

Data visualization

▼ step-1 import labirary

```
1 import seaborn as sns
2 import matplotlib.pyplot as plt
```

▼ Step-2 load data set

```
1 titanic=sns.load_dataset("titanic")
2 titanic.head()
3
```

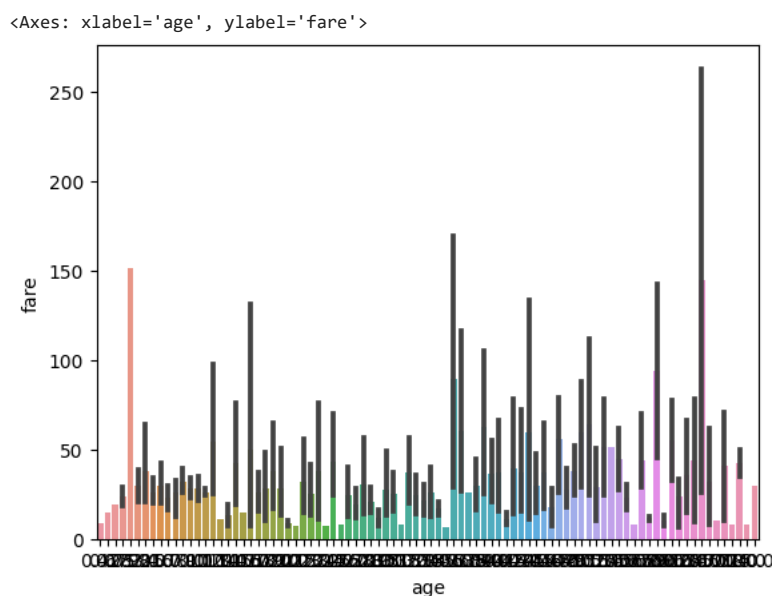
	survived	pclass	sex	age	sibsp	parch	fare	embarked	class	who	adult_male	deck	embark_town	alive	alone
0	0	3	male	22.0	1	0	7.2500	S	Third	man	True	NaN	Southampton	no	False
1	1	1	female	38.0	1	0	71.2833	C	First	woman	False	C	Cherbourg	yes	False
2	1	3	female	26.0	0	0	7.9250	S	Third	woman	False	NaN	Southampton	yes	True
3	1	1	female	35.0	1	0	53.1000	S	First	woman	False	C	Southampton	yes	False
4	0	3	male	35.0	0	0	8.0500	S	Third	man	True	NaN	Southampton	no	True

▼ Step 3 plot a graph

```
1 sns.lineplot(x="fare",y="age",data=titanic)
2 plt.xlim(3)
3 plt.ylim(4)
4 plt.title("Check")
5 plt.show()
```

▼ Bar Plot

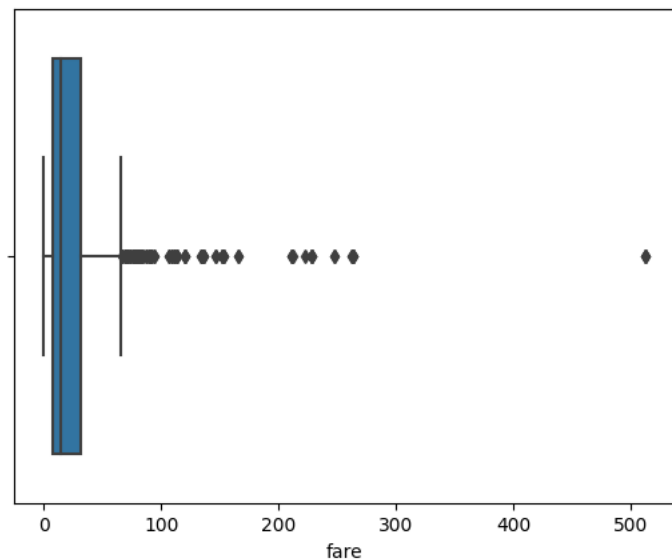
```
1 df = sns.load_dataset("titanic")
2 sns.barplot(data=df, x="age", y="fare")
```



▼ Boxplot

```
1 df = sns.load_dataset("titanic")
2 sns.boxplot(x=df["fare"])
```

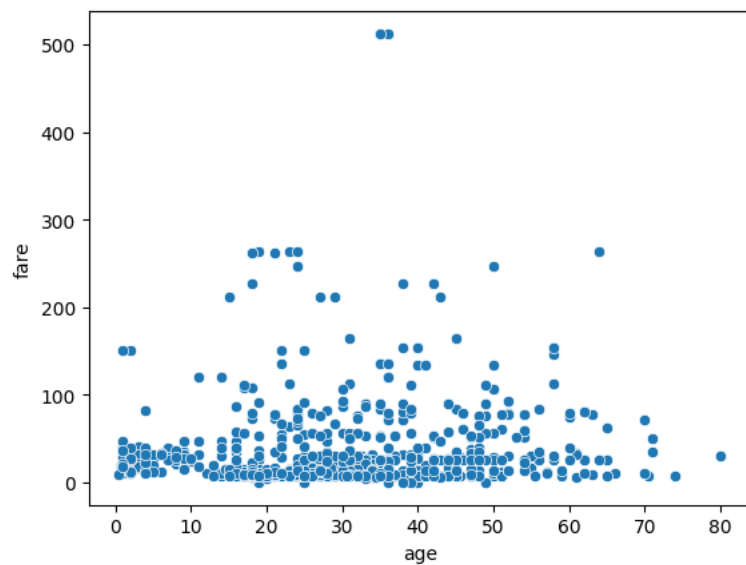
<Axes: xlabel='fare'>



▼ Scatter plot

```
1 sns.scatterplot(data=titanic, x="age", y="fare")
```

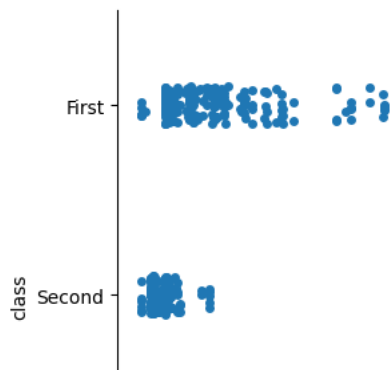
<Axes: xlabel='age', ylabel='fare'>



▼ Cat plot

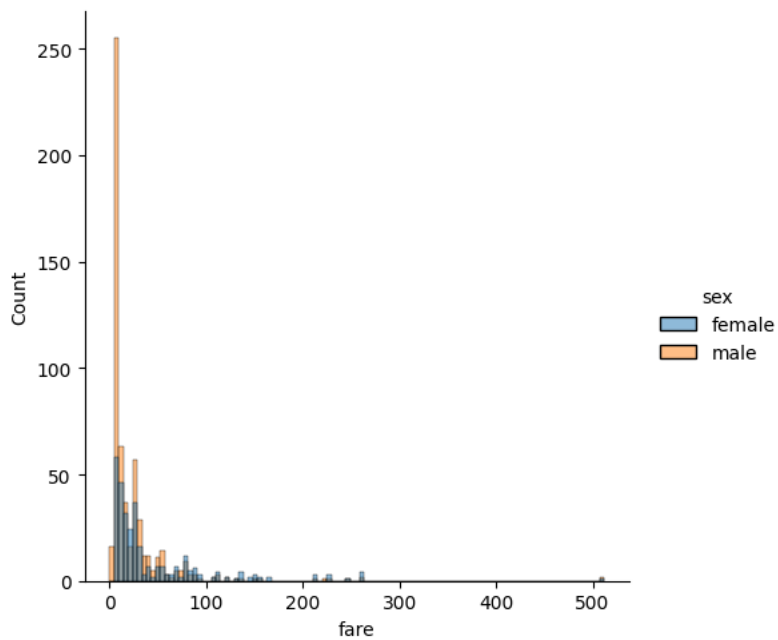
```
1 df = sns.load_dataset("titanic")
2 sns.catplot(data=df, x="fare", y="class")
```

```
<seaborn.axisgrid.FacetGrid at 0x7fc5bf9a1330>
```



```
1 sns.displot(data = titanic , x = "fare" , hue = "sex", hue_order = ['female', 'male'])
```

```
<seaborn.axisgrid.FacetGrid at 0x7fc5bf8ffb50>
```

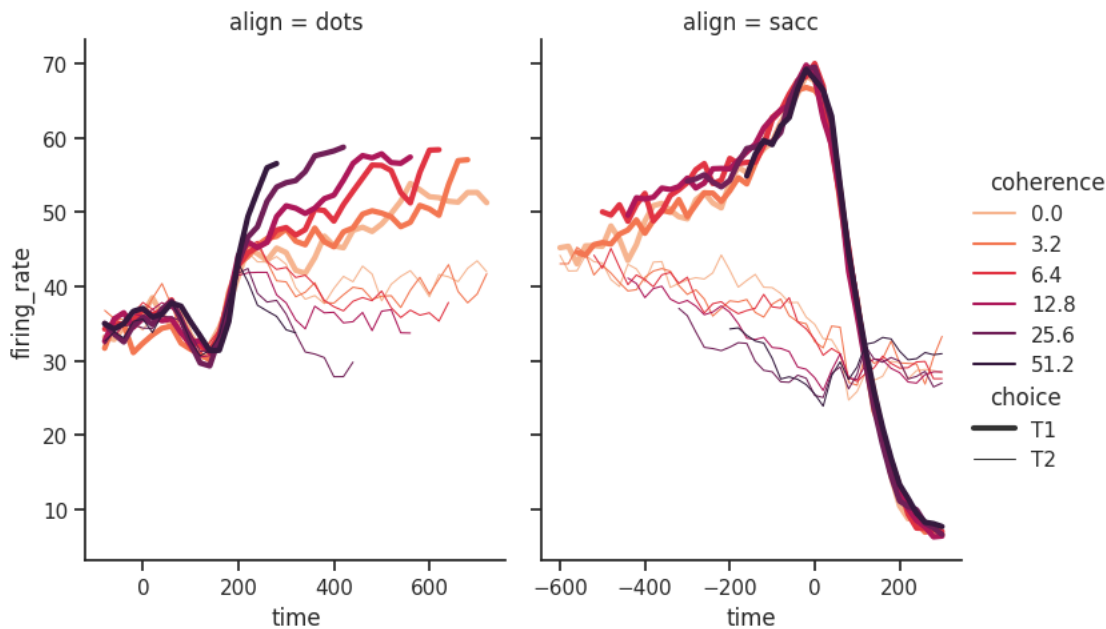


```
1 sns.heatmap(data = titanic.corr())
```

```
<ipython-input-68-f4da9c94fae6>:1: FutureWarning: The default value of numeric_only in DataFrame.corr is deprecated. In a futur
sns.heatmap(data = titanic.corr())
```

```
1 import seaborn as sns
2 sns.set_theme(style="ticks")
3
4 dots = sns.load_dataset("dots")
5
6 # Define the palette as a list to specify exact values
7 palette = sns.color_palette("rocket_r")
8
9 # Plot the lines on two facets
10 sns.relplot(
11     data=dots,
12     x="time", y="firing_rate",
13     hue="coherence", size="choice", col="align",
14     kind="line", size_order=["T1", "T2"], palette=palette,
15     height=5, aspect=.75, facet_kws=dict(sharex=False),
16 )
17
```

```
<seaborn.axisgrid.FacetGrid at 0x7fc5bf871b40>
```

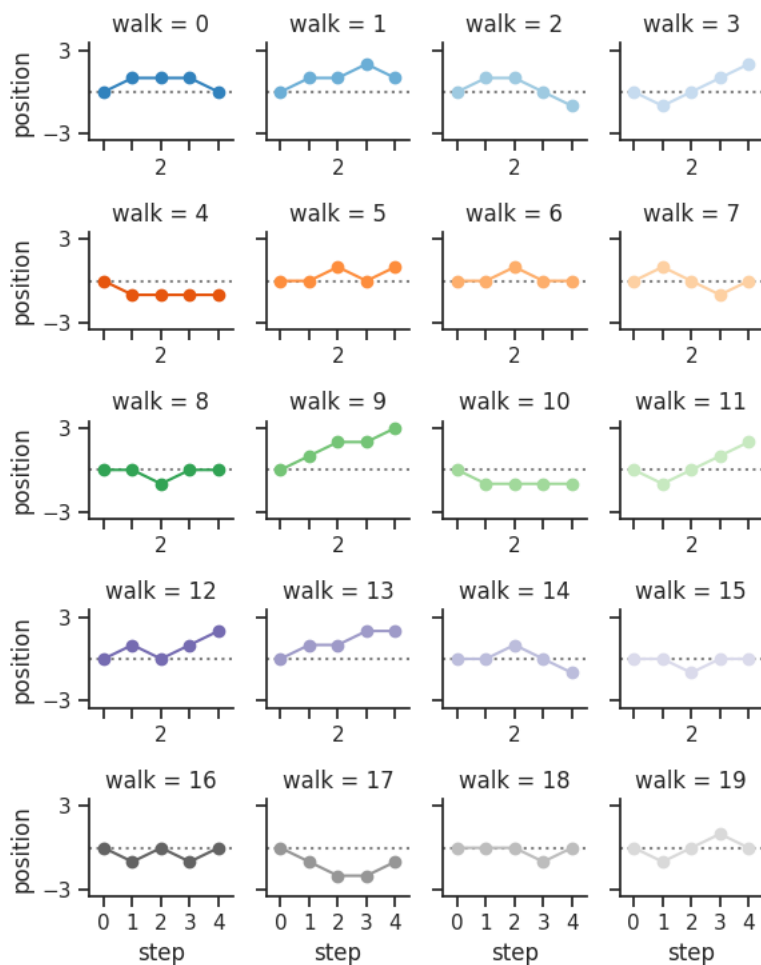


```
1 import numpy as np
2 import pandas as pd
3 import seaborn as sns
4 import matplotlib.pyplot as plt
5
6 sns.set_theme(style="ticks")
7
8 # Create a dataset with many short random walks
9 rs = np.random.RandomState(4)
10 pos = rs.randint(-1, 2, (20, 5)).cumsum(axis=1)
11 pos -= pos[:, 0, np.newaxis]
12 step = np.tile(range(5), 20)
13 walk = np.repeat(range(20), 5)
14 df = pd.DataFrame(np.c_[pos.flat, step, walk],
15                  columns=["position", "step", "walk"])
16
17 # Initialize a grid of plots with an Axes for each walk
18 grid = sns.FacetGrid(df, col="walk", hue="walk", palette="tab20c",
19                    col_wrap=4, height=1.5)
20
21 # Draw a horizontal line to show the starting point
22 grid.refline(y=0, linestyle=":")
23
24 # Draw a line plot to show the trajectory of each random walk
25 grid.map(plt.plot, "step", "position", marker="o")
26
27 # Adjust the tick positions and labels
28 grid.set(xticks=np.arange(5), yticks=[-3, 3],
29         xlim=(-5, 5), ylim=(-3, 3))
29
```

```

29 # Adjust the arrangement of the plots
30
31 # Adjust the arrangement of the plots
32 grid.fig.tight_layout(w_pad=1)
33

```

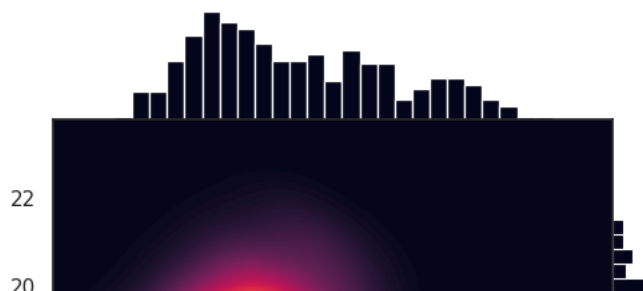


```

1 import seaborn as sns
2 sns.set_theme(style="white")
3
4 df = sns.load_dataset("penguins")
5
6 g = sns.JointGrid(data=df, x="body_mass_g", y="bill_depth_mm", space=0)
7 g.plot_joint(sns.kdeplot,
8             fill=True, clip=((2200, 6800), (10, 25)),
9             thresh=0, levels=100, cmap="rocket")
10 g.plot_marginals(sns.histplot, color="#03051A", alpha=1, bins=25)
11

```

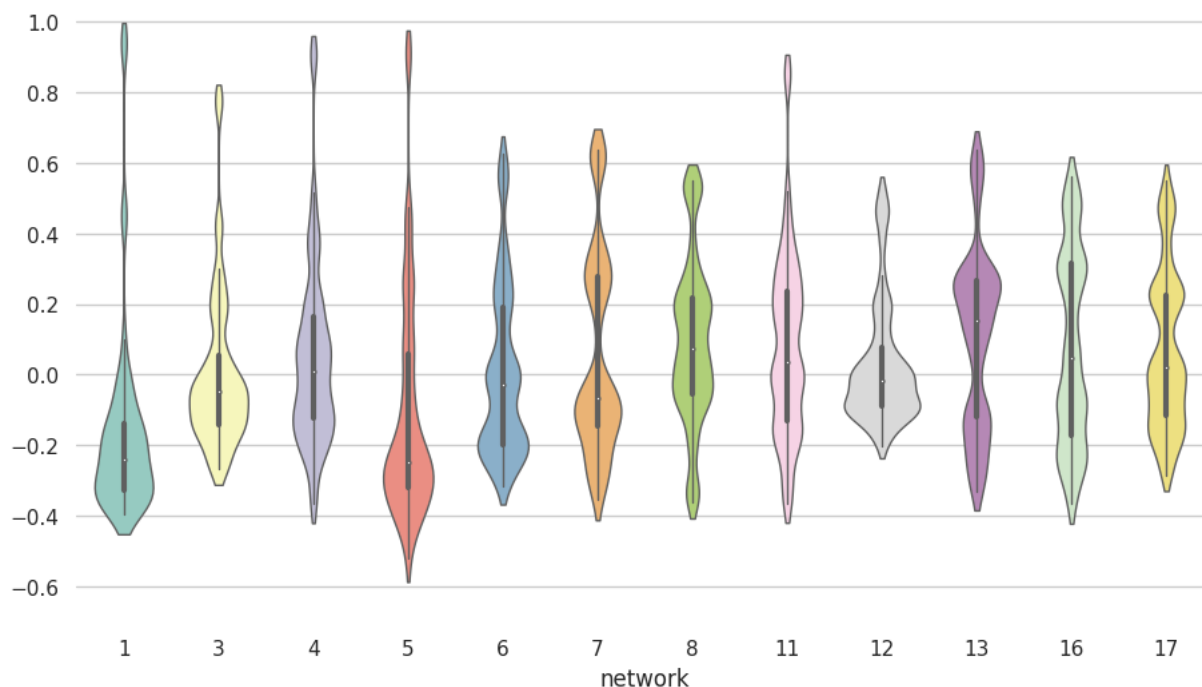
<seaborn.axisgrid.JointGrid at 0x7fc5bf4d7f40>



```

1 import seaborn as sns
2 import matplotlib.pyplot as plt
3 sns.set_theme(style="whitegrid")
4
5 # Load the example dataset of brain network correlations
6 df = sns.load_dataset("brain_networks", header=[0, 1, 2], index_col=0)
7
8 # Pull out a specific subset of networks
9 used_networks = [1, 3, 4, 5, 6, 7, 8, 11, 12, 13, 16, 17]
10 used_columns = (df.columns.get_level_values("network")
11                 .astype(int)
12                 .isin(used_networks))
13 df = df.loc[:, used_columns]
14
15 # Compute the correlation matrix and average over networks
16 corr_df = df.corr().groupby(level="network").mean()
17 corr_df.index = corr_df.index.astype(int)
18 corr_df = corr_df.sort_index().T
19
20 # Set up the matplotlib figure
21 f, ax = plt.subplots(figsize=(11, 6))
22
23 # Draw a violinplot with a narrower bandwidth than the default
24 sns.violinplot(data=corr_df, palette="Set3", bw=.2, cut=1, linewidth=1)
25
26 # Finalize the figure
27 ax.set(ylim=(-.7, 1.05))
28 sns.despine(left=True, bottom=True)
29

```



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