

# 1007\_\_basic\_\_analysis

January 29, 2023

```
[1]: import pandas as pd
import numpy as np
```

```
[2]: df= pd.read_csv('winequality-red.csv')
df.head()
```

```
[2]:   fixed acidity  volatile acidity  citric acid  residual sugar  chlorides \
0           7.4             0.70         0.00           1.9       0.076
1           7.8             0.88         0.00           2.6       0.098
2           7.8             0.76         0.04           2.3       0.092
3          11.2             0.28         0.56           1.9       0.075
4           7.4             0.70         0.00           1.9       0.076
```

```
   free sulfur dioxide  total sulfur dioxide  density  pH  sulphates \
0              11.0             34.0  0.9978  3.51       0.56
1              25.0             67.0  0.9968  3.20       0.68
2              15.0             54.0  0.9970  3.26       0.65
3              17.0             60.0  0.9980  3.16       0.58
4              11.0             34.0  0.9978  3.51       0.56
```

```
   alcohol  quality
0       9.4        5
1       9.8        5
2       9.8        5
3       9.8        6
4       9.4        5
```

```
[3]: df.dtypes # type check
```

```
[3]: fixed acidity      float64
volatile acidity      float64
citric acid           float64
residual sugar        float64
chlorides             float64
free sulfur dioxide   float64
total sulfur dioxide   float64
density               float64
pH                   float64
```

```

sulphates          float64
alcohol            float64
quality            int64
dtype: object

```

## 1 EDA

```
[4]: df.describe()
```

```

[4]:      fixed acidity  volatile acidity  citric acid  residual sugar  \
count      1599.000000      1599.000000    1599.000000      1599.000000
mean         8.319637         0.527821     0.270976         2.538806
std          1.741096         0.179060     0.194801         1.409928
min           4.600000         0.120000     0.000000         0.900000
25%           7.100000         0.390000     0.090000         1.900000
50%           7.900000         0.520000     0.260000         2.200000
75%           9.200000         0.640000     0.420000         2.600000
max          15.900000         1.580000     1.000000        15.500000

      chlorides  free sulfur dioxide  total sulfur dioxide  density  \
count      1599.000000      1599.000000      1599.000000    1599.000000
mean         0.087467         15.874922         46.467792     0.996747
std          0.047065         10.460157         32.895324     0.001887
min           0.012000         1.000000         6.000000     0.990070
25%           0.070000         7.000000         22.000000     0.995600
50%           0.079000        14.000000         38.000000     0.996750
75%           0.090000        21.000000         62.000000     0.997835
max           0.611000        72.000000        289.000000     1.003690

      pH  sulphates  alcohol  quality
count      1599.000000    1599.000000    1599.000000    1599.000000
mean         3.311113     0.658149     10.422983     5.636023
std          0.154386     0.169507     1.065668     0.807569
min           2.740000     0.330000     8.400000     3.000000
25%           3.210000     0.550000     9.500000     5.000000
50%           3.310000     0.620000    10.200000     6.000000
75%           3.400000     0.730000    11.100000     6.000000
max           4.010000     2.000000    14.900000     8.000000

```

### 1.1 Filtering

```
[5]: df['fixed acidity'] # Return as Series when single []
```

```

[5]: 0      7.4
     1      7.8
     2      7.8
     3     11.2

```

```

4          7.4
...
1594       6.2
1595       5.9
1596       6.3
1597       5.9
1598       6.0
Name: fixed acidity, Length: 1599, dtype: float64

```

```
[6]: df[['fixed acidity']] # two [], return as dataframe or list of list
```

```
[6]:      fixed acidity
0          7.4
1          7.8
2          7.8
3         11.2
4          7.4
...
1594       6.2
1595       5.9
1596       6.3
1597       5.9
1598       6.0
```

[1599 rows x 1 columns]

```
[7]: # Showing mutiple "COLUMN" as per necessary
df [['fixed acidity', 'density', 'quality']] # misddle value shown as ...
↳ later we will see how to see as full
```

```
[7]:      fixed acidity  density  quality
0          7.4  0.99780      5
1          7.8  0.99680      5
2          7.8  0.99700      5
3         11.2  0.99800      6
4          7.4  0.99780      5
...
1594       6.2  0.99490      5
1595       5.9  0.99512      6
1596       6.3  0.99574      6
1597       5.9  0.99547      5
1598       6.0  0.99549      6
```

[1599 rows x 3 columns]

```
[8]: # Test1: Check Spefic "ROW" as required or with any condition
```

```
df[df['fixed acidity'] > 9] # Check when fixed acidity > 9; syntax df inside ↪df
```

```
[8]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides \
3	11.2	0.28	0.56	1.9	0.075
56	10.2	0.42	0.57	3.4	0.070
68	9.3	0.32	0.57	2.0	0.074
74	9.7	0.32	0.54	2.5	0.094
88	9.3	0.39	0.44	2.1	0.107
...	...	...	...	...	...
1470	10.0	0.69	0.11	1.4	0.084
1474	9.9	0.50	0.50	13.8	0.205
1476	9.9	0.50	0.50	13.8	0.205
1543	11.1	0.44	0.42	2.2	0.064
1548	11.2	0.40	0.50	2.0	0.099

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates \
3	17.0	60.0	0.99800	3.16	0.58
56	4.0	10.0	0.99710	3.04	0.63
68	27.0	65.0	0.99690	3.28	0.79
74	28.0	83.0	0.99840	3.28	0.82
88	34.0	125.0	0.99780	3.14	1.22
...	...	...	...	...	...
1470	8.0	24.0	0.99578	2.88	0.47
1474	48.0	82.0	1.00242	3.16	0.75
1476	48.0	82.0	1.00242	3.16	0.75
1543	14.0	19.0	0.99758	3.25	0.57
1548	19.0	50.0	0.99783	3.10	0.58

	alcohol	quality
3	9.8	6
56	9.6	5
68	10.7	5
74	9.6	5
88	9.5	5
...	...	...
1470	9.7	5
1474	8.8	5
1476	8.8	5
1543	10.4	6
1548	10.4	5

[441 rows x 12 columns]

```
[9]: # Test 2
df[(df['fixed acidity'] > 9) & (df['citric acid'] > 0.5)] # Multiple ↪
condition over row, more condition can be added inside parenthesis
```

```
[9]:      fixed acidity  volatile acidity  citric acid  residual sugar  chlorides  \
3          11.2          0.28          0.56          1.9          0.075
56         10.2          0.42          0.57          3.4          0.070
68          9.3          0.32          0.57          2.0          0.074
74          9.7          0.32          0.54          2.5          0.094
151         9.2          0.52          1.00          3.4          0.610
...
1221        10.9          0.32          0.52          1.8          0.132
1319         9.1          0.76          0.68          1.7          0.414
1414        10.0          0.32          0.59          2.2          0.077
1416        10.0          0.32          0.59          2.2          0.077
1454        11.7          0.45          0.63          2.2          0.073
```

```
      free sulfur dioxide  total sulfur dioxide  density  pH  sulphates  \
3          17.0          60.0  0.99800  3.16          0.58
56           4.0          10.0  0.99710  3.04          0.63
68          27.0          65.0  0.99690  3.28          0.79
74          28.0          83.0  0.99840  3.28          0.82
151         32.0          69.0  0.99960  2.74          2.00
...
1221          17.0          44.0  0.99734  3.28          0.77
1319          18.0          64.0  0.99652  2.90          1.33
1414           3.0          15.0  0.99940  3.20          0.78
1416           3.0          15.0  0.99940  3.20          0.78
1454           7.0          23.0  0.99974  3.21          0.69
```

```
      alcohol  quality
3          9.8        6
56         9.6        5
68        10.7        5
74         9.6        5
151        9.4        4
...
1221        11.5        6
1319         9.1        6
1414         9.6        5
1416         9.6        5
1454        10.9        6
```

[142 rows x 12 columns]

```
[10]: df [(df['fixed acidity'] > 9) & (df['citric acid'] > 0.5) & (df['pH'] >=3)] #L
↳ Test 3
```

```
[10]:      fixed acidity  volatile acidity  citric acid  residual sugar  chlorides  \
3          11.2          0.28          0.56          1.9          0.075
56         10.2          0.42          0.57          3.4          0.070
```

68	9.3	0.32	0.57	2.0	0.074
74	9.7	0.32	0.54	2.5	0.094
197	11.5	0.30	0.60	2.0	0.067
...	...	...	...	...	...
1220	10.9	0.32	0.52	1.8	0.132
1221	10.9	0.32	0.52	1.8	0.132
1414	10.0	0.32	0.59	2.2	0.077
1416	10.0	0.32	0.59	2.2	0.077
1454	11.7	0.45	0.63	2.2	0.073

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates \
3	17.0	60.0	0.99800	3.16	0.58
56	4.0	10.0	0.99710	3.04	0.63
68	27.0	65.0	0.99690	3.28	0.79
74	28.0	83.0	0.99840	3.28	0.82
197	12.0	27.0	0.99810	3.11	0.97
...	...	...	...	...	...
1220	17.0	44.0	0.99734	3.28	0.77
1221	17.0	44.0	0.99734	3.28	0.77
1414	3.0	15.0	0.99940	3.20	0.78
1416	3.0	15.0	0.99940	3.20	0.78
1454	7.0	23.0	0.99974	3.21	0.69

	alcohol	quality
3	9.8	6
56	9.6	5
68	10.7	5
74	9.6	5
197	10.1	6
...	...	...
1220	11.5	6
1221	11.5	6
1414	9.6	5
1416	9.6	5
1454	10.9	6

[131 rows x 12 columns]

```
[11]: # In case of row wise selection, its returning all
df [(df['fixed acidity'] > 9) | (df['citric acid'] > 0.5)] # Or Function
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides \
3	11.2	0.28	0.56	1.9	0.075
16	8.5	0.28	0.56	1.8	0.092
19	7.9	0.32	0.51	1.8	0.341
47	8.7	0.29	0.52	1.6	0.113
56	10.2	0.42	0.57	3.4	0.070

...	...	...	...	...	...	...
1548	11.2	0.40	0.50	2.0	0.099	
1566	6.7	0.16	0.64	2.1	0.059	
1570	6.4	0.36	0.53	2.2	0.230	
1574	5.6	0.31	0.78	13.9	0.074	
1576	8.0	0.30	0.63	1.6	0.081	

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	\
3	17.0	60.0	0.99800	3.16	0.58	
16	35.0	103.0	0.99690	3.30	0.75	
19	17.0	56.0	0.99690	3.04	1.08	
47	12.0	37.0	0.99690	3.25	0.58	
56	4.0	10.0	0.99710	3.04	0.63	
...	...	...	...	...	...	...
1548	19.0	50.0	0.99783	3.10	0.58	
1566	24.0	52.0	0.99494	3.34	0.71	
1570	19.0	35.0	0.99340	3.37	0.93	
1574	23.0	92.0	0.99677	3.39	0.48	
1576	16.0	29.0	0.99588	3.30	0.78	

	alcohol	quality
3	9.8	6
16	10.5	7
19	9.2	6
47	9.5	5
56	9.6	5
...	...	...
1548	10.4	5
1566	11.2	6
1570	12.4	6
1574	10.5	6
1576	10.8	6

[489 rows x 12 columns]

```
[12]: # Row Wise File - with only selective columns as per conditio
df.loc[df['fixed acidity'] == 9.2, ['fixed acidity', 'citric acid', 'pH']] #
    ↳syntax: loc means, we are locationg, then conditio as ==9.2 & name of col as
    ↳want to show
# as per given conditio, the total retunr number / qnt not show, we can see it
    ↳as next code
```

```
[12]:    fixed acidity  citric acid    pH
151          9.2         1.00  2.74
457          9.2         0.21  3.28
460          9.2         0.52  3.35
491          9.2         0.50  3.34
```

524	9.2	0.49	3.23
540	9.2	0.24	3.26
614	9.2	0.18	2.87
691	9.2	0.24	3.48
741	9.2	0.24	3.21
765	9.2	0.10	3.31
880	9.2	0.18	3.15
905	9.2	0.20	3.23
1093	9.2	0.36	3.33
1170	9.2	0.34	3.20
1225	9.2	0.23	3.15
1360	9.2	0.31	3.24

```
[13]: data = df.loc[df['fixed acidity'] == 9.2, ['fixed acidity', 'citric acid',
↪ 'pH']] # just bring the previous code in a var and see the shape
data.shape
```

```
[13]: (16, 3)
```

```
[14]: df.tail(7)
```

```
[14]:
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides \
1592	6.3	0.510	0.13	2.3	0.076
1593	6.8	0.620	0.08	1.9	0.068
1594	6.2	0.600	0.08	2.0	0.090
1595	5.9	0.550	0.10	2.2	0.062
1596	6.3	0.510	0.13	2.3	0.076
1597	5.9	0.645	0.12	2.0	0.075
1598	6.0	0.310	0.47	3.6	0.067

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates \
1592	29.0	40.0	0.99574	3.42	0.75
1593	28.0	38.0	0.99651	3.42	0.82
1594	32.0	44.0	0.99490	3.45	0.58
1595	39.0	51.0	0.99512	3.52	0.76
1596	29.0	40.0	0.99574	3.42	0.75
1597	32.0	44.0	0.99547	3.57	0.71
1598	18.0	42.0	0.99549	3.39	0.66

	alcohol	quality
1592	11.0	6
1593	9.5	6
1594	10.5	5
1595	11.2	6
1596	11.0	6
1597	10.2	5
1598	11.0	6



```
[15]: df[5:11] # want to see from index 5 to 10
```

```
[15]:      fixed acidity  volatile acidity  citric acid  residual sugar  chlorides \
5          7.4          0.66          0.00          1.8          0.075
6          7.9          0.60          0.06          1.6          0.069
7          7.3          0.65          0.00          1.2          0.065
8          7.8          0.58          0.02          2.0          0.073
9          7.5          0.50          0.36          6.1          0.071
10         6.7          0.58          0.08          1.8          0.097

      free sulfur dioxide  total sulfur dioxide  density  pH  sulphates \
5          13.0          40.0  0.9978  3.51          0.56
6          15.0          59.0  0.9964  3.30          0.46
7          15.0          21.0  0.9946  3.39          0.47
8           9.0          18.0  0.9968  3.36          0.57
9          17.0         102.0  0.9978  3.35          0.80
10         15.0          65.0  0.9959  3.28          0.54

      alcohol  quality
5          9.4        5
6          9.4        5
7         10.0        7
8          9.5        7
9         10.5        5
10         9.2        5
```

View Full Rows & Column, middle ... will not be shown

```
[16]: # Required codes for see full, but its not recommended actually or no needed
# pd.set_option('display.max_rows',None)
# pd.set_option('display.max_columns', None)
# df
```

## 1.2 Processed File Export

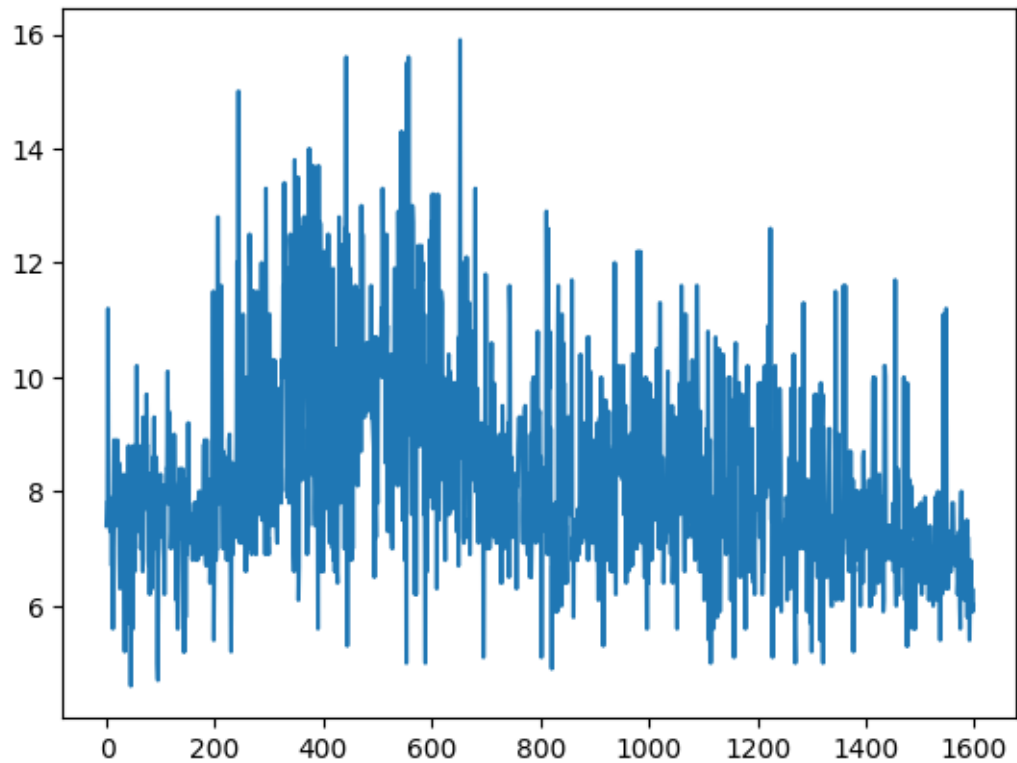
```
[17]: # x = df [(df['fixed acidity'] > 9) | (df['citric acid'] > 0.5)] # processed
      ↪ file stored into a variable file as x
# x.to_csv('1007_processedfile.csv')
```

## 2 Visualization

Very basic or Ordinary Visualization

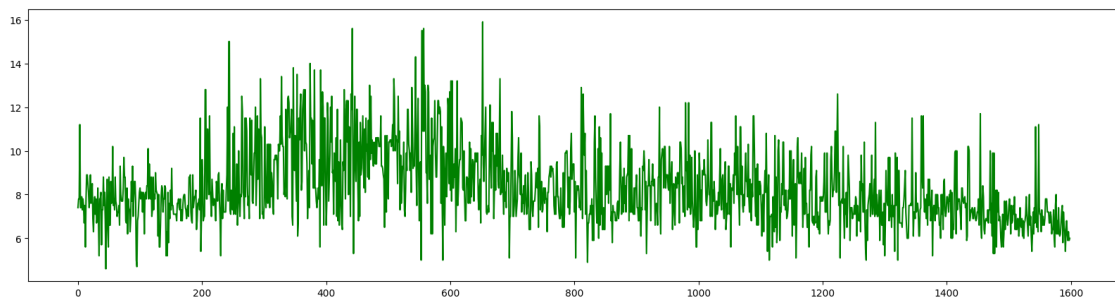
```
[18]: # Plot with a single columns
df['fixed acidity'].plot.line()
```

```
[18]: <AxesSubplot:>
```



```
[19]: df['fixed acidity'].plot(figsize=(20,5), color='green')
```

```
[19]: <AxesSubplot:>
```



## 3 Better Visualization

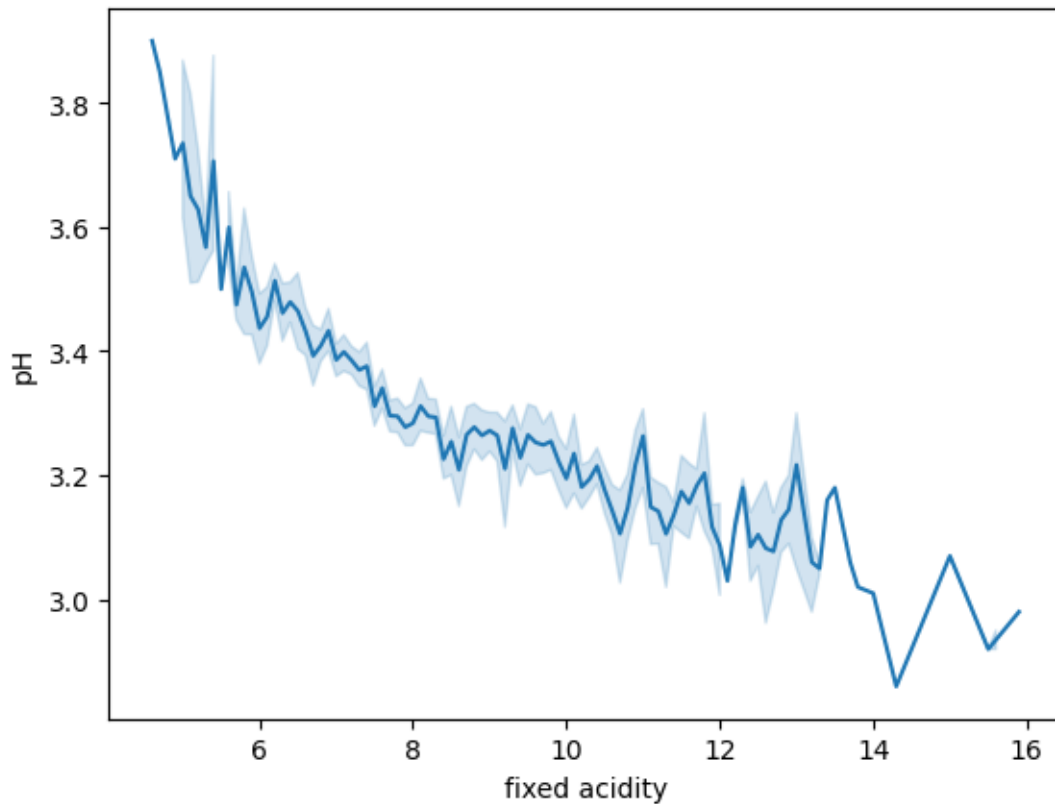
### 3.1 Seaborn

```
[20]: import seaborn as sns
```

- There is Negative Relationship between Acidity & Ph, lets visualize the relationship between these two
- Sns lineplot must require independent & dependent variable, here x is independent & y is dependent over x

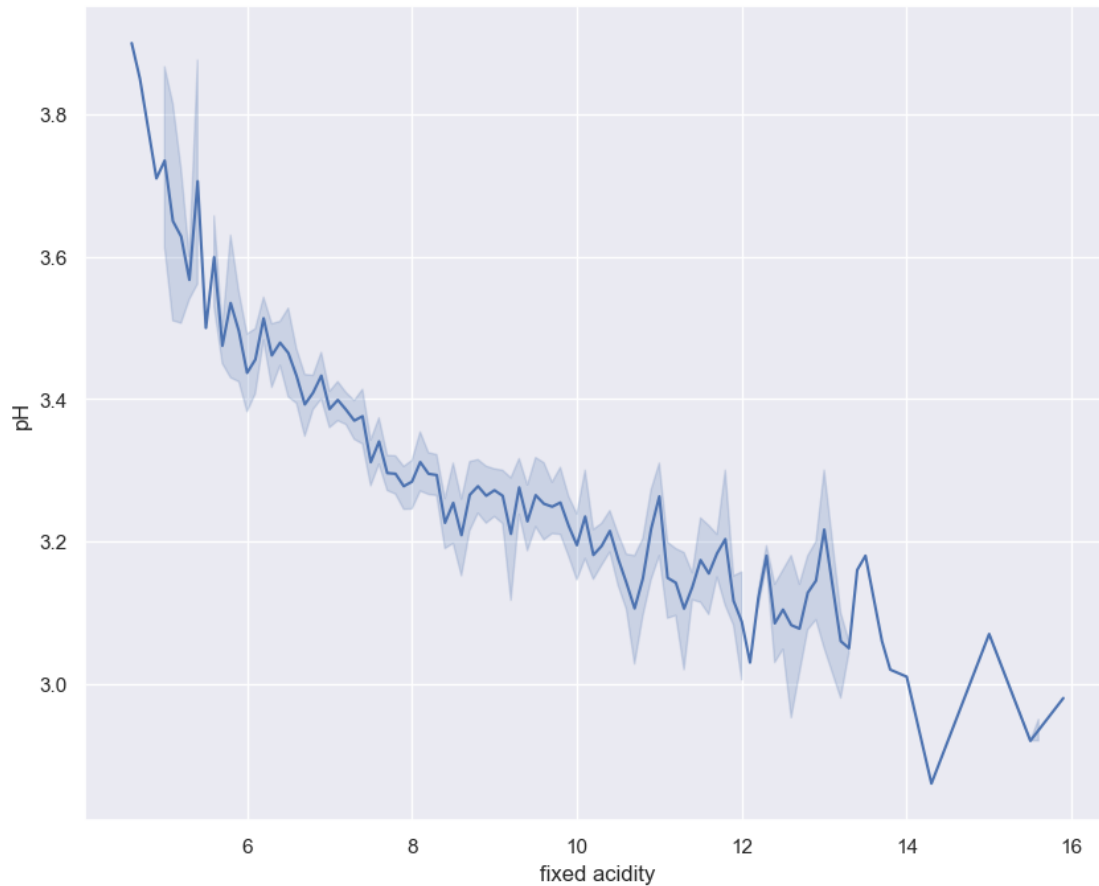
```
[21]: sns.lineplot(data=df, x='fixed acidity', y='pH')
```

```
[21]: <AxesSubplot:xlabel='fixed acidity', ylabel='pH'>
```



```
[32]: # Figure Size Customization
sns.set(rc={'figure.figsize':(10, 8)})
sns.lineplot(data=df, x='fixed acidity', y='pH')
# The shadow represent the deviation / standard deviation / error rate or how
↳ much deviation is there
```

```
[32]: <AxesSubplot:xlabel='fixed acidity', ylabel='pH'>
```

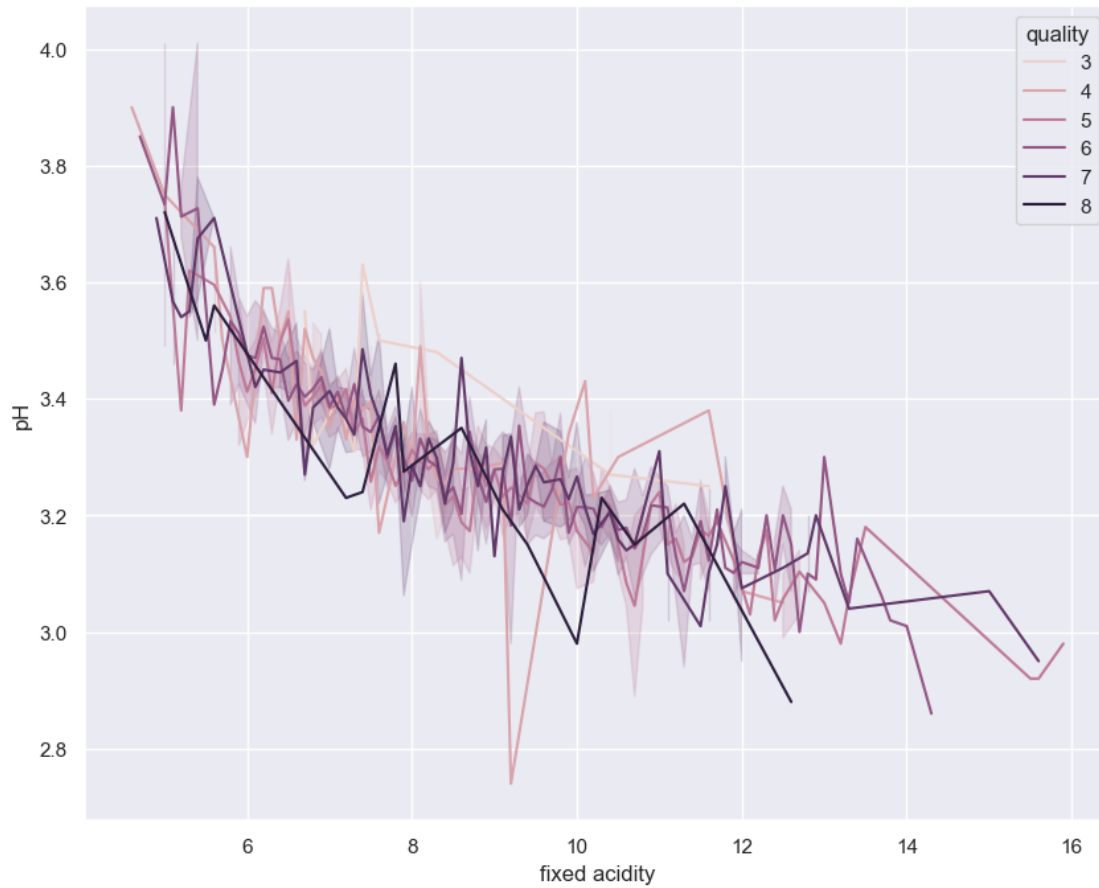


### 3.1.1 Categorical Data Visualization

- Male / Female type
- In this dataset, quality is categorical data
- Event
- By the category you want to represent data, that must be passed in “hue”
- Checking Facebook traffic for day & night

```
[34]: sns.lineplot(data=df, x='fixed acidity', y='pH', hue='quality') # We have 6
      ↪ type of category
```

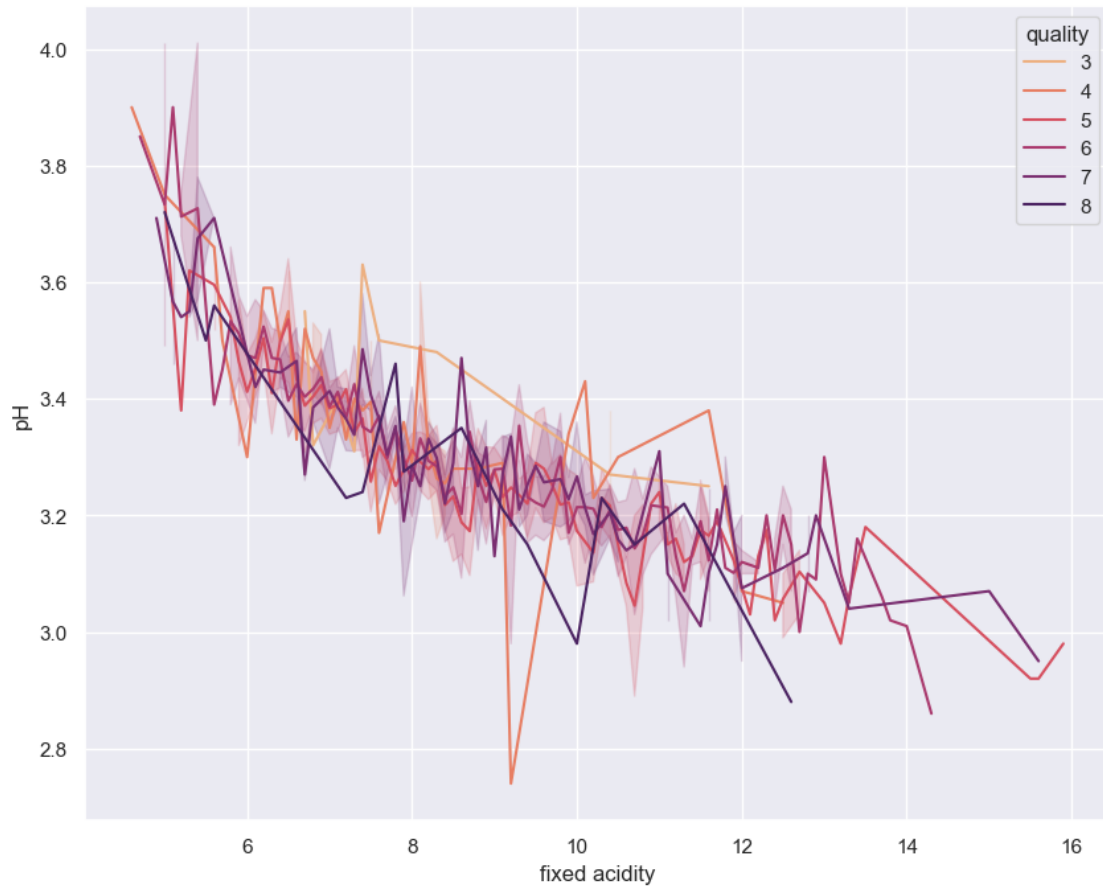
```
[34]: <AxesSubplot:xlabel='fixed acidity', ylabel='pH'>
```



### 3.2 Color Customization

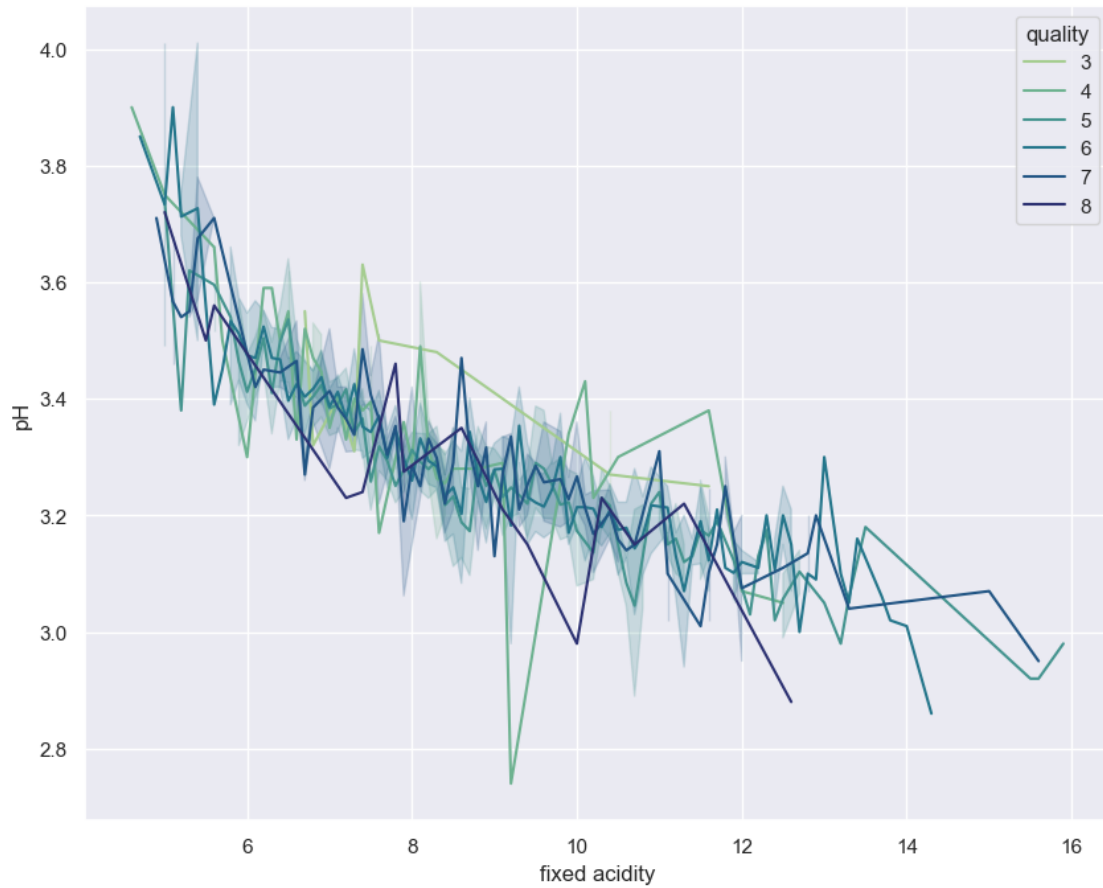
```
[36]: p = sns.color_palette("flare", as_cmap=True) # customized color palette stored
      ↪ in p & then pass to following code for color customization
      sns.lineplot(data=df, x='fixed acidity', y='pH', hue='quality', palette=p)
```

```
[36]: <AxesSubplot:xlabel='fixed acidity', ylabel='pH'>
```



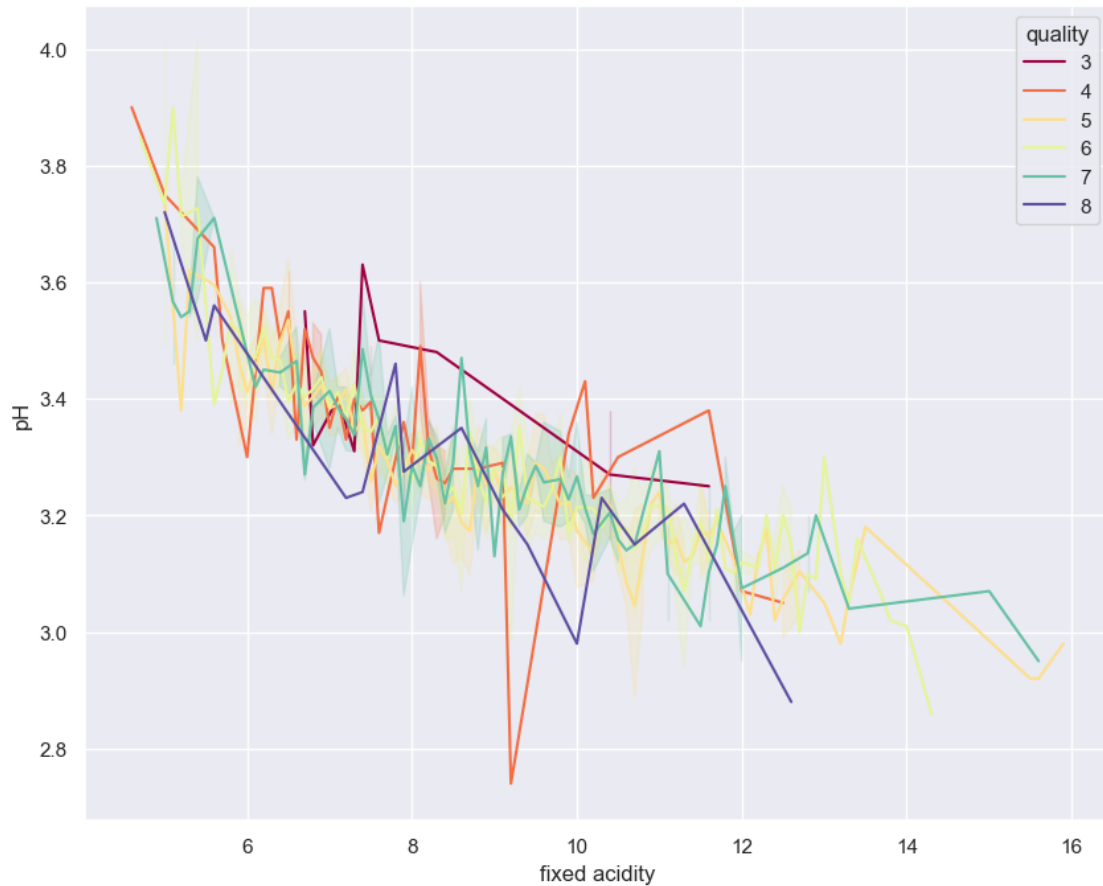
```
[38]: p = sns.color_palette("crest", as_cmap=True) # Another color palette
      sns.lineplot(data=df, x='fixed acidity', y='pH', hue='quality', palette=p)
```

```
[38]: <AxesSubplot:xlabel='fixed acidity', ylabel='pH'>
```



```
[39]: p = sns.color_palette("Spectral", as_cmap=True) # Another color palette
      sns.lineplot(data=df, x='fixed acidity', y='pH', hue='quality', palette=p)
```

```
[39]: <AxesSubplot:xlabel='fixed acidity', ylabel='pH'>
```



### 3.3 Creating New Dataset for Visualization

```
[52]: df2 = [[ 'Alice', 30, 'Male', 55 ], ['Bobe', 17, 'Female', 25 ], ['Jeba', 11, 'Female', 12], ['Tom', 45, 'Male', 72], ['Rita', 21, 'Female', 35], ['Jackline', 51, 'Female', 65], ['Peter', 55, 'Male', 72]]
df2 = pd.DataFrame(df2, columns=['Name', 'Age', 'Gender', 'Weight in KG']) # Column declaration
df2
```

```
[52]:
```

	Name	Age	Gender	Weight in KG
0	Alice	30	Male	55
1	Bobe	17	Female	25
2	Jeba	11	Female	12
3	Tom	45	Male	72
4	Rita	21	Female	35
5	Jackline	51	Female	65
6	Peter	55	Male	72

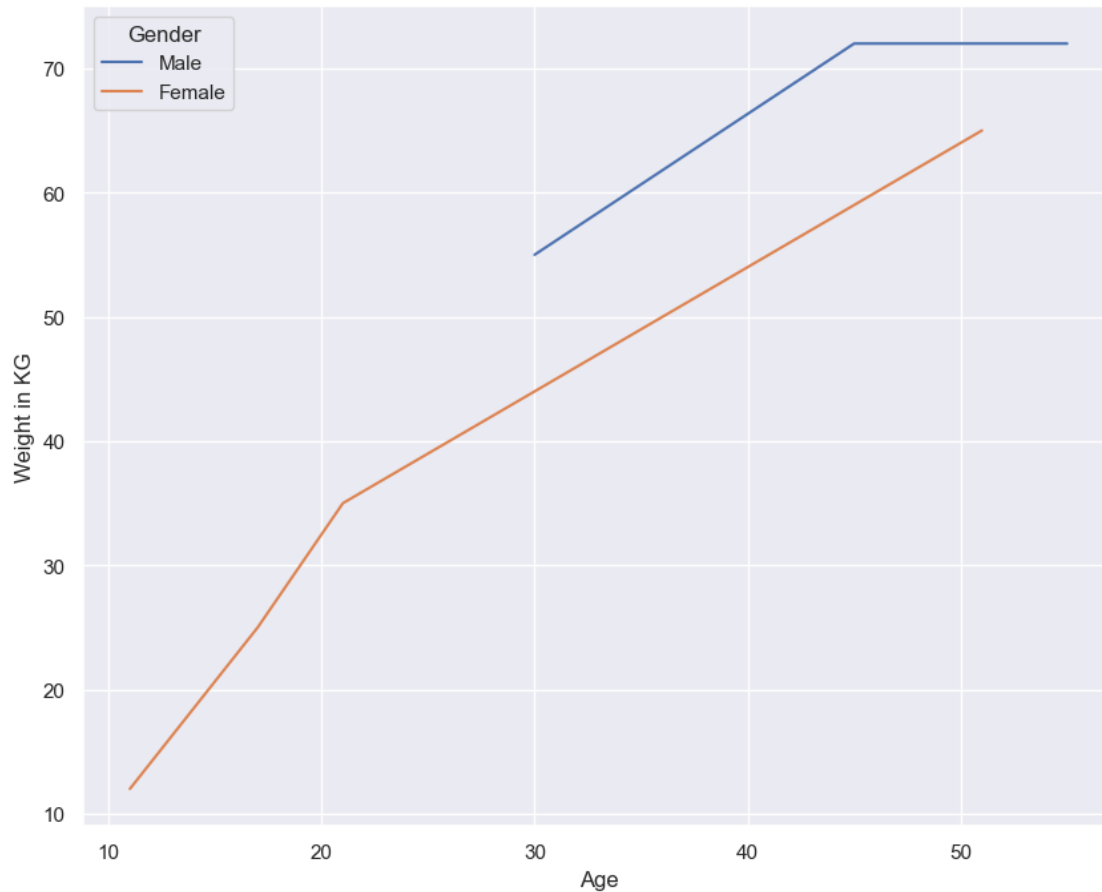
- We will see a visualization for Age & Weight in respect of Gender



- More Application: facebook traffic day & night

```
[55]: sns.lineplot(data=df2, x='Age', y='Weight in KG', hue='Gender') # hue param is useful when divided by category
```

```
[55]: <AxesSubplot:xlabel='Age', ylabel='Weight in KG'>
```



### 3.4 Creating Test Data

```
[60]: tips = [
    [16.99, 1.01, 'Female', 'No', 'Sun', 'Dinner', 2],
    [10.34, 1.66, 'Male', 'No', 'Sun', 'Dinner', 3],
    [21.01, 3.50, 'Male', 'No', 'Sun', 'Dinner', 3],
    [23.65, 3.50, 'Male', 'No', 'Mon', 'Dinner', 4],
    [24.59, 3.61, 'Female', 'No', 'Sun', 'Dinner', 3],
]
tips = pd.DataFrame(data=tips, columns=['Total Bill', 'Tips', 'Gender', 'Smoker', 'Day', 'Time', 'Size'])
tips
```

```
[60]:
```

	Total Bill	Tips	Gender	Smoker	Day	Time	Size
0	16.99	1.01	Female	No	Sun	Dinner	2
1	10.34	1.66	Male	No	Sun	Dinner	3
2	21.01	3.50	Male	No	Sun	Dinner	3
3	23.65	3.50	Male	No	Mon	Dinner	4
4	24.59	3.61	Female	No	Sun	Dinner	3

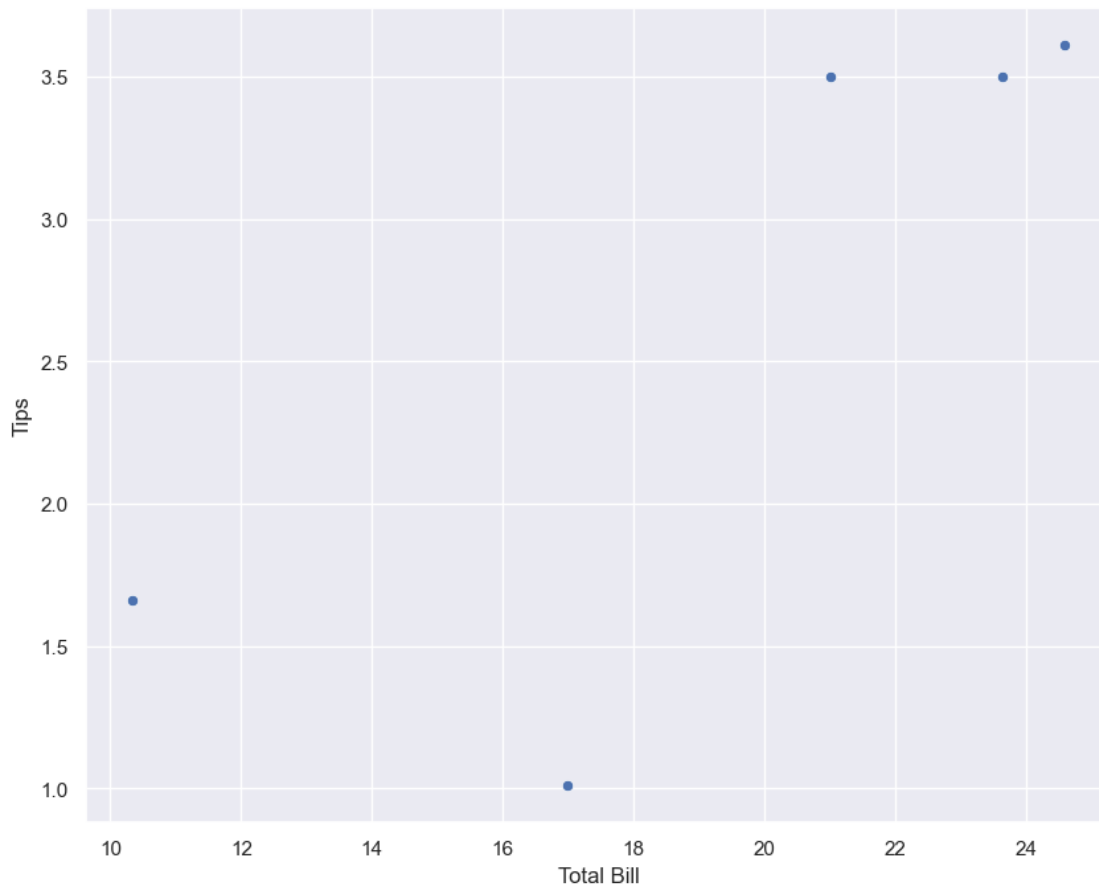
```
[64]: # Alternative Way for Dataset creation with seperate col
tips2 = [
    [16.99, 1.01, 'Female', 'No', 'Sun', 'Dinner', 2],
    [10.34, 1.66, 'Male', 'No', 'Sun', 'Dinner', 3],
    [21.01, 3.50, 'Male', 'No', 'Sun', 'Dinner', 3],
    [23.65, 3.50, 'Male', 'No', 'Mon', 'Dinner', 4],
    [24.59, 3.61, 'Female', 'No', 'Sun', 'Dinner', 3],
]
c = ['Total Bill', 'Tips', 'Gender', 'Smoker', 'Day', 'Time', 'Size'] # Just_
    ↳ keep a column var as c & keep the col name here
tips2 = pd.DataFrame(data=tips, columns= c ) # Pass the c here
tips2
```

```
[64]:
```

	Total Bill	Tips	Gender	Smoker	Day	Time	Size
0	16.99	1.01	Female	No	Sun	Dinner	2
1	10.34	1.66	Male	No	Sun	Dinner	3
2	21.01	3.50	Male	No	Sun	Dinner	3
3	23.65	3.50	Male	No	Mon	Dinner	4
4	24.59	3.61	Female	No	Sun	Dinner	3

```
[65]: sns.scatterplot(data=tips, x='Total Bill', y='Tips')
```

```
[65]: <AxesSubplot:xlabel='Total Bill', ylabel='Tips'>
```



```
[71]: tips = pd.read_csv('tips.csv')
      tips.head()
```

```
[71]:   total_bill   tip     sex smoker  day    time  size
0      16.99   1.01  Female     No   Sun  Dinner     2
1      10.34   1.66    Male     No   Sun  Dinner     3
2      21.01   3.50    Male     No   Sun  Dinner     3
3      23.68   3.31    Male     No   Sun  Dinner     2
4      24.59   3.61  Female     No   Sun  Dinner     4
```

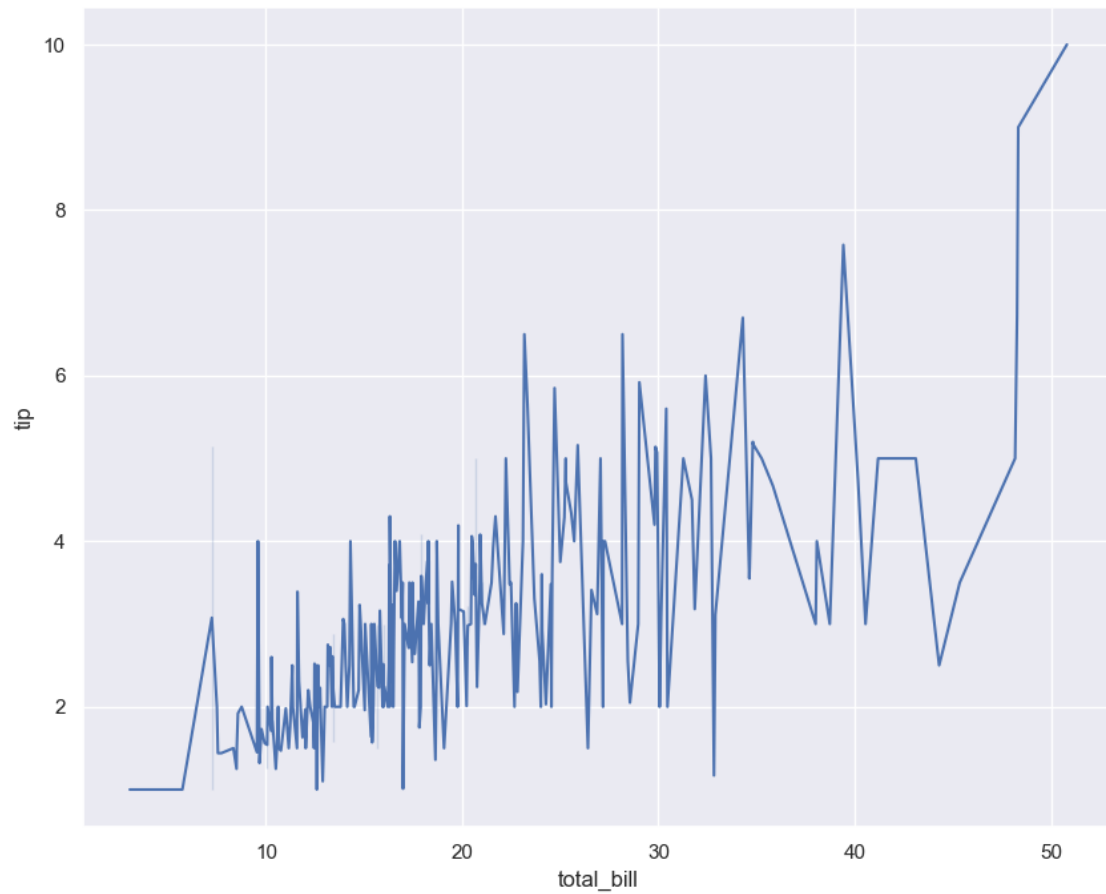
```
[74]: tips.shape
```

```
[74]: (244, 7)
```

### 3.4.1 Line Plot

```
[75]: sns.lineplot(data=tips, x='total_bill', y='tip')
```

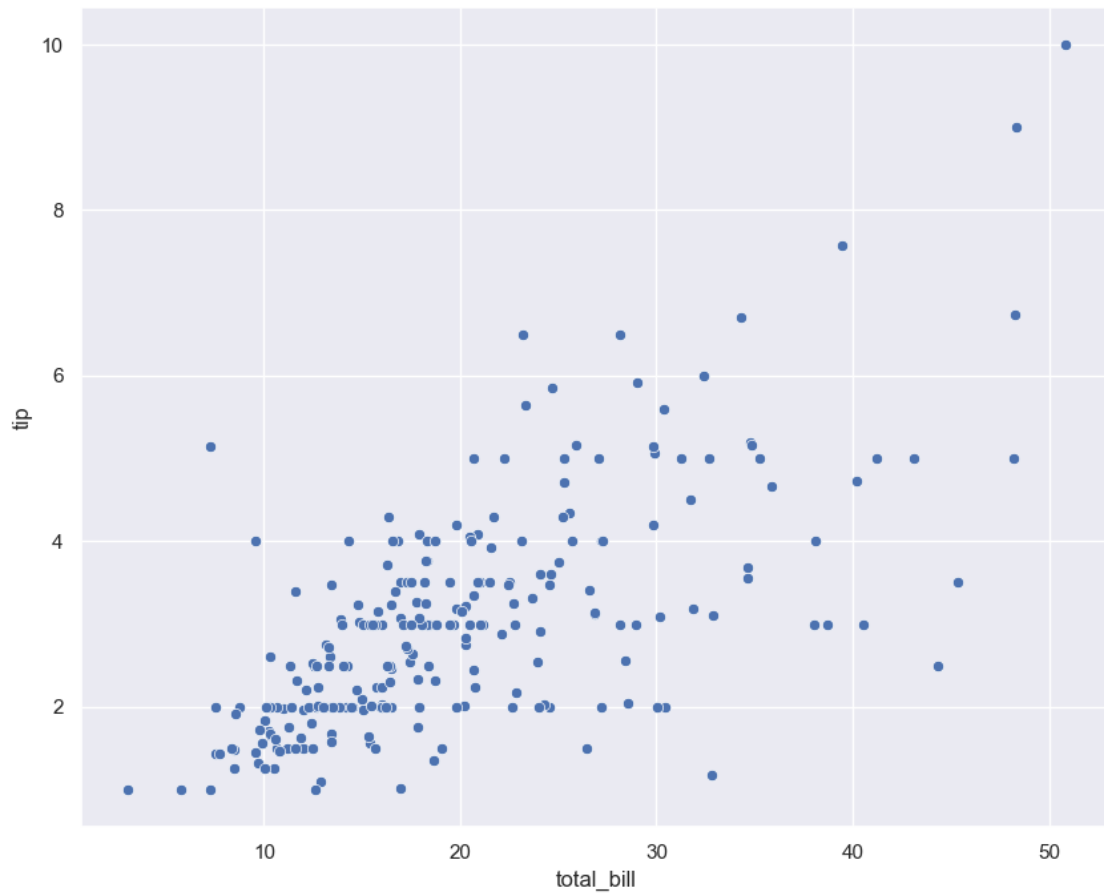
```
[75]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



### 3.4.2 Scatter plot

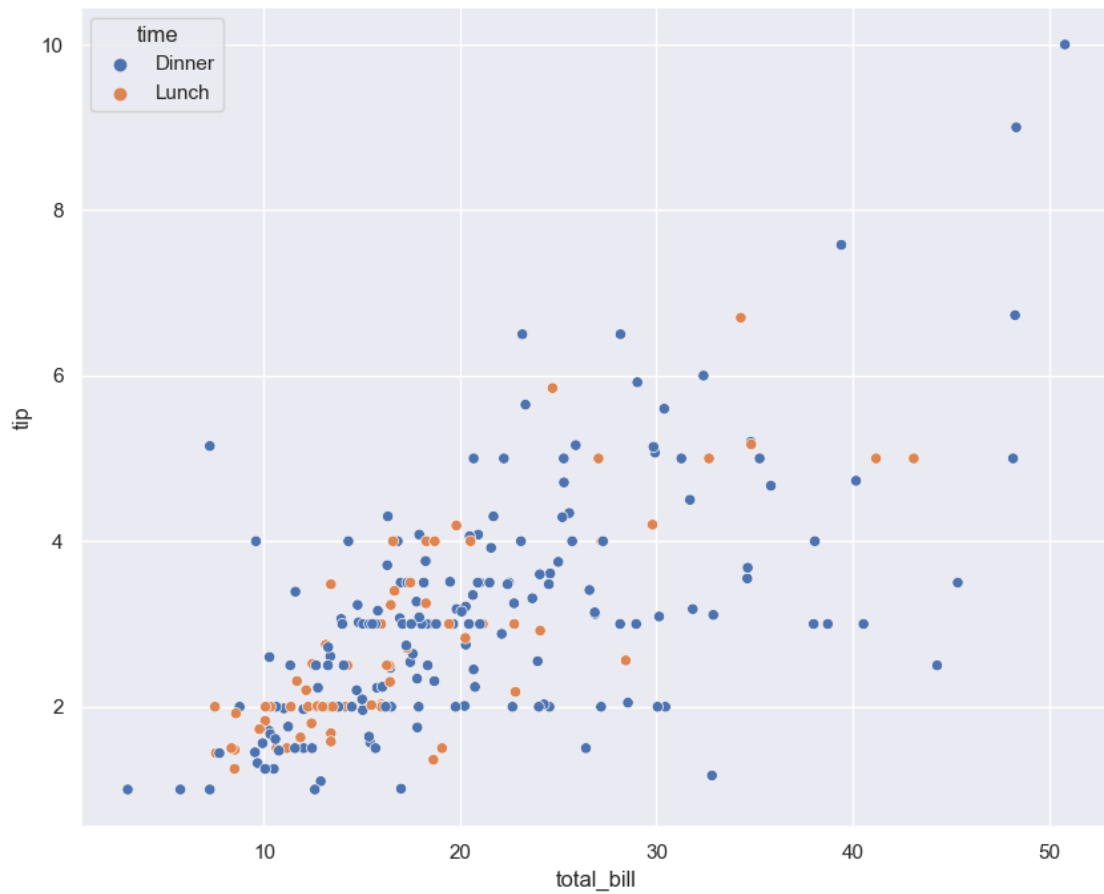
```
[73]: sns.scatterplot(data=tips, x='total_bill', y='tip')
```

```
[73]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



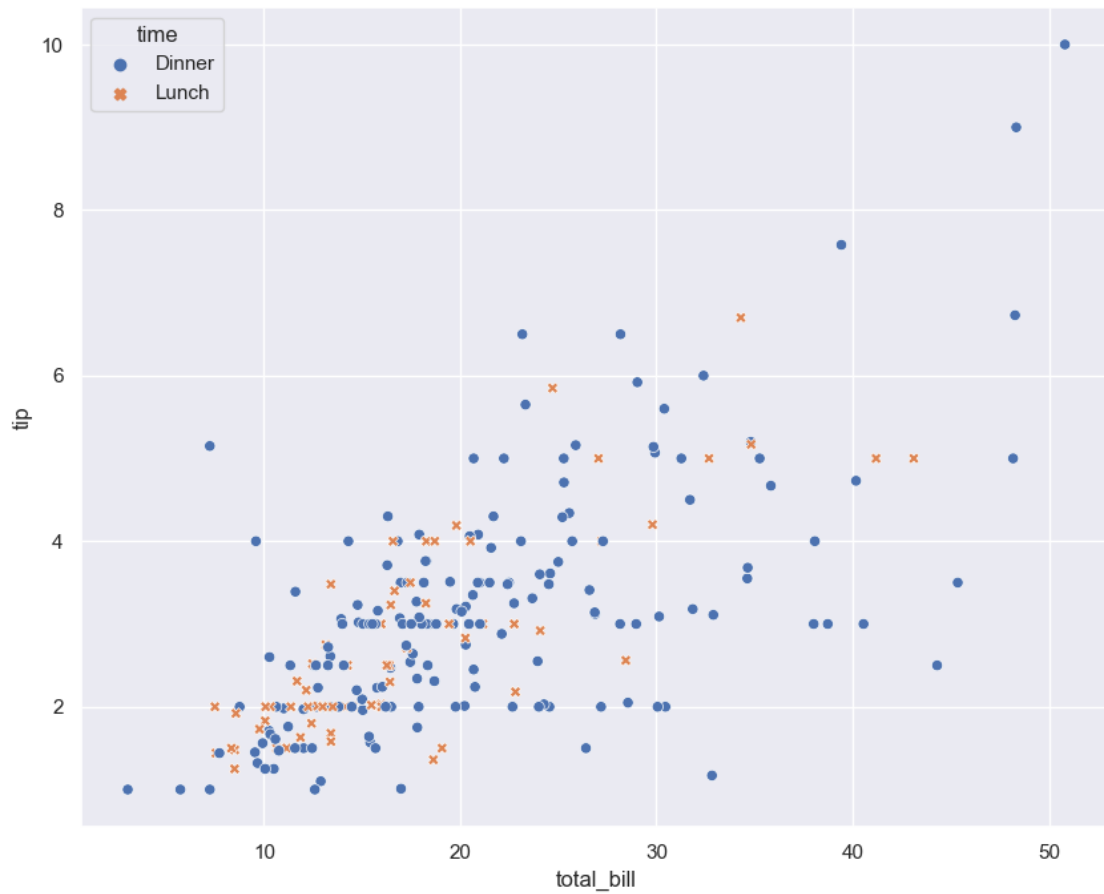
```
[76]: sns.scatterplot(data=tips, x='total_bill', y='tip', hue='time')
```

```
[76]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



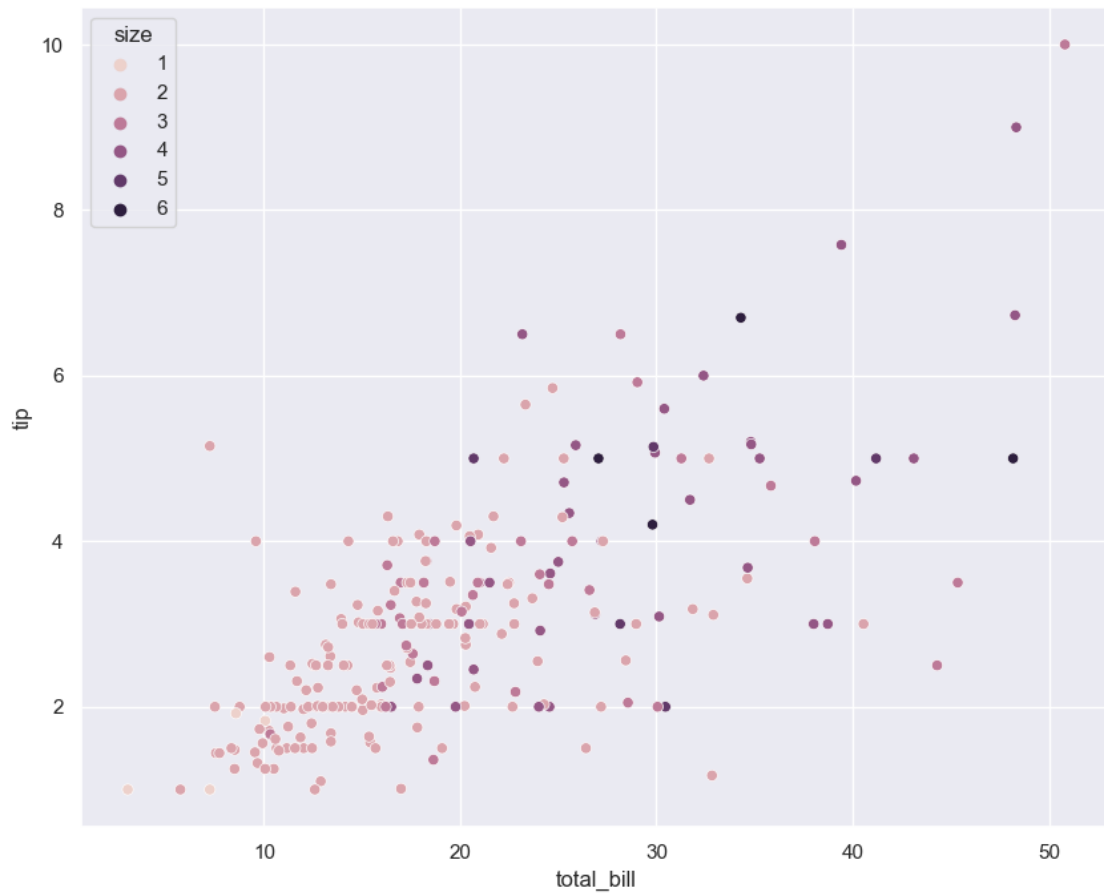
```
[77]: sns.scatterplot(data=tips, x='total_bill', y='tip', hue='time', style='time')
```

```
[77]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



```
[78]: sns.scatterplot(data=tips, x='total_bill', y='tip', hue='size')
```

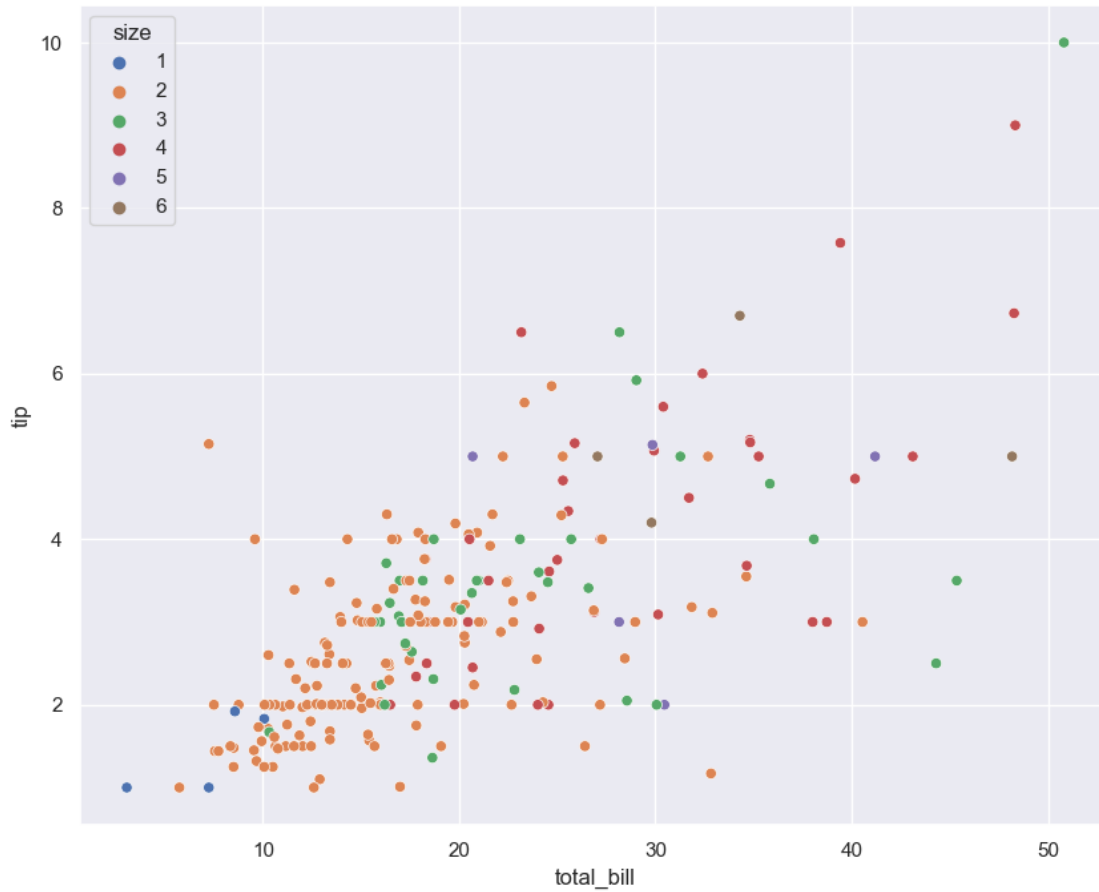
```
[78]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



```
[79]: sns.scatterplot(data=tips, x='total_bill', y='tip', hue='size', palette="deep")
```

```
[79]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



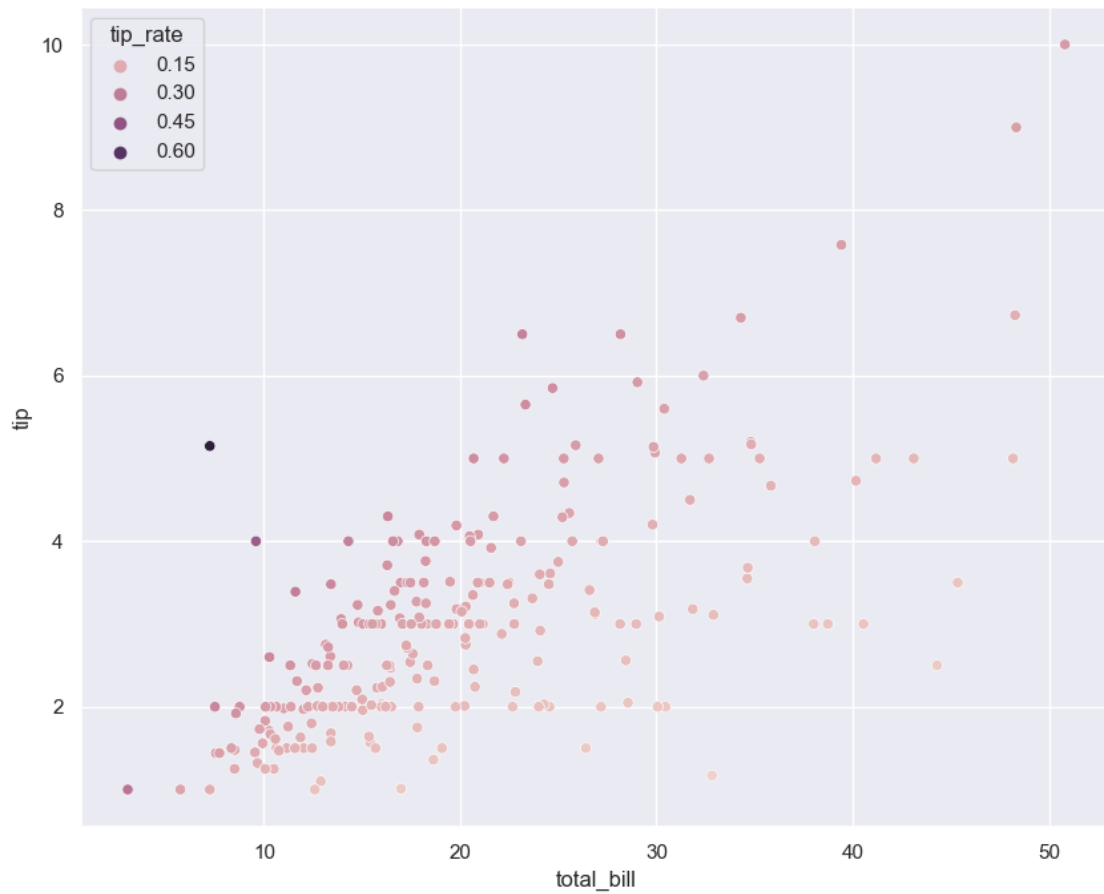


### 3.4.3 Tip rate

- If there are a large number of unique numeric values, the legend will show a representative, evenly-spaced set:

```
[84]: tip_rate = tips.eval("tip / total_bill").rename("tip_rate")
      sns.scatterplot(data=tips, x="total_bill", y="tip", hue=tip_rate)
```

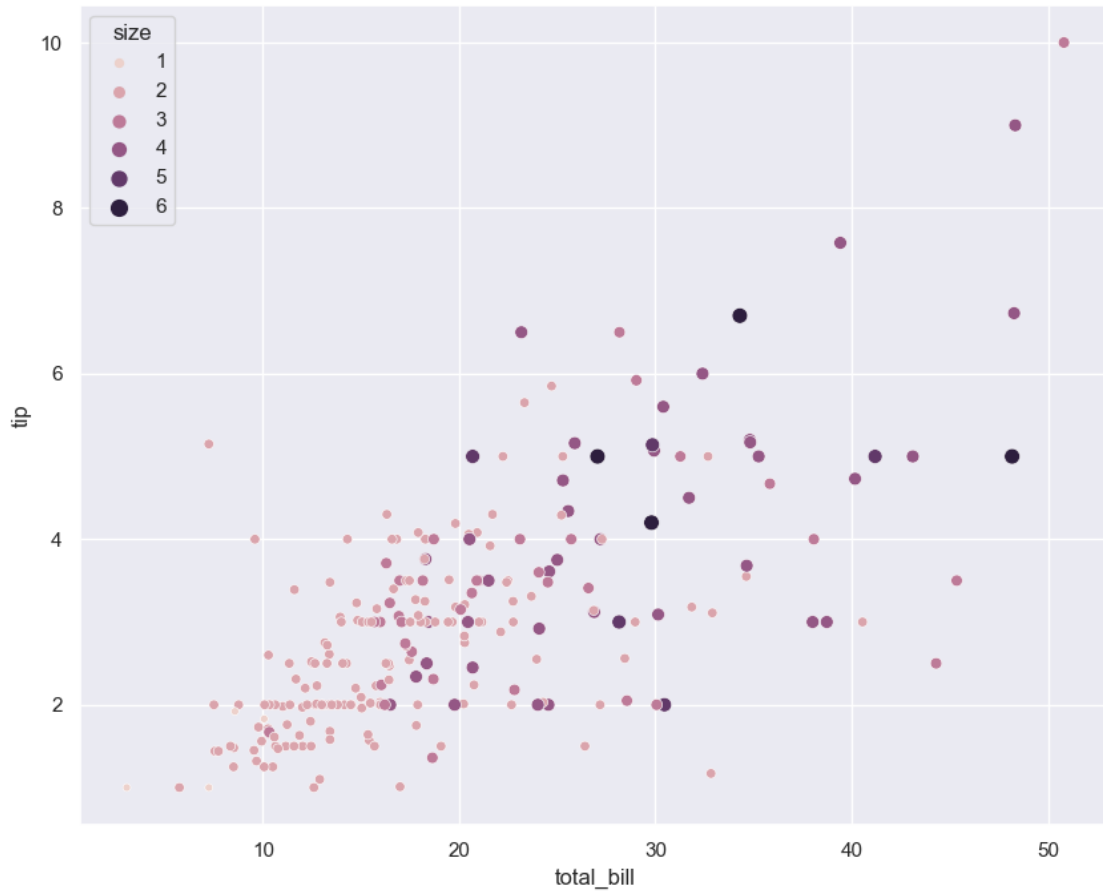
```
[84]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



- A numeric variable can also be assigned to size to apply a semantic mapping to the areas of the points:

```
[85]: sns.scatterplot(data=tips, x='total_bill', y='tip', hue='size', size="size")
```

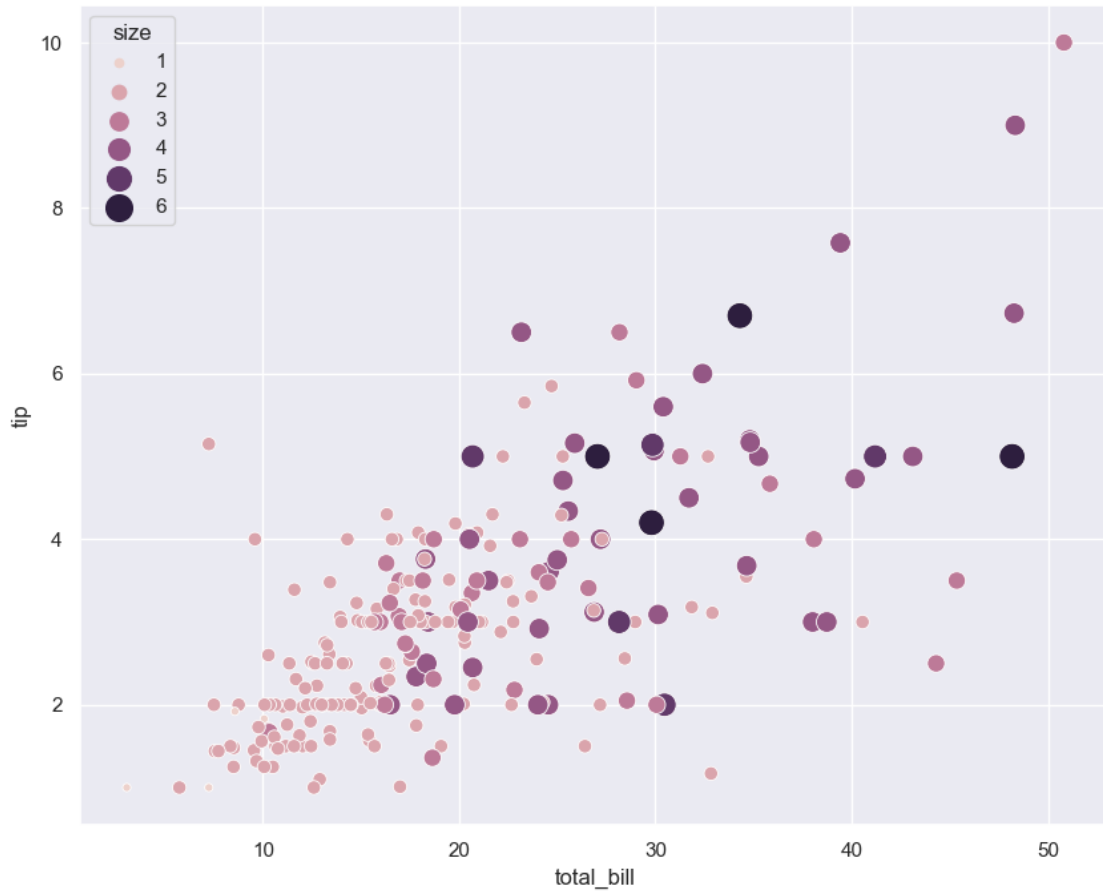
```
[85]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



- Control the range of marker areas with sizes, and set legend="full" to force every unique value to appear in the legend:

```
[86]: sns.scatterplot(  
    data=tips, x="total_bill", y="tip", hue="size", size="size",  
    sizes=(20, 200), legend="full"  
)
```

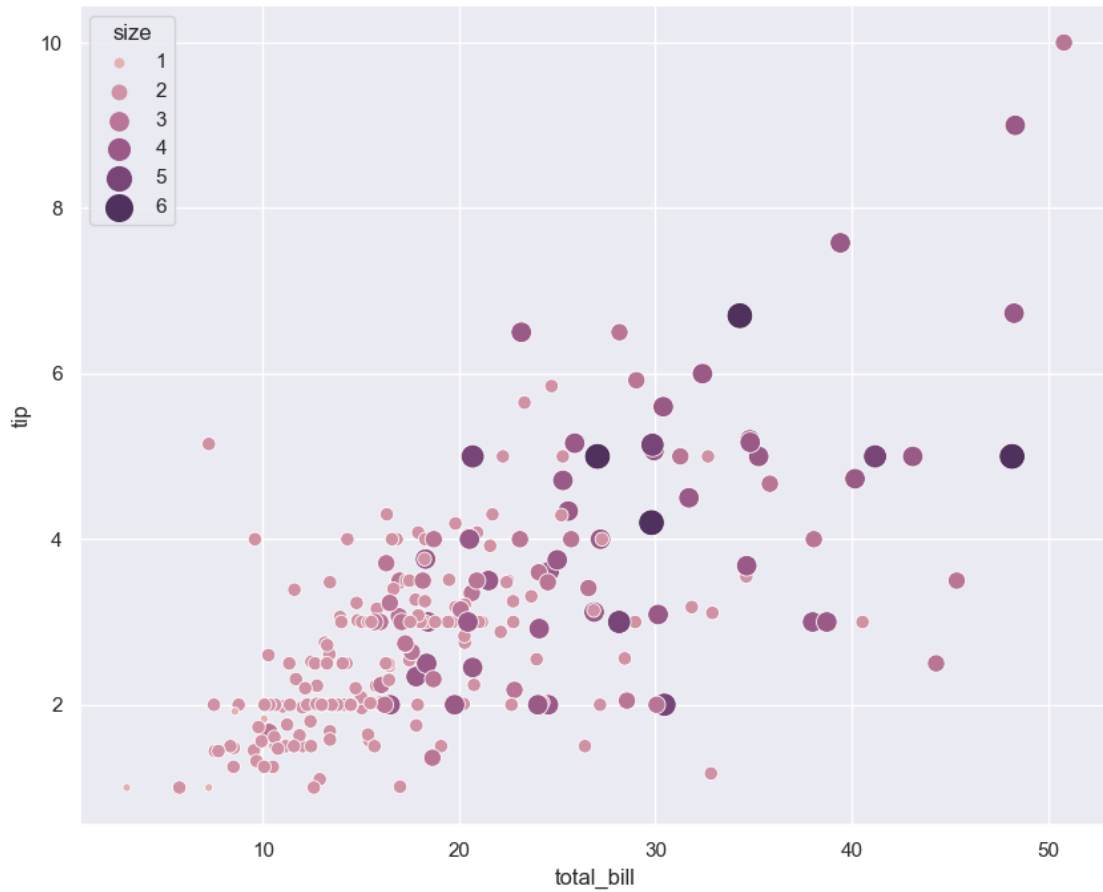
```
[86]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



- Pass a tuple of values or a `matplotlib.colors.Normalize` object to `hue_norm` to control the quantitative hue mapping:

```
[87]: sns.scatterplot(
      data=tips, x="total_bill", y="tip", hue="size", size="size",
      sizes=(20, 200), hue_norm=(0, 7), legend="full"
    )
```

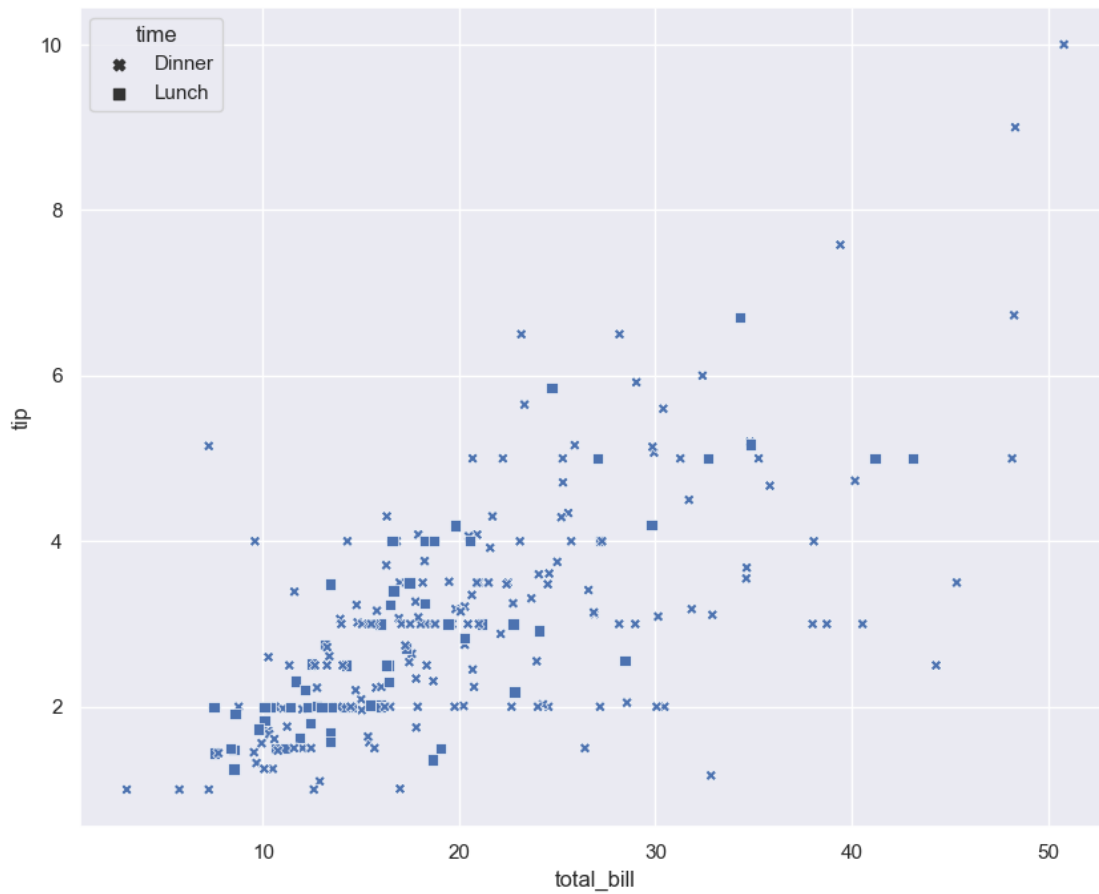
```
[87]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



- Control the specific markers used to map the style variable by passing a Python list or dictionary of marker codes:

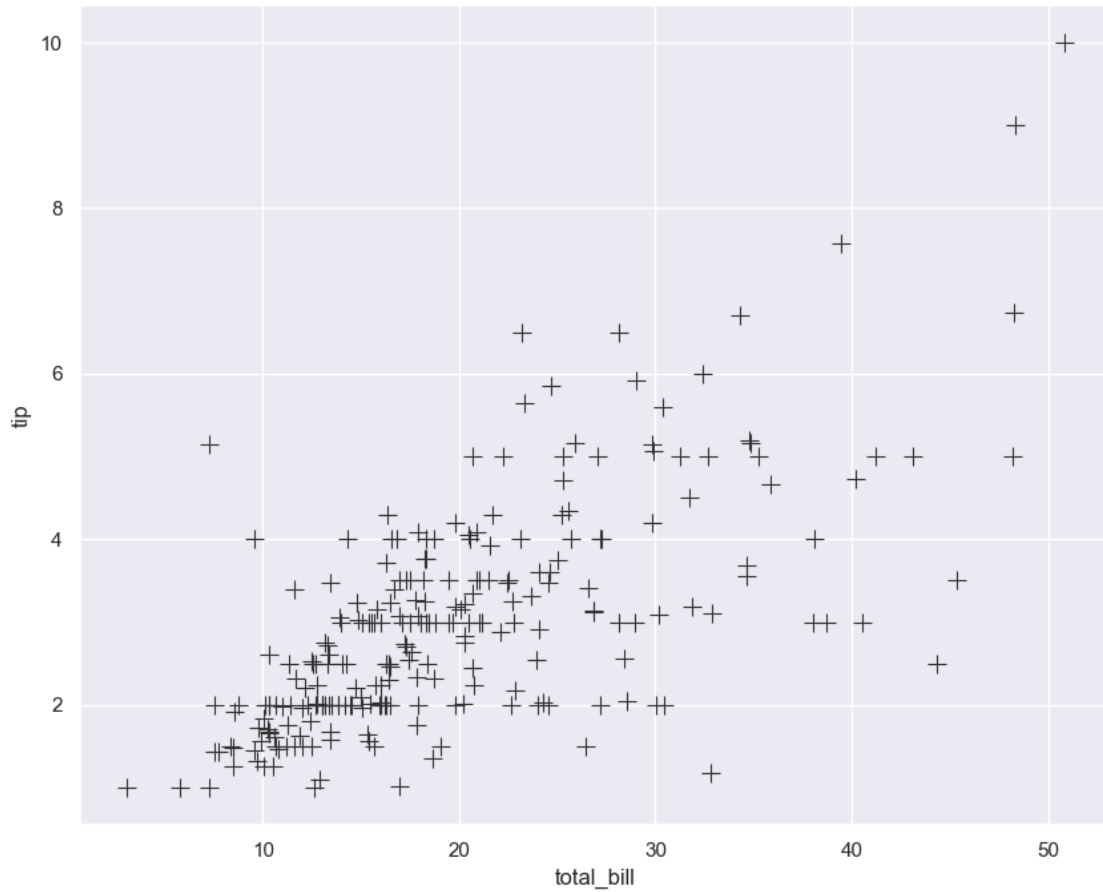
```
[88]: markers = {"Lunch": "s", "Dinner": "X"}
sns.scatterplot(data=tips, x="total_bill", y="tip", style="time",
               ↪ markers=markers)
```

```
[88]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



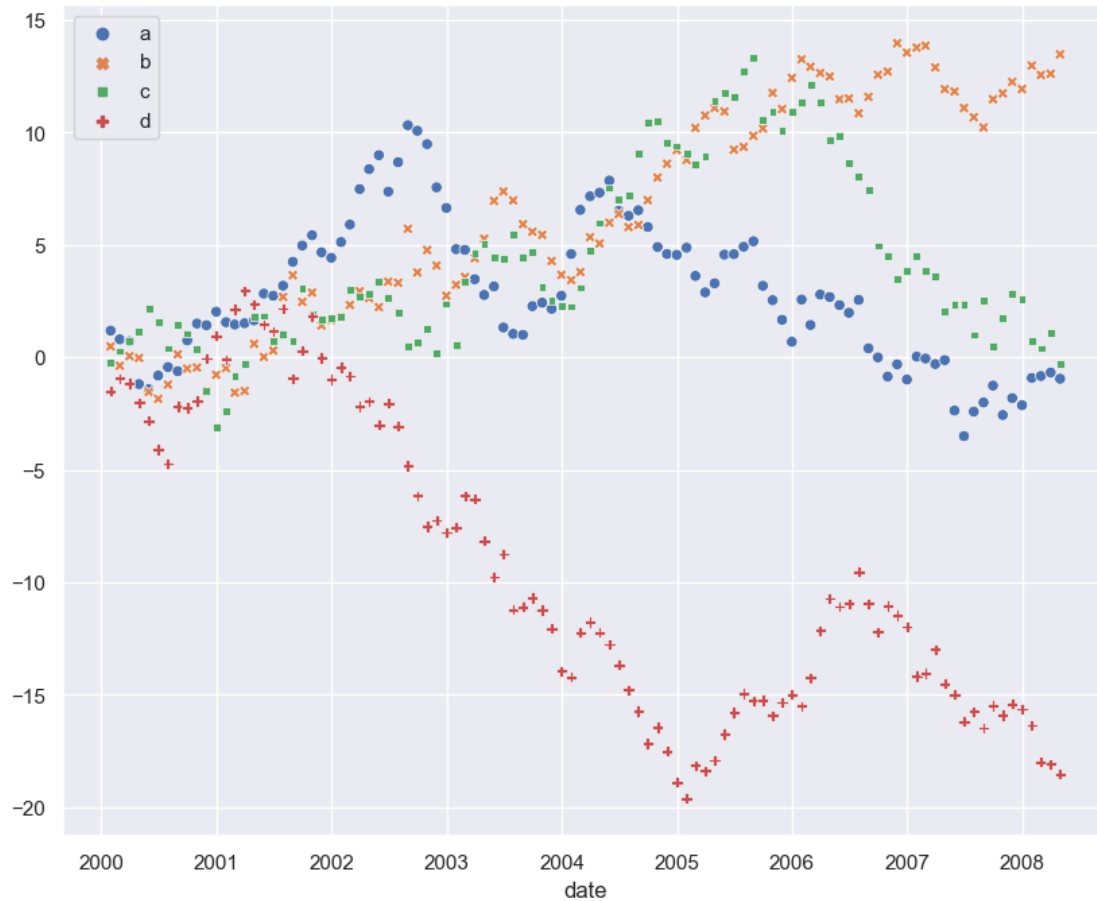
```
[89]: sns.scatterplot(data=tips, x="total_bill", y="tip", s=100, color=".2",  
    ↪marker="+")
```

```
[89]: <AxesSubplot:xlabel='total_bill', ylabel='tip'>
```



```
[90]: index = pd.date_range("1 1 2000", periods=100, freq="m", name="date")
      data = np.random.randn(100, 4).cumsum(axis=0)
      wide_df = pd.DataFrame(data, index, ["a", "b", "c", "d"])
      sns.scatterplot(data=wide_df)
```

```
[90]: <AxesSubplot:xlabel='date'>
```



- Use `relplot()` to combine `scatterplot()` and `FacetGrid`. This allows grouping within additional categorical variables, and plotting them across multiple subplots.
- Using `relplot()` is safer than using `FacetGrid` directly, as it ensures synchronization of the semantic mappings across facets.

```
[91]: sns.relplot(
      data=tips, x="total_bill", y="tip",
      col="time", hue="day", style="day",
      kind="scatter"
    )
```

```
[91]: <seaborn.axisgrid.FacetGrid at 0x21614e09e40>
```





```
[93]: # https://pandas.pydata.org/docs/user\_guide/10min.html # ***
```