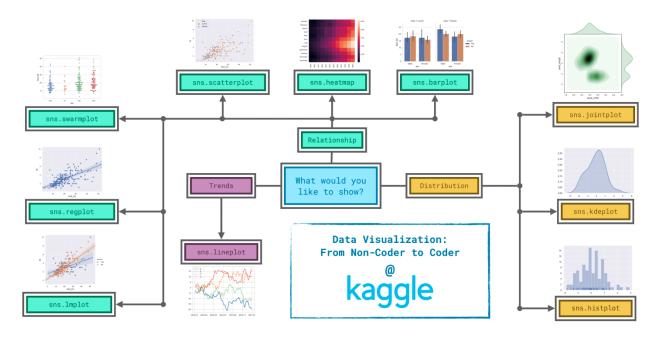
Data Visualization

https://www.kaggle.com/code

https://www.kaggle.com/datasets



Since it's not always easy to decide how to best tell the story behind your data, we've broken the chart types into three broad categories to help with this.

- Trends A trend is defined as a pattern of change.
 - sns.lineplot **Line charts** are best to show trends over a period of time, and multiple lines can be used to show trends in more than one group.
- **Relationship** There are many different chart types that you can use to understand relationships between variables in your data.
 - sns.barplot Bar charts are useful for comparing quantities corresponding to different groups.
 - sns.heatmap Heatmaps can be used to find color-coded patterns in tables of numbers.
 - sns.scatterplot Scatter plots show the relationship between two continuous variables; if color-coded, we can also show the relationship with a third <u>categorical</u> variable.
 - sns.regplot Including a **regression line** in the scatter plot makes it easier to see any linear relationship between two variables.
 - sns.lmplot This command is useful for drawing multiple regression lines, if the scatter plot contains multiple, color-coded groups.
 - sns.swarmplot Categorical scatter plots show the relationship between a continuous variable and a categorical variable.

- **Distribution** We visualize distributions to show the possible values that we can expect to see in a variable, along with how likely they are.
 - sns.histplot **Histograms** show the distribution of a single numerical variable.
 - sns.kdeplot KDE plots (or 2D KDE plots) show an estimated, smooth distribution of a single numerical variable (or two numerical variables).
 - sns.jointplot This command is useful for simultaneously displaying a 2D KDE plot with the corresponding KDE plots for each individual variable.

Setting up the notebook:

```
import pandas as pd
pd.plotting.register_matplotlib_converters()
import matplotlib.pyplot as plt
%matplotlib inline
import seaborn as sns
print("Setup Complete")
```

Load the data

```
# Path of the file to read
fifa_filepath = "../input/fifa.csv"

# Read the file into a variable fifa_data
fifa_data = pd.read_csv(fifa_filepath, index_col="Date", parse_dates=True)
```

Examine the data

```
# Print the first 5 rows of the data
fifa_data.head()
```

Lineplot

```
# Set the width and height of the figure
plt.figure(figsize=(16,6))
# Line chart showing how FIFA rankings evolved over time
sns.lineplot(data=fifa_data)
```

For instance, we use sns.lineplot to make line charts. Soon, you'll learn that we use sns.barplot and sns.heatmap to make bar charts and heatmaps, respectively.

Plot a subset of the data

```
list(spotify_data.columns)
# Set the width and height of the figure
plt.figure(figsize=(14,6))

# Add title
plt.title("Daily Global Streams of Popular Songs in 2017-2018")

# Line chart showing daily global streams of 'Shape of You'
sns.lineplot(data=spotify_data['Shape of You'], label="Shape of You")

# Line chart showing daily global streams of 'Despacito'
sns.lineplot(data=spotify_data['Despacito'], label="Despacito")

# Add label for horizontal axis
plt.xlabel("Date")
```

Setting theme:

```
# Change the style of the figure to the "dark" theme
sns.set_style("dark")
Seaborn has five different themes: (1)"darkgrid", (2)"whitegrid", (3)"dark", (4)"white",
and (5)"ticks",
```

Bar chart

```
# Set the width and height of the figure
plt.figure(figsize=(10,6))

# Add title
plt.title("Average Arrival Delay for Spirit Airlines Flights, by Month")

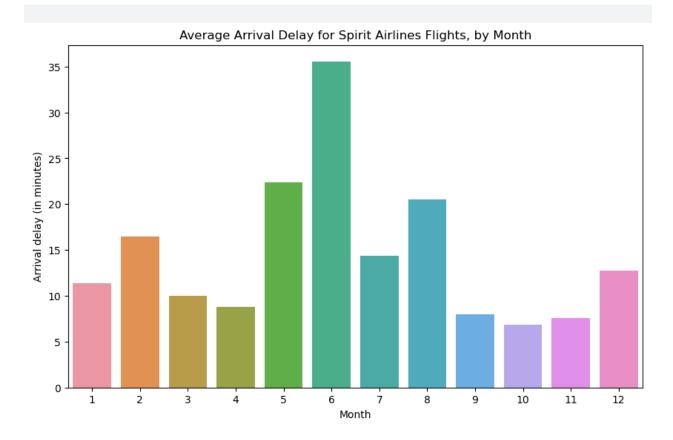
# Bar chart showing average arrival delay for Spirit Airlines flights by mo
nth
sns.barplot(x=flight_data.index, y=flight_data['NK'])

# Add label for vertical axis
plt.ylabel("Arrival delay (in minutes)")
It has three main components:
```

- sns.barplot This tells the notebook that we want to create a bar chart.
 - Remember that sns refers to the <u>seaborn</u> package, and all of the commands that you use to create charts in this course will start with this prefix.

- x=flight_data.index This determines what to use on the horizontal axis. In this case, we have selected the column that *index*es the rows (in this case, the column containing the months).
- y=flight_data['NK'] This sets the column in the data that will be used to determine the height of each bar. In this case, we select the 'NK' column.

Important Note: You must select the indexing column with flight_data.index, and it is not possible to use flight_data['Month'] (which will return an error). This is because when we loaded the dataset, the "Month" column was used to index the rows. We always have to use this special notation to select the indexing column.



Heatmap

```
# Set the width and height of the figure
plt.figure(figsize=(14,7))

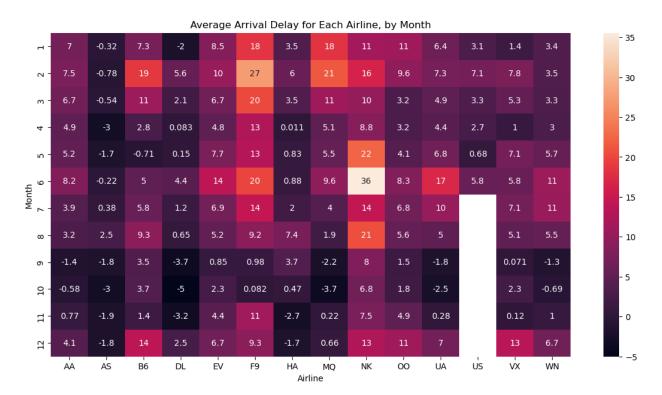
# Add title
plt.title("Average Arrival Delay for Each Airline, by Month")

# Heatmap showing average arrival delay for each airline by month
sns.heatmap(data=flight_data, annot=True)
```

```
# Add label for horizontal axis
plt.xlabel("Airline")
```

This code has three main components:

- sns.heatmap This tells the notebook that we want to create a heatmap.
- data=flight_data This tells the notebook to use all of the entries in flight_data to create the heatmap.
- annot=True This ensures that the values for each cell appear on the chart. (Leaving this out removes the numbers from each of the cells!)

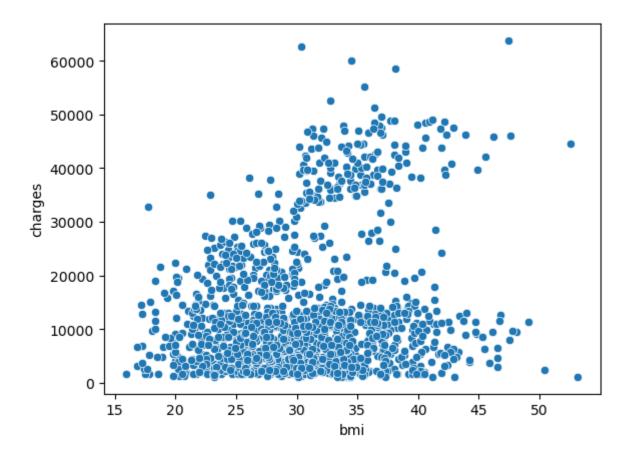


Scatter plots:

Usually, we use scatter plots to highlight the relationship between two continuous variables (like "bmi" and "charges")

To create a simple **scatter plot**, we use the sns.scatterplot command and specify the values for:

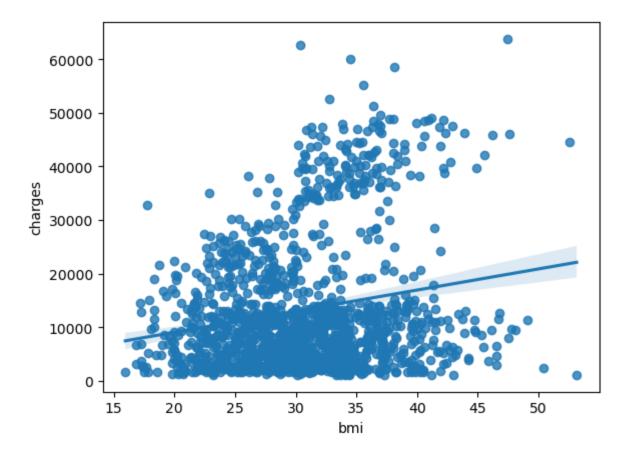
- the horizontal x-axis (x=insurance_data['bmi']), and
- the vertical y-axis (y=insurance_data['charges']).



The scatterplot above suggests that body mass index (BMI) and insurance charges are **positively correlated**, where customers with higher BMI typically also tend to pay more in insurance costs. (*This pattern makes sense, since high BMI is typically associated with higher risk of chronic disease.*)

To double-check the strength of this relationship, you might like to add a **regression line**, or the line that best fits the data. We do this by changing the command to sns.regplot.

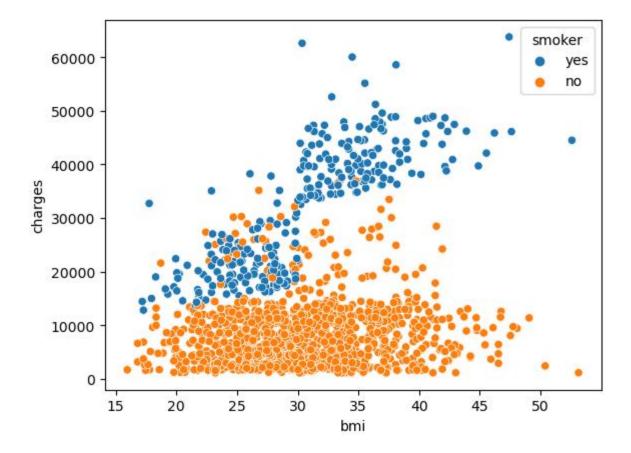
```
sns.regplot(x=insurance_data['bmi'], y=insurance_data['charges'])
```



Color-coded scatter plots

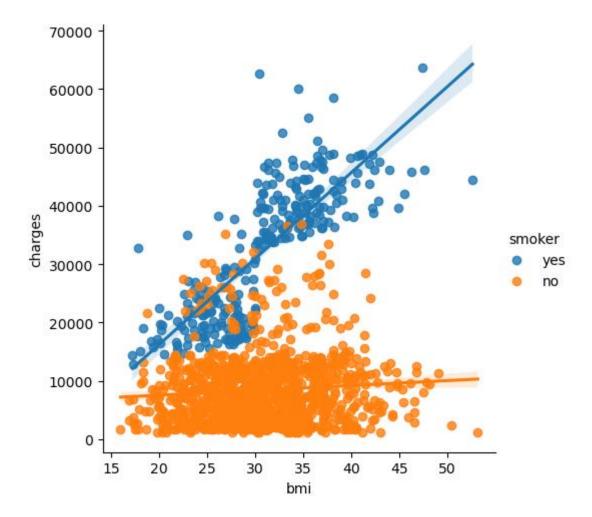
We can use scatter plots to display the relationships between (*not two, but...*) three variables! One way of doing this is by color-coding the points.

sns.scatterplot(x=insurance_data['bmi'], y=insurance_data['charges'], hue=
insurance_data['smoker'])



To further emphasize this fact, we can use the sns.lmplot command to add two regression lines, corresponding to smokers and nonsmokers. (You'll notice that the regression line for smokers has a much steeper slope, relative to the line for nonsmokers!)

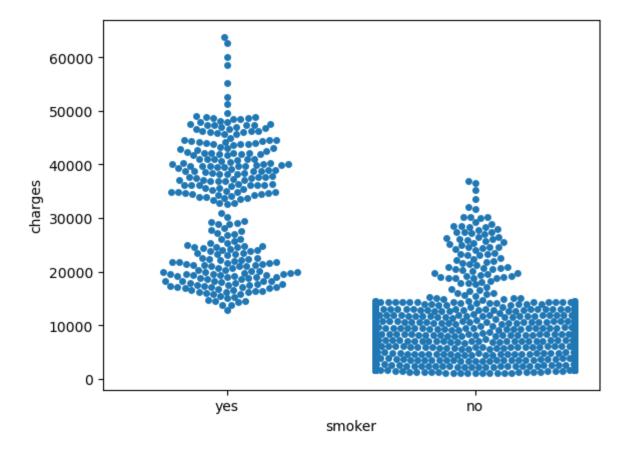
```
sns.lmplot(x="bmi", y="charges", hue="smoker", data=insurance_data)
```



The sns.lmplot command above works slightly differently than the commands you have learned about so far:

- Instead of setting x=insurance_data['bmi'] to select the 'bmi' column in insurance_data, we set x="bmi" to specify the name of the column only.
- Similarly, y="charges" and hue="smoker" also contain the names of columns.
- We specify the dataset with data=insurance_data

we can adapt the design of the scatter plot to feature a categorical variable (like "smoker") on one of the main axes. We'll refer to this plot type as a **Categorical scatter plot**, and we build it with the sns.swarmplot command.

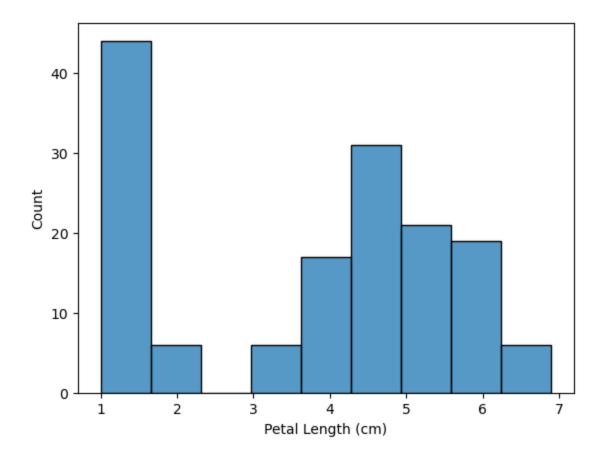


Among other things, this plot shows us that:

- on average, non-smokers are charged less than smokers, and
- the customers who pay the most are smokers; whereas the customers who pay the least are non-smokers.

Histograms

```
# Histogram
sns.histplot(iris_data['Petal Length (cm)'])
```

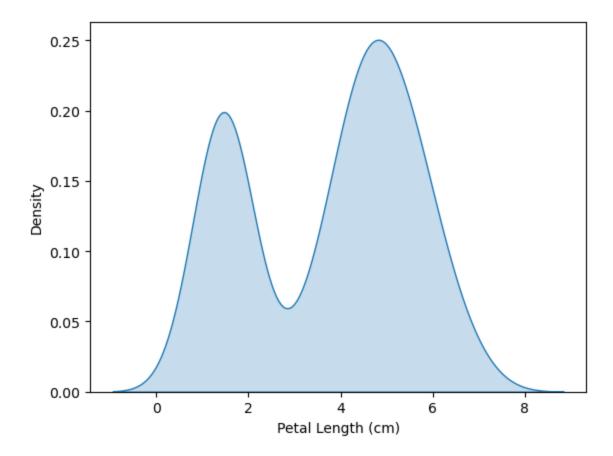


Density plots

The next type of plot is a **kernel density estimate (KDE)** plot. In case you're not familiar with KDE plots, you can think of it as a smoothed histogram.

To make a KDE plot, we use the sns.kdeplot command. Setting shade=True colors the area below the curve (and data= chooses the column we would like to plot).

```
# KDE plot
sns.kdeplot(data=iris_data['Petal Length (cm)'], shade=True)
```

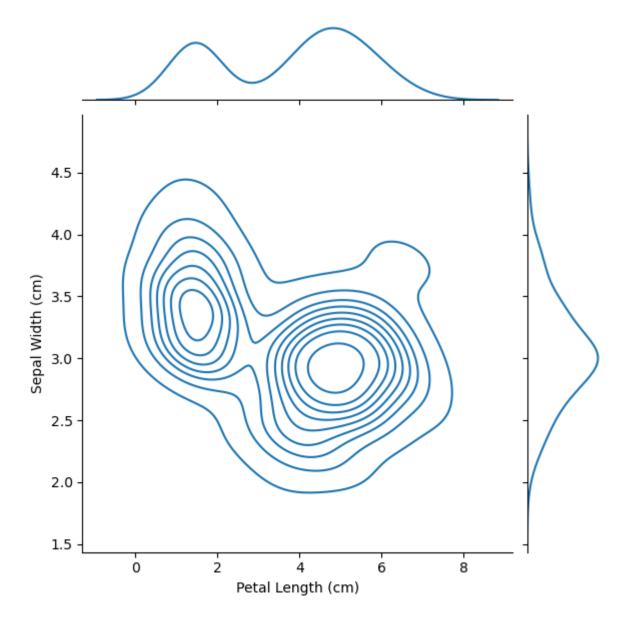


2D KDE plots

We're not restricted to a single column when creating a KDE plot. We can create a **two-dimensional (2D) KDE plot** with the sns.jointplot command.

In the plot below, the color-coding shows us how likely we are to see different combinations of sepal width and petal length, where darker parts of the figure are more likely.

```
In [5]:
# 2D KDE plot
sns.jointplot(x=iris_data['Petal Length (cm)'], y=iris_data['Sepal Width (
cm)'], kind="kde")
```

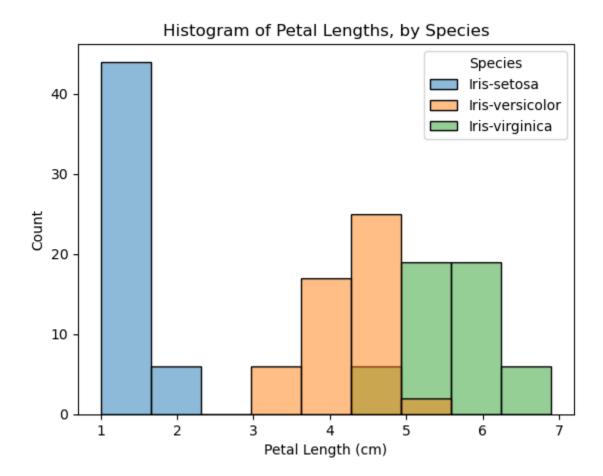


Color-coded plots

We can create three different histograms (one for each species) of petal length by using the sns.histplot command (as above).

- data= provides the name of the variable that we used to read in the data
- x= sets the name of column with the data we want to plot
- hue= sets the column we'll use to split the data into different histograms

```
# Histograms for each species
sns.histplot(data=iris_data, x='Petal Length (cm)', hue='Species')
```



We can also create a KDE plot for each species by using sns.kdeplot (as above). The functionality for data, x, and hue are identical to when we used sns.histplot above. Additionally, we set shade=True to color the area below each curve.

```
# KDE plots for each species
sns.kdeplot(data=iris_data, x='Petal Length (cm)', hue='Species', shade=Tr
ue)
# Add title
plt.title("Distribution of Petal Lengths, by Species")
```



