YEDDU DIVYA SRI

Mail:divyasri252005@gmail.com

Infosys Springboard Virtual Internship 6.0 (Batch 4 & 5)

BudgetWise AI based Expense Forecasting Tool

❖ **NUMPY**: NumPy stands for Numerical Python.

It is a Python library used for performing fast mathematical, statistical, and array-based operations.

```
#~is output
Pip install Numpy
import numpy as np
n1=np.array([10,20,30,40,50,60])
n2=np.array([60,40,90,100,150])
np.intersect1d(n1,n2)
~array([40, 60])
np.setdiff1d(n1,n2)
~array([10, 20, 30, 50])
np.setdiff1d(n2,n1)
~array([ 90, 100, 150])
n1=np.array([1340,130])
n2=np.array([4,5])
np.sum([n1,n2])
~np.int64(1479)
np.sum([n1,n2],axis=0)
~array([1344, 135])
np.sum([n1,n2],axis=1)
~array([1470, 9])
n1=n1+1
n1
~array([1341, 131])
n1=n1*2
n1
~array([2682, 262])
```

```
n2=n2-1
n2
~array([2, 3])
n1=n1/5
n1
~array([536.4, 52.4])
np.mean(n1)
~np.float64(294.4)
np.std(n1)
~np.float64(241.999999999999)
np.median(n1)
~np.float64(294.4)
import numpy as np
n1=np.array([[1,2,3],[4,5,6],[7,8,9]])
n1
-array([[1, 2, 3],
    [4, 5, 6],
    [7, 8, 9]])
n1[0]
\simarray([1, 2, 3])
n1[1]
~array([4, 5, 6])
n1[:,1]
~array([2, 5, 8])
n1[::]
-array([[1, 2, 3],
    [4, 5, 6],
    [7, 8, 9]])
n1[:,2]
~array([3, 6, 9])
import numpy as np
n1=np.array([[1,2,3],[3,4,5],[7,8,9]])
~array([[1, 2, 3],
    [3, 4, 5],
```

```
[7, 8, 9]]
n1.transpose()
\simarray([[1, 3, 7],
    [2, 4, 8],
    [3, 5, 9]])
n1=np.array([[1,2,3],[3,4,5],[7,8,9]])
n2=np.array([[10,20,30],[30,40,50],[70,80,90]])
n1.dot(n2)
~array([[ 280, 340, 400],
    [500, 620, 740],
    [ 940, 1180, 1420]])
n2.dot(n1)
~array([[ 280, 340, 400],
    [500, 620, 740],
    [ 940, 1180, 1420]])
import numpy as np
n1=np.array([10,20,30,40])
n1
~array([10, 20, 30, 40])
import numpy as np
n2=np.array([[10,20,30,40],[50,60,70,80]])
n2
~array([[10, 20, 30, 40],
    [50, 60, 70, 80]])
n1=np.zeros((1,2))
n1
\simarray([[0., 0.]])
n1=np.zeros((5,5))
~array([[0., 0., 0., 0., 0.],
    [0., 0., 0., 0., 0.]
    [0., 0., 0., 0., 0.]
    [0., 0., 0., 0., 0.]
    [0., 0., 0., 0., 0.]
n1=np.full((5,5),10)
n1
~array([[10, 10, 10, 10, 10],
    [10, 10, 10, 10, 10],
    [10, 10, 10, 10, 10],
```

```
[10, 10, 10, 10, 10],
    [10, 10, 10, 10, 10]])
n1=np.arange(10,20)
n1
~array([10, 11, 12, 13, 14, 15, 16, 17, 18, 19])
n1=np.arange(10,50,5)
~array([10, 15, 20, 25, 30, 35, 40, 45])
n1=np.random.randint(1,100,5)
~array([32, 81, 88, 8, 11], dtype=int32)
n1=np.array([10,20,30])
n2=np.array([40,50,60])
np.vstack((n1,n2))
~array([[10, 20, 30],
    [40, 50, 60]])
n1=np.array([10,20,30])
n2=np.array([40,50,60])
np.hstack((n1,n2))
~array([10, 20, 30, 40, 50, 60])
n1=np.array([[1,3,4],[45,65,66]])
n1.shape
\sim(2, 3)
n1.shape=(3,2)
n1.shape
\sim(3, 2)
n1=np.array([10,20,30])
n2=np.array([40,50,60])
np.column_stack((n1,n2))
~array([[10, 40],
    [20, 50],
    [30, 60]])
```

❖ PANDAS: Pandas is a Python library used for data manipulation and analysis. It provides data structures like Series and DataFrame to handle and analyze data easily.

import pandas as pd

```
s1=pd.Series([1,2,3,4,5])
s1
~0 1
1 2
2 3
3 4
4 5
dtype: int64
s1=pd.Series([1,2,3,4,5],index=['a','b','c','d','e'])
s1
~a 1
b 2
c 3
d 4
e 5
dtype: int64
pd.Series({'a':10,'b':20,'c':30})
~a 10
b 20
c 30
dtype: int64
pd.Series({'a':10,'b':20,'c':30},index=['c','a','g','k'])
~c 30.0
a 10.0
g
   NaN
k NaN
dtype: float64
s1=pd.Series([1,2,3,4,5,6,7,8,9])
s1[3]
~np.int64(4)
s1[::]
~0 1
1 2
2 3
3 4
4 5
5 6
6 7
7 8
8 9
dtype: int64
```

```
s1[:-1]
~0 1
1
   2
2
   3
  4
3
4
  5
5
  6
6 7
7
   8
dtype: int64
s1[:-2]
~0 1
1
   2
2
   3
3
  4
4 5
5 6
6 7
dtype: int64
s1[:-3]
s1[-3:]
~6 7
7 8
8
   9
dtype: int64
s1[:4]
~0 1
1 2
2 3
3 4
dtype: int64
import pandas as pd
df = pd.read_csv("iris.csv")
df.head()
    sepal_length sepal_width petal_length petal_width
                                                      species
0
       5.1
                3.5
                         1.4
                                 0.2 setosa
        4.9
                3.0
1
                         1.4
                                  0.2 setosa
2
        4.7
                3.2
                         1.3
                                  0.2 setosa
3
        4.6
                3.1
                         1.5
                                  0.2 setosa
4
        5.0
                3.6
                         1.4
                                  0.2 setosa
```

```
df.tail()
```

```
~ sepal_length sepal_width petal_length petal_width
                                                         species
145
          6.7
                    3.0
                              5.2
                                       2.3 virginica
                                        1.9 virginica
146
           6.3
                    2.5
                              5.0
147
           6.5
                    3.0
                              5.2
                                        2.0 virginica
                                        2.3 virginica
148
           6.2
                    3.4
                              5.4
149
           5.9
                    3.0
                              5.1
                                        1.8 virginica
```

df.shape

 $\sim (150, 5)$

df.columns

~ Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width', 'species'], dtype='object')

df.dtypes

~ sepal_length float64 sepal_width float64 petal_length float64 petal_width float64 species object dtype: object

df.describe()

sepal_length sepal_width petal_length petal_width 150.000000 150.000000 150.000000 150.000000 count 1.199333 5.843333 3.057333 3.758000 mean 0.828066 0.435866 1.765298 0.762238 std 4.300000 min 2.000000 1.000000 0.100000 25% 5.100000 2.800000 1.600000 0.300000 50% 5.800000 3.000000 4.350000 1.300000 75% 6.400000 3.300000 5.100000 1.800000 7.900000 4.400000 6.900000 2.500000 max

df.info()

~ <class 'pandas.core.frame.DataFrame'>

RangeIndex: 150 entries, 0 to 149 Data columns (total 5 columns):

Column Non-Null Count Dtype

dtypes: float64(4), object(1)

⁰ sepal_length 150 non-null float64

¹ sepal_width 150 non-null float64

² petal_length 150 non-null float64

³ petal_width 150 non-null float64

⁴ species 150 non-null object

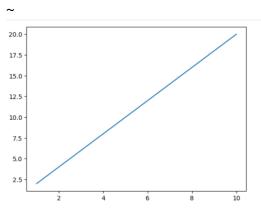
❖ Plotting using Matplotlib Pyplot: Pyplot is a module in the Matplotlib library that provides simple functions for creating plots and visualizing data easily. It allows users to generate line graphs, bar charts, histograms, scatter plots, and more with simple commands.

import numpy as np from matplotlib import pyplot as plt

```
import numpy as np
x = np.arange(1, 11)
x
~ array([ 1, 2, 3, 4, 5, 6, 7, 8, 9, 10])
y=2*x
y
~ array([ 2, 4, 6, 8, 10, 12, 14, 16, 18, 20])
```

• Line Plot: A **line plot** is a type of chart used to show data points connected by straight lines.

plt.plot(x,y)
plt.show()

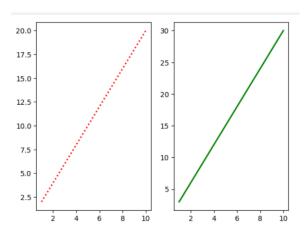


plt.plot(x,y)
plt.title("Line plot")
plt.xlabel("x-label")
plt.ylabel("y-label")
plt.show()

Line plot

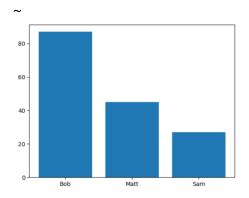
20.0
17.5
15.0
12.5
2 4 5 8 10

```
plt.plot(x,y,color='red',linestyle=':',linewidth=2)
plt.show()
20.0
15.0
12.5
 7.5
 5.0
x=np.arange(1,11)
y1=2*x
y2=3*x
plt.plot(x,y1,color='red',linestyle=':',linewidth=2)
plt.plot(x,y2,color='orange',linestyle='-',linewidth=2)
plt.title("Line plot")
plt.xlabel('xlabel')
plt.ylabel('ylabel')
plt.grid(True)
plt.show()
                       Line plot
   25
   20
   10
plt.subplot(1,2,1)
plt.plot(x,y1,color='red',linestyle=':',linewidth=2)
plt.subplot(1,2,2)
plt.plot(x,y2,color='g',linestyle='-',linewidth=2)
plt.show()
```

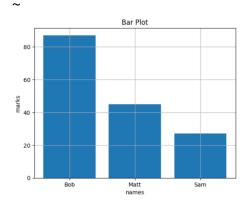


• Bar Plot: A bar plot (or bar chart) is a graph that represents categorical data with rectangular bars.

```
student={"Bob":87,"Matt":45,"Sam":27}
names=list(student.keys())
values=list(student.values())
plt.bar(names,values)
plt.show()
```

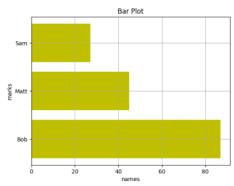


plt.bar(names,values)
plt.title("Bar Plot")
plt.xlabel('names')
plt.ylabel('marks')
plt.grid(True)
plt.show()



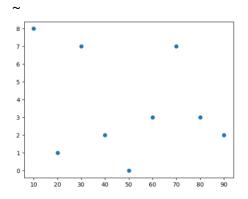
plt.barh(names, values, color='y')

```
plt.title("Bar Plot")
plt.xlabel('names')
plt.ylabel('marks')
plt.grid(True)
plt.show()
```

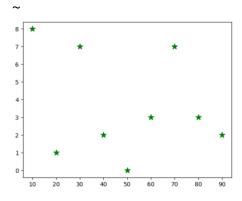


• Scatter Plot: A scatter plot is used to display the relationship between two variables using dots on a graph.

```
x=[10,20,30,40,50,60,70,80,90]
a=[8,1,7,2,0,3,7,3,2]
plt.scatter(x,a)
plt.show()
```



x=[10,20,30,40,50,60,70,80,90] a=[8,1,7,2,0,3,7,3,2] plt.scatter(x,a,marker="*",c="g",s=100) plt.show()



```
x=[10,20,30,40,50,60,70,80,90]

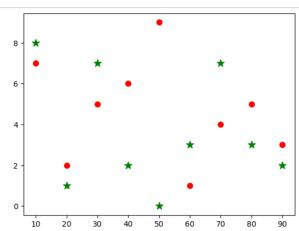
a=[8,1,7,2,0,3,7,3,2]

b=[7,2,5,6,9,1,4,5,3]
```

plt.scatter(x,a,marker="*",c="g",s=100) plt.scatter(x,b,marker=".",c="r",s=200)

plt.show()

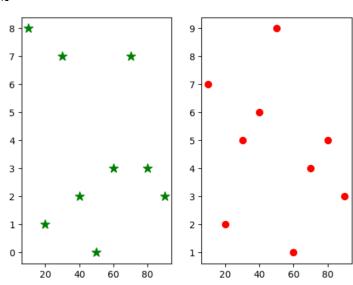
~



$$\begin{split} x &= [10, 20, 30, 40, 50, 60, 70, 80, 90] \\ a &= [8, 1, 7, 2, 0, 3, 7, 3, 2] \\ b &= [7, 2, 5, 6, 9, 1, 4, 5, 3] \\ plt.subplot(1, 2, 1) \\ plt.scatter(x, a, marker = "*", c = "g", s = 100) \\ plt.subplot(1, 2, 2) \\ plt.scatter(x, b, marker = ".", c = "r", s = 200) \end{split}$$

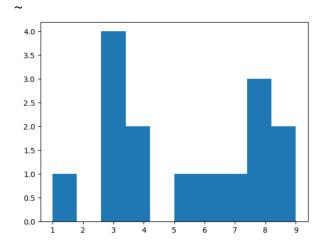
plt.show()

~

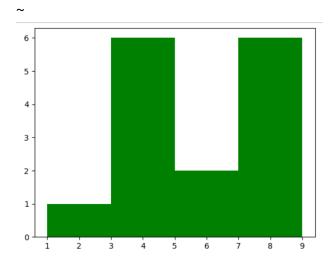


• Histogram Plot: A histogram plot is used to show the frequency distribution of continuous data using adjacent rectangular bars.

```
#histogram plot
#creating data
data=[1,3,3,3,3,9,9,5,4,4,8,8,8,6,7]
#making histogram
plt.hist(data)
plt.show()
```



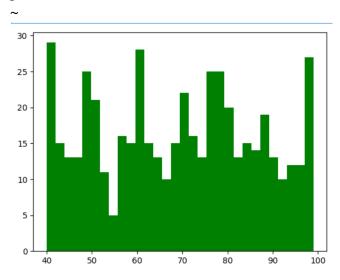
plt.hist(data,color="g",bins=4)
plt.show()



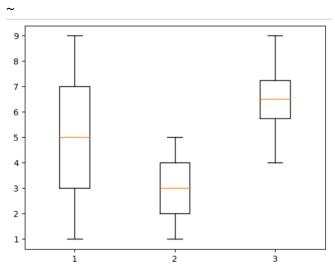
import pandas as pd
soil=pd.read_csv('soil_crop_dataset.csv')
soil.head()

:		рН	Nitrogen	Phosphorus	Potassium	Moisture	Temperature	Rainfall	Soil_Type	Recommended_Crop
	0	5.53	72	52	60	22.01	24.91	249	Sandy,Peaty,Loamy	Peanut, Maize, Tomato, Barley, Carrot
	1	6.71	87	18	45	21.85	29.03	135	Clayey,Peaty,Red,Black	Peanut, Onion, Potato, Millet
	2	8.49	88	13	45	20.36	18.88	52	Red,Black,Clayey	Potato, Millet, Tomato, Cotton
	3	7.27	46	60	31	15.64	19.61	77	Silty,Red,Clayey	Potato,Rice,Tomato,Wheat,Onion
	4	8.22	60	60	33	8.81	33.09	171	Silty,Sandy,Peaty	Tomato,Potato,Wheat,Rice

plt.hist(soil['Nitrogen'],bins=30,color='g')
plt.show()

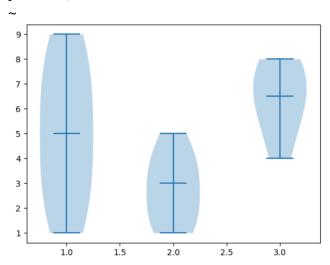


• Box Plot: A box plot is used to display the distribution, spread, and outliers of a dataset using quartiles.



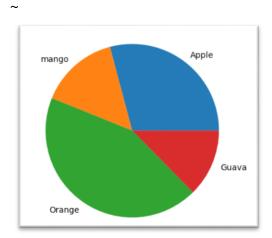
• Violin Plot: A violin plot is used to show the distribution of data across categories, combining features of a box plot and a density plot.

```
one=[1,2,3,4,5,6,7,8,9]
two=[1,2,3,4,5,4,3,2,1]
three=[6,7,8,7,8,6,5,4]
data=list([one,two,three])
plt.violinplot(data,showmedians=True)
plt.show()
```

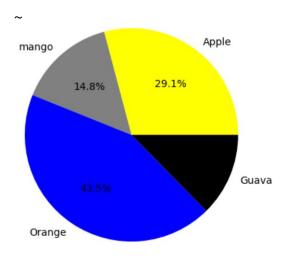


• Pie Plot: A pie plot is used to show the proportion or percentage of different categories as slices of a whole circle.

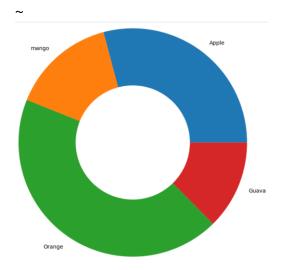
```
fruit=['Apple','mango',"Orange","Guava"]
quantity=[67,34,100,29]
plt.pie(quantity,labels=fruit)
plt.show()
```



 $plt.pie(quantity,labels=fruit,autopct='\%0.1f\%\%',colors=['yellow','grey','blue','black'])\\ plt.show()$



plt.pie(quantity,labels=fruit,radius=2)
plt.pie([1],colors=['w'],radius=1)
plt.show()



❖ Plotting using Seaborn and Matplotlib Pyplot:

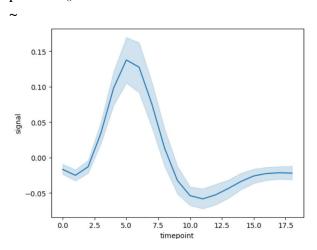
Seaborn is a Python data visualization library built on top of Matplotlib.

import seaborn as sns from matplotlib import pyplot as plt

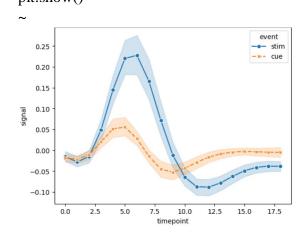
fmri = sns.load_dataset("fmri")
fmri.head()

~					
	subject	timepoint	event	region	signal
0	s13	18	stim	parietal	-0.017552
1	s5	14	stim	parietal	-0.080883
2	s12	18	stim	parietal	-0.081033
3	s11	18	stim	parietal	-0.046134
4	s10	18	stim	parietal	-0.037970

sns.lineplot(x="timepoint",y="signal",data=fmri)
plt.show()



sns.lineplot(x="timepoint",y="signal",hue="event",style="event",markers=Tru
e,data=fmri)
plt.show()



fmri=sns.load_dataset("fmri")
fmri.head()

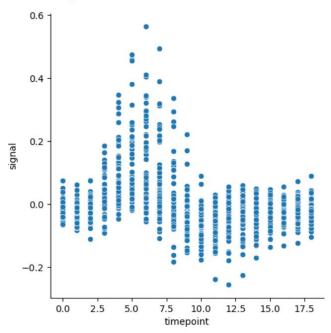
~	subject	timepoint	event	region	signal
0	s13	18	stim	parietal	-0.017552
1	s5	14	stim	parietal	-0.080883
2	s12	18	stim	parietal	-0.081033
3	s11	18	stim	parietal	-0.046134
4	s10	18	stim	parietal	-0.037970

• Relplot: A relplot (relationship plot) in Seaborn is used to visualize relationships between two variables, often as scatter or line plots.

fmri.shape ~ (1064, 5)

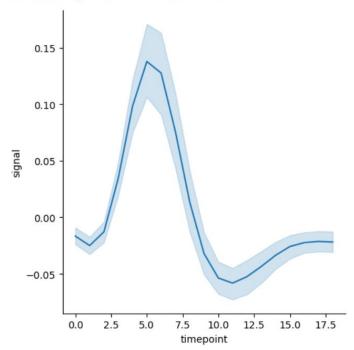
sns.relplot(data=fmri,x="timepoint",y="signal")

<seaborn.axisgrid.FacetGrid at 0x26ceb742780>



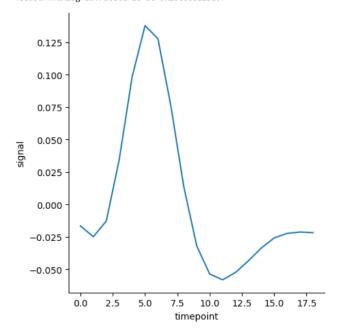
sns.relplot(data=fmri,x="timepoint",y="signal",kind="line")

<seaborn.axisgrid.FacetGrid at 0x26cec7f9bb0>



```
sns.relplot(
   data=fmri,kind="line",
   x="timepoint",y="signal",errorbar=None
)
```





Hue: It is used to add a third variable in a plot by coloring the data points according to different categories.

Kind: It specifies the type of plot to draw, such as 'scatter' or 'line' in functions like relplot().

style: It is used to change the appearance of data points or lines (like marker shapes or line styles) based on a categorical variable.

```
sns.relplot(
   data=fmri,kind="line",
   x="timepoint",y="signal",
   hue="event"
)
 <seaborn.axisgrid.FacetGrid at 0x26cec8ca780>
    0.25
    0.20
    0.15
     0.10
                                                        event
                                                          stim
    0.05
     0.00
    -0.05
   -0.10
                                          15.0 17.5
          0.0
                     5.0
                               10.0
                                     12.5
                           timepoint
```

```
sns.relplot(
   data=fmri,kind="line",
   x="timepoint",y="signal",
   hue="region",style="event"
)
 <seaborn.axisgrid.FacetGrid at 0x26cec7c6930>
     0.3
     0.2
                                                             region
 signal
                                                             parietal
     0.1
                                                             frontal
                                                             stim
                                                         --- cue
     0.0
    -0.1
               2.5
                     5.0
                           7.5
                                10.0
                                       12.5
                                            15.0
```

Dashes: Used to control or remove the line style (solid, dashed, etc.) in line plots. **Markers:** Used to display symbols at data points (like circles, squares, or triangles) in line or scatter plots.

```
sns.relplot(
   data=fmri,kind="line",
   x="timepoint",y="signal",
   hue="region",style="event",
   dashes=False,markers=True
)
 <seaborn.axisgrid.FacetGrid at 0x26cec937860>
    0.3
    0.2
                                                    region
                                                    parietal
    0.1
                                                    frontal
                                                    event
    0.0
   -0.1
                       7.5
                           10.0
                                      15.0
```

***** Categorical scatterplots

- Categorical scatterplots are used to show the relationship between a categorical variable and a numerical variable using points instead of bars.
- Categorical data represents values that belong to distinct groups or categories, such as colors, names, or types.

two types of categorical data:

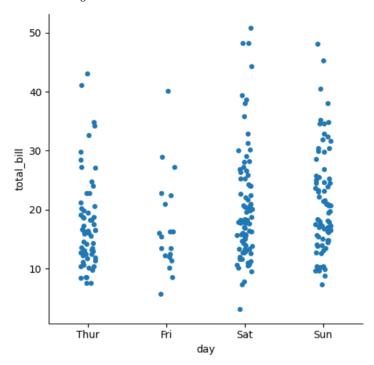
- **Nominal Data:** Categories that **do not have any specific order** (e.g., colors red, blue, green).
- Ordinal Data: Categories that have a meaningful order or ranking (e.g., ratings poor, average, good, excellent).

import seaborn as sns
tips=sns.load_dataset("tips")
tips.head()

~							
	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

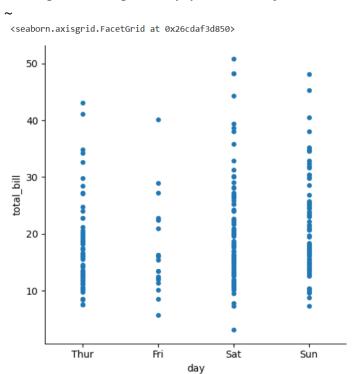
sns.catplot(data=tips,x='day',y='total_bill')

<seaborn.axisgrid.FacetGrid at 0x26ceabf75f0>

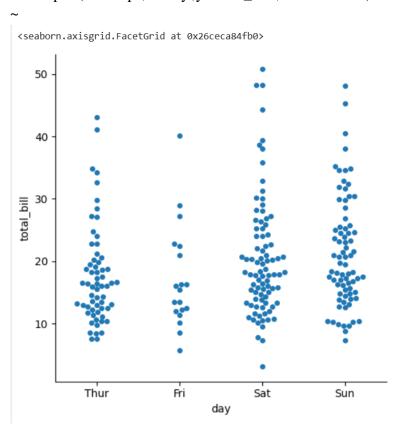


Jitter: adds small random noise to data points in a plot to prevent overlapping and make the distribution clearer.

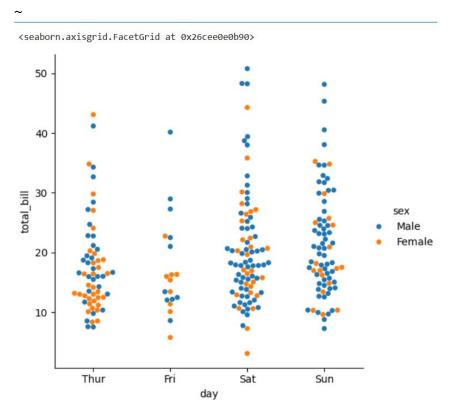
 $sns.catplot(data \!\!=\!\! tips, \!x \!\!=\!\! 'day', \!y \!\!=\!\! 'total_bill', \!jitter \!\!=\!\! False)$



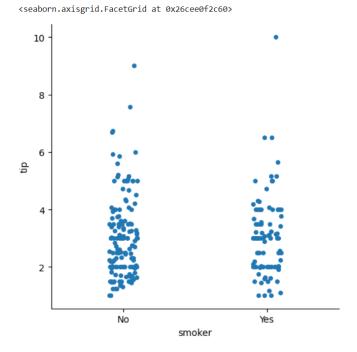
#prevent from overlapping (swarm plot)
sns.catplot(data=tips,x='day',y='total_bill',kind='swarm')



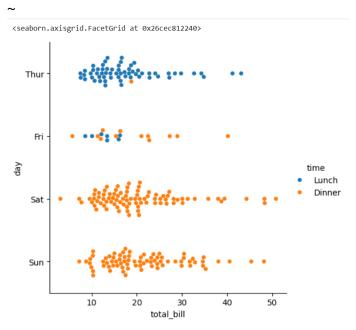
#add the hue sematic sns.catplot(data=tips,x='day',y='total_bill',hue='sex',kind='swarm')



#order parameter - to display multiple categorical plot in same figure sns.catplot(data=tips,x='smoker',y='tip',order=['No','Yes'])



#plot on vertical sns.catplot(data=tips,x='total_bill',y='day',hue='time',kind='swarm')

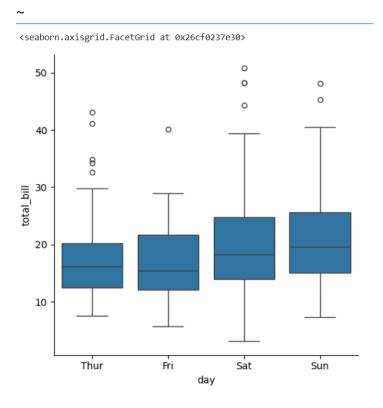


• **Comparing distribution** means analyzing how data values are spread across different groups or categories to identify patterns or differences.

#comparing distribution

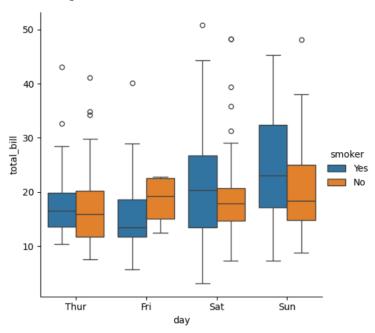
 box plot: A box plot is used to display the distribution of data based on five summary statistics — minimum, first quartile, median, third quartile, and maximum.

#sns.catplot(data=tips,x='total_bill',y='day',kind='box')
sns.catplot(data=tips,x='day',y='total_bill',kind='box')



#adding the hue sematic sns.catplot(data=tips,x='day',y='total_bill',hue='smoker',kind='box')

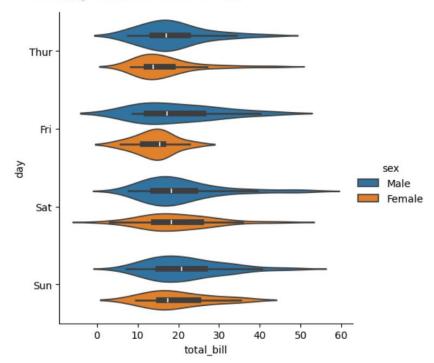
<seaborn.axisgrid.FacetGrid at 0x26cf1c3eb10>



• **violin plot**: A violin plot is used to visualize the distribution of data and its probability density, combining features of a box plot and a kernel density plot.

sns.catplot(data=tips,x='total_bill',y='day',hue='sex',kind='violin')

<seaborn.axisgrid.FacetGrid at 0x26cf1d8a480>



#split in the violin plot sns.catplot(data=tips,x='day',y='total_bill',hue='sex',kind='violin',split=True)

<seaborn.axisgrid.FacetGrid at 0x26cf57783e0>

