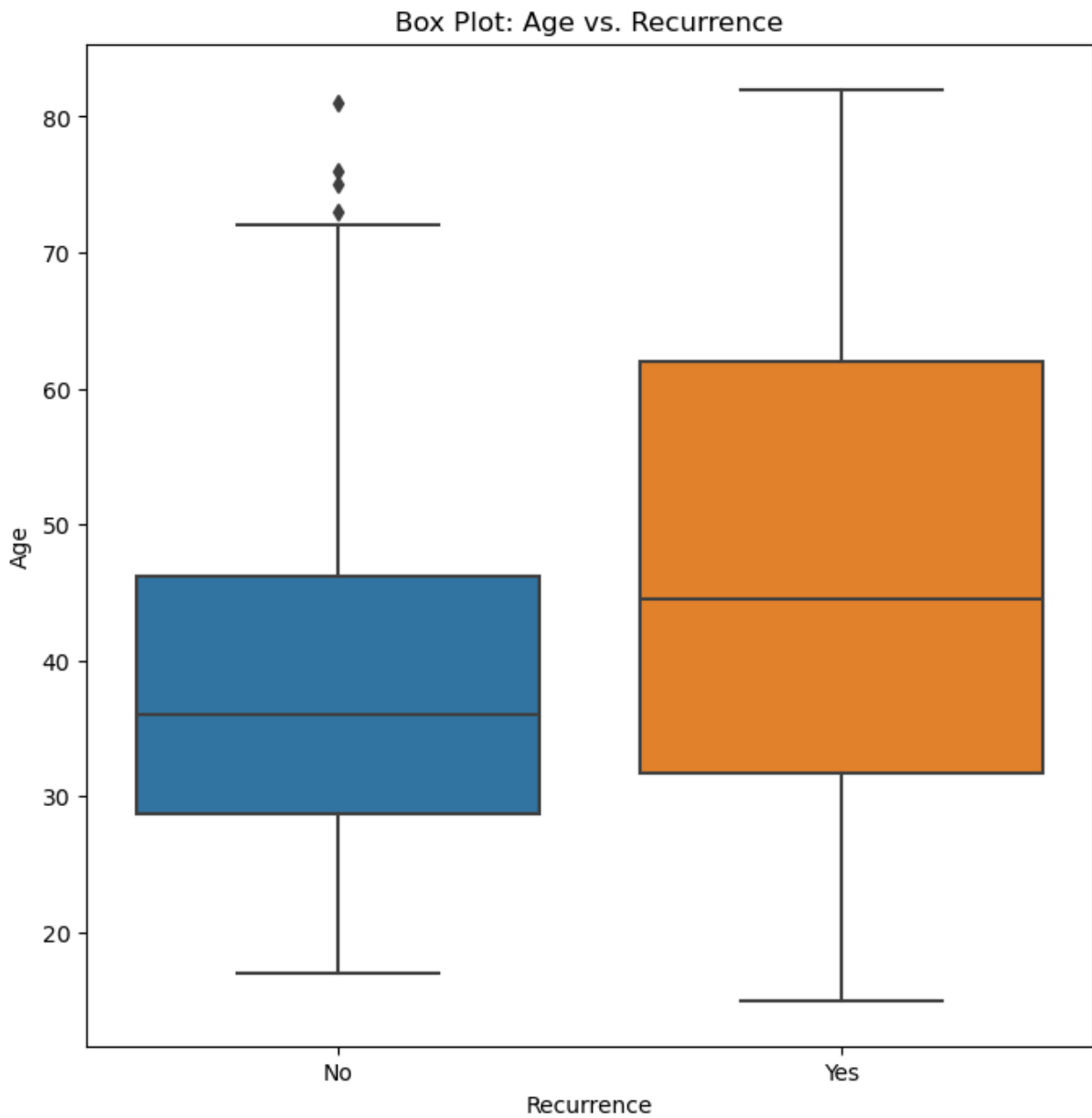
- M1 (metastasis present): Every patient **in** this group experienced recurrence
- M0 (no metastasis): Even though a portion of patients without metastasis

The test strongly indicates that the presence of metastasis (M1) **is** associated with recurrence.

This finding suggests that metastasis **is** a key factor influencing the outcome of thyroid cancer.

In [ ]: All the above Chi-square values show that Tumor size(T), Lymph node involvement

```
In [27]: plt.figure(figsize=(8, 8))
sns.boxplot(x='Recurred', y='Age', data=patient_data)
plt.title('Box Plot: Age vs. Recurrence')
plt.xlabel('Recurrence')
plt.ylabel('Age')
plt.show()
```



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In [ ]: Most of the cases without recurrence fall below 40 years of age. There are f
Most cases with recurrence are of the ages 30 and above, with the biggest nu
```