Making informative visualizations (sometimes called plots) is one of the most important tasks in data analysis. It may be a part of the exploratory process—for example, to help identify outliers or needed data transformations, or as a way of generating ideas for models. For others, building an interactive visualization for the web may be the end goal. Python has many add-on libraries for making static or dynamic visualizations, but I'll be mainly focused on matplotlib and libraries that build on top of it.

matplotlib is a desktop plotting package designed for creating (mostly two dimensional) publication-quality plots. The project was started by John Hunter in 2002 to enable a MATLAB-like plotting interface in Python. The matplotlib and IPython communities have collaborated to simplify interactive plotting from the IPython shell (and now, Jupyter notebook). matplotlib supports various GUI backends on all operating systems and additionally can export visualizations to all of the common vector and raster graphics formats (PDF, SVG, JPG, PNG, BMP, GIF, etc.). With the exception of a few diagrams, nearly all of the graphics in this book were produced using matplotlib.

Over time, matplotlib has spawned a number of add-on toolkits for data visualization that use matplotlib for their underlying plotting. One of these is seaborn, which we explore later in this chapter.

The simplest way to follow the code examples in the chapter is to use interactive plotting in the Jupyter notebook. To set this up, execute the following statement in a Jupyter notebook:

In [1]: %matplotlib notebook

### 9.1 A Brief matplotlib API Primer

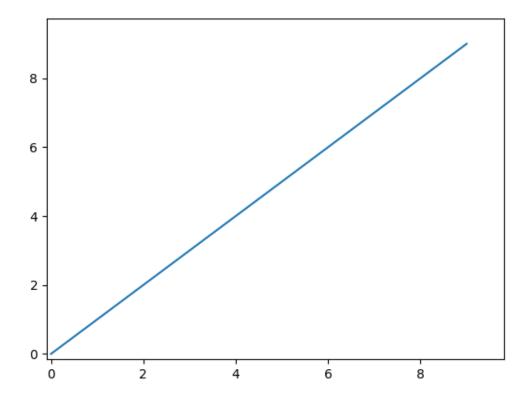
With matplotlib, we use the following import convention:

```
In [2]: import matplotlib.pyplot as plt
```

After running %matplotlib notebook in Jupyter (or simply %matplotlib in IPython), we can try creating a simple plot. If everything is set up right, a line plot like Figure 9-1 should appear:

```
In [3]: import numpy as np
    data = np.arange(10)
    data
Out[3]: array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
```

```
In [5]: plt.plot(data)
```

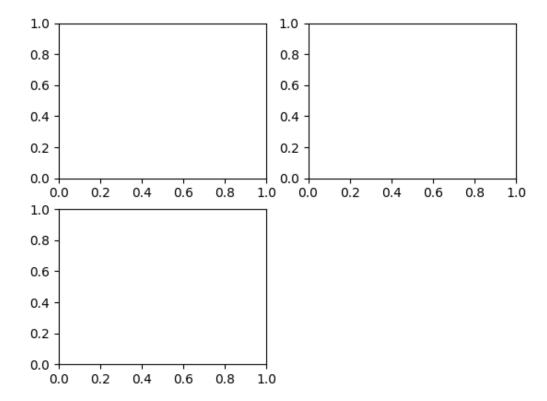


Out[5]: [<matplotlib.lines.Line2D at 0x188fda6fc50>]

While libraries like seaborn and pandas's built-in plotting functions will deal with many of the mundane details of making plots, should you wish to customize them beyond the function options provided, you will need to learn a bit about the matplotlib API.

### **Figures and Subplots**

Plots in matplotlib reside within a Figure object. You can create a new figure with plt.figure:



In IPython, an empty plot window will appear, but in Jupyter nothing will be shown until we use a few more commands. plt.figure has a number of options; notably, figsize will guarantee the figure has a certain size and aspect ratio if saved to disk.

You can't make a plot with a blank figure. You have to create one or more subplots using add\_subplot:

```
In [8]: ax1 = fig.add_subplot(2, 2, 1)
```

This means that the figure should be  $2 \times 2$  (so up to four plots in total), and we're selecting the first of four subplots (numbered from 1). If you create the next two subplots, you'll end up with a visualization that looks like Figure 9-2

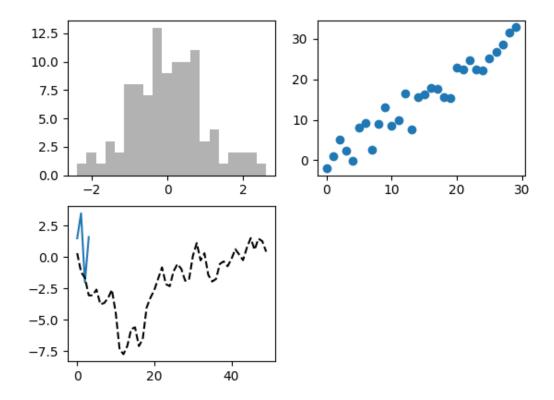
```
In [9]: ax2 = fig.add_subplot(2, 2, 2)
ax3 = fig.add_subplot(2, 2, 3)
```

One nuance of using Jupyter notebooks is that plots are reset after each cell is evaluated, so for more complex plots you must put all of the plotting commands in a single notebook cell.

Here we run all of these commands in the same cell.

```
In [10]: fig = plt.figure()
ax1 = fig.add_subplot(2, 2, 1)
```

```
ax2 = fig.add_subplot(2, 2, 2)
ax3 = fig.add_subplot(2, 2, 3)
```



When you issue a plotting command like plt.plot([1.5, 3.5, -2, 1.6]), matplotlib draws on the last figure and subplot used (creating one if necessary), thus hiding the figure and subplot creation. So if we add the following command, you'll get something like Figure 9-3:

```
In [11]: plt.plot([1.5, 3.5, -2, 1.6])
Out[11]: [<matplotlib.lines.Line2D at 0x188fe5be438>]
In [12]: plt.plot(np.random.randn(50).cumsum(), 'k--')
Out[12]: [<matplotlib.lines.Line2D at 0x188fe27b198>]
```

The 'k--' is a style option instructing matplotlib to plot a black dashed line. The objects returned by fig.add\_subplot here are AxesSubplot objects, on which you can directly plot on the other empty subplots by calling each one's instance method (see Figure 9-4):

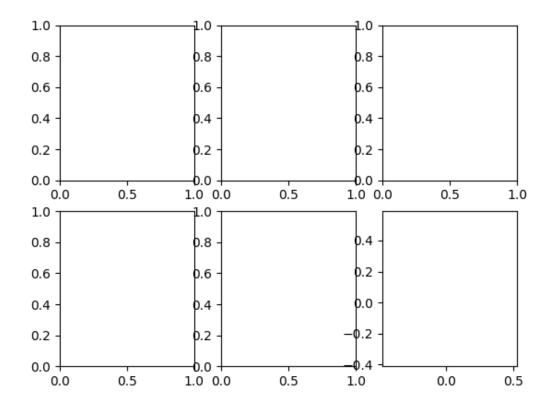
```
In [13]: _ = ax1.hist(np.random.randn(100), bins=20, color='k', alpha=0.3)
ax2.scatter(np.arange(30), np.arange(30) + 3 * np.random.randn(30))
```

Out[13]: <matplotlib.collections.PathCollection at 0x188fe5d4780>

You can find a comprehensive catalog of plot types in the matplotlib documentation.

Creating a figure with a grid of subplots is a very common task, so matplotlib includes a convenience method, plt.subplots, that creates a new figure and returns a NumPy array containing the created subplot objects

```
In [14]: fig, axes = plt.subplots(2, 3)
axes
```



This is very useful, as the axes array can be easily indexed like a two-dimensional array; for example, axes[0, 1]. You can also indicate that subplots should have the same x- or y-axis using sharex and sharey, respectively. This is especially useful when you're comparing data on the same scale; otherwise, matplotlib autoscales plot limits independently. See Table 9-1 for more on this method

### Table 9-1. pyplot.subplots options

Argument --> Description

nrows --> Number of rows of subplots

ncols --> Number of columns of subplots

sharex --> All subplots should use the same x-axis ticks (adjusting the xlim will affect all subplots)

sharey --> All subplots should use the same y-axis ticks (adjusting the ylim will affect all subplots)

subplot\_kw --> Dict of keywords passed to add\_subplot call used to create each subplot

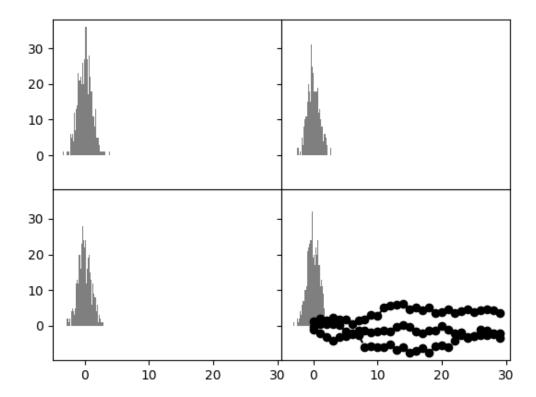
\*\*fig\_kw --> Additional keywords to subplots are used when creating the figure, such as plt.subplots(2, 2, figsize=(8, 6))

### Adjusting the spacing around subplots

By default matplotlib leaves a certain amount of padding around the outside of the subplots and spacing between subplots. This spacing is all specified relative to the height and width of the plot, so that if you resize the plot either programmatically or manually using the GUI window, the plot will dynamically adjust itself. You can change the spacing using the subplots\_adjust method on Figure objects, also available as a top-level function:

wspace and hspace controls the percent of the figure width and figure height, respectively, to use as spacing between subplots. Here is a small example where I shrink the spacing all the way to zero (see Figure 9-5):

```
In [16]:
    fig, axes = plt.subplots(2, 2, sharex=True, sharey=True)
    for i in range(2):
        for j in range(2):
            axes[i, j].hist(np.random.randn(500), bins=50, color='k', alpha=0.5)
    plt.subplots_adjust(wspace=0, hspace=0)
```



You may notice that the axis labels overlap. matplotlib doesn't check whether the labels overlap, so in a case like this you would need to fix the labels yourself by specifying explicit tick locations and tick labels (we'll look at how to do this in the following sections).

# Colors, Markers, and Line Styles

Matplotlib's main plot function accepts arrays of x and y coordinates and optionally a string abbreviation indicating color and line style. For example, to plot x versus y with green dashes, you would execute:

```
In [19]: # ax.plot(x, y, 'g--')
```

This way of specifying both color and line style in a string is provided as a convenience; in practice if you were creating plots programmatically you might prefer not to have to munge strings together to create plots with the desired style. The same plot could also have been expressed more explicitly as:

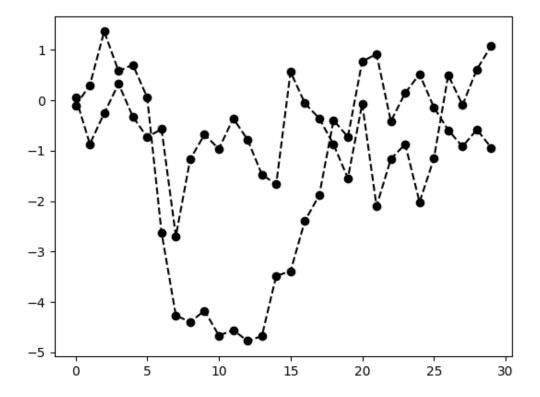
```
In [18]: # ax.plot(x, y, linestyle='--', color='g')
```

There are a number of color abbreviations provided for commonly used colors, but you can use any color on the spectrum by specifying its hex code (e.g., '#CECECE'). You can see the full set of line styles by looking at the docstring for plot (use plot? in IPython or Jupyter).

```
In [22]: plt?
```

Line plots can additionally have markers to highlight the actual data points. Since matplotlib creates a continuous line plot, interpolating between points, it can occasionally be unclear where the points lie. The marker can be part of the style string, which must have color followed by marker type and line style (see Figure 9-6):

```
In [27]: from numpy.random import randn
  import matplotlib.pyplot as plt
  fig = plt.figure()
  plt.plot(randn(30).cumsum(), 'ko--')
```



Out[27]: [<matplotlib.lines.Line2D at 0x188ffd181d0>]

This could also have been written more explicitly as:

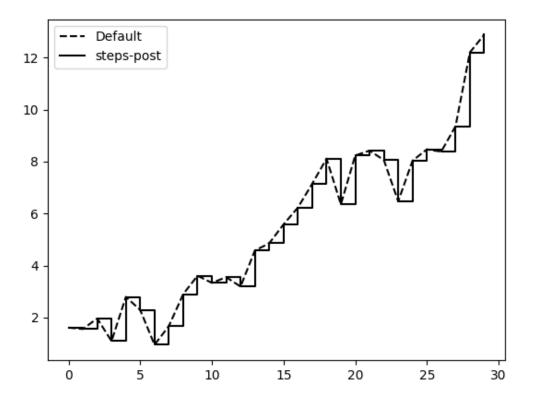
```
In [33]: # #import matplotlib.puy
# matplotlib.pyplot.plot(randn(30).cumsum(), color='k', linestyle='dashed', marker=
```

Out[33]: [<matplotlib.lines.Line2D at 0x188800929b0>]

For line plots, you will notice that subsequent points are linearly interpolated by default. This can be altered with the drawstyle option (Figure 9-7):

```
In [34]: data = np.random.randn(30).cumsum()
    fig=plt.figure()
```

```
plt.plot(data, 'k--', label='Default')
```



```
Out[34]: [<matplotlib.lines.Line2D at 0x188818e2278>]
```

```
In [35]: plt.plot(data, 'k-', drawstyle='steps-post', label='steps-post')
```

Out[35]: [<matplotlib.lines.Line2D at 0x188800ba0b8>]

```
In [36]: plt.legend(loc='best')
```

Out[36]: <matplotlib.legend.Legend at 0x18881905358>

You may notice output like <matplotlib.lines.Line2D at ...> when you run this. matplotlib returns objects that reference the plot subcomponent that was just added. A lot of the time you can safely ignore this output. Here, since we passed the label arguments to plot, we are able to create a plot legend to identify each line using plt.legend.

You must call plt.legend (or ax.legend, if you have a reference to the axes) to create the legend, whether or not you passed the label options when plotting the data.

### Ticks, Labels, and Legends

For most kinds of plot decorations, there are two main ways to do things: using the procedural pyplot interface (i.e., matplotlib.pyplot) and the more object-oriented native matplotlib API.

The pyplot interface, designed for interactive use, consists of methods like xlim, xticks, and xticklabels. These control the plot range, tick locations, and tick labels, respectively. They can be used in two ways

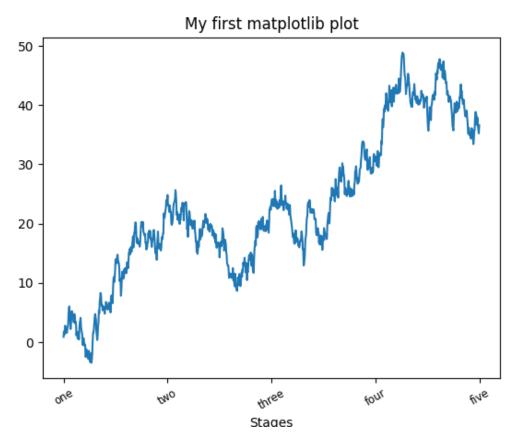
- Called with no arguments returns the current parameter value (e.g., plt.xlim() returns the current x-axis plotting range)
- Called with parameters sets the parameter value (e.g., plt.xlim([0, 10]), sets the x-axis range to 0 to 10)

All such methods act on the active or most recently created AxesSubplot. Each of them corresponds to two methods on the subplot object itself; in the case of xlim these are ax.get\_xlim and ax.set\_xlim. I prefer to use the subplot instance methods myself in the interest of being explicit (and especially when working with multiple subplots), but you can certainly use whichever you find more convenient.

#### Setting the title, axis labels, ticks, and ticklabels

To illustrate customizing the axes, I'll create a simple figure and plot of a random walk (see Figure 9-8):

```
In [37]: fig = plt.figure()
    ax = fig.add_subplot(1, 1, 1)
    ax.plot(np.random.randn(1000).cumsum())
```



Out[37]: [<matplotlib.lines.Line2D at 0x18881cba780>]

To change the x-axis ticks, it's easiest to use set\_xticks and set\_xticklabels. The former instructs matplotlib where to place the ticks along the data range; by default these locations will also be the labels. But we can set any other values as the labels using set\_xticklabels:

```
In [38]: ticks = ax.set_xticks([0, 250, 500, 750, 1000])
labels = ax.set_xticklabels(['one', 'two', 'three', 'four', 'five'],rotation=30, four
```

The rotation option sets the x tick labels at a 30-degree rotation. Lastly, set\_xlabel gives a name to the x-axis and set\_title the subplot title (see Figure 9-9 for the resulting figure):

```
In [39]: ax.set_title('My first matplotlib plot')
Out[39]: Text(0.5, 1.0, 'My first matplotlib plot')
In [40]: ax.set_xlabel('Stages')
Out[40]: Text(0.5, 10.888891973024519, 'Stages')
```

Modifying the y-axis consists of the same process, substituting y for x in the above. The axes class has a set method that allows batch setting of plot properties. From the prior example, we could also have written:

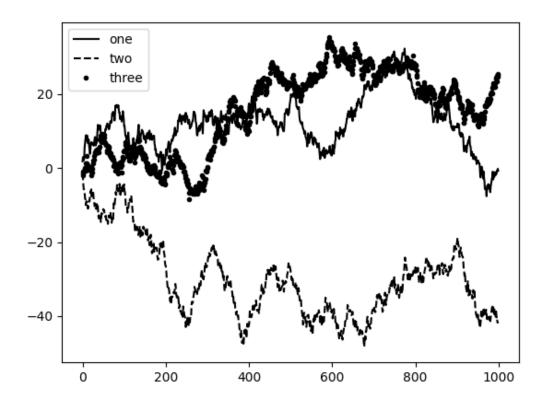
```
In [41]: props = {
   'title': 'My first matplotlib plot',
   'xlabel': 'Stages'
}
ax.set(**props)
```

```
Out[41]: [Text(0.5, 1.0, 'My first matplotlib plot'),
Text(0.5, 10.888891973024519, 'Stages')]
```

#### **Adding legends**

Legends are another critical element for identifying plot elements. There are a couple of ways to add one. The easiest is to pass the label argument when adding each piece of the plot:

```
In [42]: from numpy.random import randn
    fig = plt.figure(); ax = fig.add_subplot(1, 1, 1)
    ax.plot(randn(1000).cumsum(), 'k', label='one')
    ax.plot(randn(1000).cumsum(), 'k--', label='two')
    ax.plot(randn(1000).cumsum(), 'k.', label='three')
```



Out[42]: [<matplotlib.lines.Line2D at 0x1888069b278>]

Once you've done this, you can either call ax.legend() or plt.legend() to automatically create a legend. The resulting plot is in Figure 9-10:

```
In [43]: ax.legend(loc='best')
```

Out[43]: <matplotlib.legend.Legend at 0x18881bc78d0>

The legend method has several other choices for the location loc argument. See the docstring (with ax.legend?) for more information.

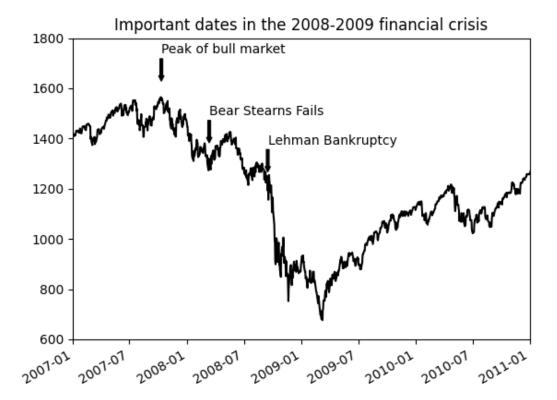
The loc tells matplotlib where to place the plot. If you aren't picky, 'best' is a good option, as it will choose a location that is most out of the way. To exclude one or more elements from the legend, pass no label or label='nolegend'.

### Annotations and Drawing on a Subplot

In addition to the standard plot types, you may wish to draw your own plot annotations, which could consist of text, arrows, or other shapes. You can add annotations and text using the text, arrow, and annotate functions. text draws text at given coordinates (x, y) on the plot with optional custom styling

Annotations can draw both text and arrows arranged appropriately. As an example, let's plot the closing S&P 500 index price since 2007 (obtained from Yahoo! Finance) and annotate it with some of the important dates from the 2008–2009 financial crisis. You can most easily reproduce this code example in a single cell in a Jupyter notebook. See Figure 9-11 for the result:

```
In [47]: import pandas as pd
         from datetime import datetime
         fig = plt.figure()
         ax = fig.add_subplot(1, 1, 1)
         data = pd.read_csv('spx.csv', index_col=0, parse_dates=True)
         spx = data['SPX']
         spx.plot(ax=ax, style='k-')
         crisis_data = [
          (datetime(2007, 10, 11), 'Peak of bull market'),
          (datetime(2008, 3, 12), 'Bear Stearns Fails'),
          (datetime(2008, 9, 15), 'Lehman Bankruptcy')
         for date, label in crisis data:
             ax.annotate(label, xy=(date, spx.asof(date) + 75),xytext=(date, spx.asof(date)
             horizontalalignment='left', verticalalignment='top')
         # Zoom in on 2007-2010
         ax.set_xlim(['1/1/2007', '1/1/2011'])
         ax.set ylim([600, 1800])
         ax.set_title('Important dates in the 2008-2009 financial crisis')
```



Out[47]: Text(0.5, 1.0, 'Important dates in the 2008-2009 financial crisis')

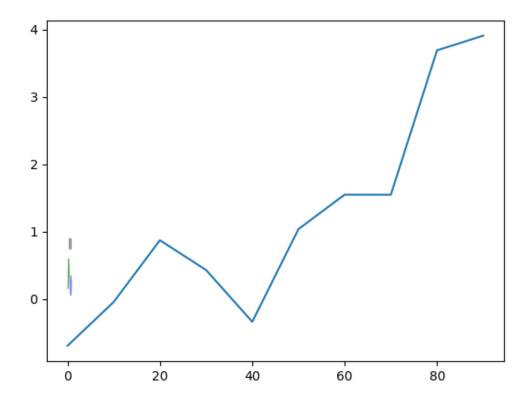
There are a couple of important points to highlight in this plot: the ax.annotate method can draw labels at the indicated x and y coordinates. We use the set\_xlim and set\_ylim methods to manually set the start and end boundaries for the plot rather than using matplotlib's default. Lastly, ax.set\_title adds a main title to the plot.

See the online matplotlib gallery for many more annotation examples to learn from.

Drawing shapes requires some more care. matplotlib has objects that represent many common shapes, referred to as patches. Some of these, like Rectangle and Circle, are found in matplotlib.pyplot, but the full set is located in matplotlib.patches.

To add a shape to a plot, you create the patch object shp and add it to a subplot by calling ax.add\_patch(shp) (see Figure 9-12):

```
In [48]: fig = plt.figure()
    ax = fig.add_subplot(1, 1, 1)
    rect = plt.Rectangle((0.2, 0.75), 0.4, 0.15, color='k', alpha=0.3)
    circ = plt.Circle((0.7, 0.2), 0.15, color='b', alpha=0.3)
    pgon = plt.Polygon([[0.15, 0.15], [0.35, 0.4], [0.2, 0.6]],
        color='g', alpha=0.5)
    ax.add_patch(rect)
    ax.add_patch(circ)
    ax.add_patch(pgon)
```



Out[48]: <matplotlib.patches.Polygon at 0x1888365d8d0>

If you look at the implementation of many familiar plot types, you will see that they are assembled from patches.

### Saving Plots to File

You can save the active figure to file using plt.savefig. This method is equivalent to the figure object's savefig instance method. For example, to save an SVG version of a figure, you need only type:

```
In [49]: plt.savefig('figpath.svg')
```

The file type is inferred from the file extension. So if you used .pdf instead, you would get a PDF. There are a couple of important options that I use frequently for publishing graphics: dpi, which controls the dots-per-inch resolution, and bbox\_inches, which can trim the whitespace around the actual figure. To get the same plot as a PNG with minimal whitespace around the plot and at 400 DPI, you would do:

```
In [50]: plt.savefig('figpath.png', dpi=400, bbox_inches='tight')
```

savefig doesn't have to write to disk; it can also write to any file-like object, such as a BytesIO:

```
In [51]: from io import BytesIO
    buffer = BytesIO()
    plt.savefig(buffer)
    plot_data = buffer.getvalue()
In [53]: # plot data
```

# Table 9-2. Figure.savefig options

Argument --> Description

fname --> String containing a filepath or a Python file-like object. The figure format is inferred from the file extension (e.g., .pdf for PDF or .png for PNG)

dpi --> The figure resolution in dots per inch; defaults to 100 out of the box but can be configured

facecolor,edgecolor --> The color of the figure background outside of the subplots; 'w' (white), by default

format --> The explicit file format to use ('png', 'pdf', 'svg', 'ps', 'eps', ...)

bbox\_inches --> The portion of the figure to save; if 'tight' is passed, will attempt to trim the empty space around the figure

### matplotlib Configuration

matplotlib comes configured with color schemes and defaults that are geared primarily toward preparing figures for publication. Fortunately, nearly all of the default behavior can be customized via an extensive set of global parameters governing figure size, subplot spacing, colors, font sizes, grid styles, and so on. One way to modify the configuration programmatically from Python is to use the rc method; for example, to set the global default figure size to be  $10 \times 10$ , you could enter:

```
In [54]: plt.rc('figure', figsize=(10, 10))
```

The first argument to rc is the component you wish to customize, such as 'figure', 'axes', 'xtick', 'ytick', 'grid', 'legend', or many others. After that can follow a sequence of keyword arguments indicating the new parameters. An easy way to write down the options in your program is as a dict:

```
In [57]: font_options = {'family' : 'monospace',
    'weight' : 'bold'}
plt.rc('font', **font_options)
```

For more extensive customization and to see a list of all the options, matplotlib comes with a configuration file matplotlibrc in the matplotlib/mpl-data directory. If you customize this file and place it in your home directory titled .matplotlibrc, it will be loaded each time you use matplotlib.

As we'll see in the next section, the seaborn package has several built-in plot themes or styles that use matplotlib's configuration system internally.

## 9.2 Plotting with pandas and seaborn

matplotlib can be a fairly low-level tool. You assemble a plot from its base components: the data display (i.e., the type of plot: line, bar, box, scatter, contour, etc.), leg- end, title, tick labels, and other annotations.

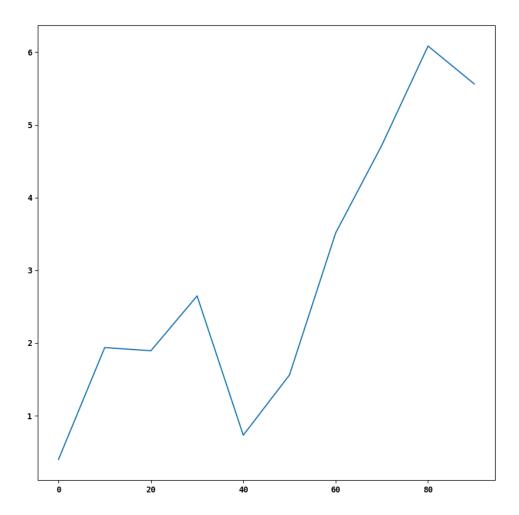
In pandas we may have multiple columns of data, along with row and column labels. pandas itself has built-in methods that simplify creating visualizations from DataFrame and Series objects. Another library is seaborn, a statistical graphics library created by Michael Waskom. Seaborn simplifies creating many common visualization types.

Importing seaborn modifies the default matplotlib color schemes and plot styles to improve readability and aesthetics. Even if you do not use the seaborn API, you may prefer to import seaborn as a simple way to improve the visual aesthetics of general matplotlib plots

#### **Line Plots**

Series and DataFrame each have a plot attribute for making some basic plot types. By default, plot() makes line plots (see Figure 9-13):

```
In [59]: fig = plt.figure()
s = pd.Series(np.random.randn(10).cumsum(), index=np.arange(0, 100, 10))
s.plot()
```

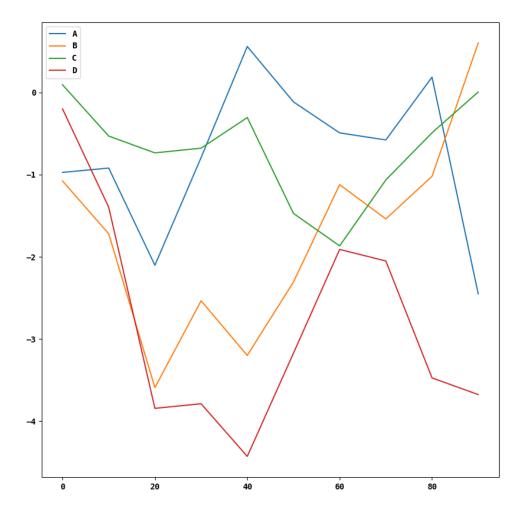


#### Out[59]: <AxesSubplot:>

The Series object's index is passed to matplotlib for plotting on the x-axis, though you can disable this by passing use\_index=False. The x-axis ticks and limits can be adjusted with the xticks and xlim options, and y-axis respectively with yticks and ylim. See Table 9-3 for a full listing of plot options. I'll comment on a few more of them throughout this section and leave the rest to you to explore.

Most of pandas's plotting methods accept an optional ax parameter, which can be a matplotlib subplot object. This gives you more flexible placement of subplots in a grid layout.

DataFrame's plot method plots each of its columns as a different line on the same subplot, creating a legend automatically (see Figure 9-14):



Out[60]: <AxesSubplot:>

The plot attribute contains a "family" of methods for different plot types. For example, df.plot() is equivalent to df.plot.line(). We'll explore some of these methods next.

Additional keyword arguments to plot are passed through to the respective matplotlib plotting function, so you can further custom- ize these plots by learning more about the matplotlib API.

## Table 9-3. Series.plot method arguments

Argument --> Description

label --> Label for plot legend

ax --> matplotlib subplot object to plot on; if nothing passed, uses active matplotlib subplot

```
style --> Style string, like 'ko--', to be passed to matplotlib
alpha --> The plot fill opacity (from 0 to 1)
kind --> Can be 'area', 'bar', 'barh', 'density', 'hist', 'kde', 'line', 'pie'
logy --> Use logarithmic scaling on the y-axis
use_index --> Use the object index for tick labels
rot --> Rotation of tick labels (0 through 360)
xticks --> Values to use for x-axis ticks
yticks --> Values to use for y-axis ticks
xlim --> x-axis limits (e.g., [0, 10])
ylim --> y-axis limits
grid --> Display axis grid (on by default)
```

DataFrame has a number of options allowing some flexibility with how the columns are handled; for example, whether to plot them all on the same subplot or to create separate subplots. See Table 9-4 for more on these.

### Table 9-4. DataFrame-specific plot arguments

```
Argument --> Description

subplots --> Plot each DataFrame column in a separate subplot

sharex --> If subplots=True, share the same x-axis, linking ticks and limits

sharey --> If subplots=True, share the same y-axis

figsize --> Size of figure to create as tuple

title --> Plot title as string

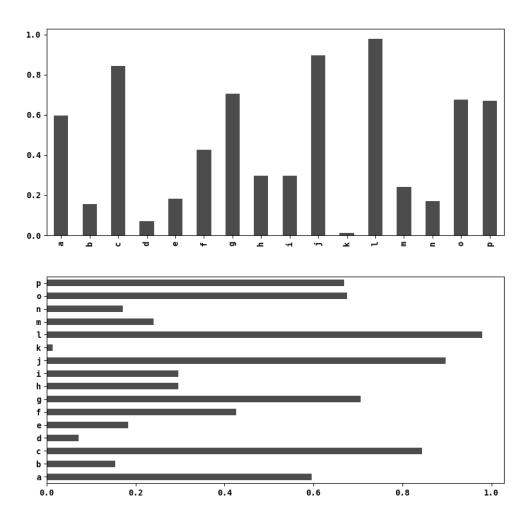
legend --> Add a subplot legend (True by default)

sort_columns --> Plot columns in alphabetical order; by defaul it uses existing column order
```

### **Bar Plots**

The plot.bar() and plot.barh() make vertical and horizontal bar plots, respectively. In this case, the Series or DataFrame index will be used as the x (bar) or y (barh) ticks (see Figure 9-15):

```
In [62]: fig, axes = plt.subplots(2, 1)
    data = pd.Series(np.random.rand(16), index=list('abcdefghijklmnop'))
    data.plot.bar(ax=axes[0], color='k', alpha=0.7)
```



```
Out[62]: <AxesSubplot:>
```

```
In [63]: data.plot.barh(ax=axes[1], color='k', alpha=0.7)
```

#### Out[63]: <AxesSubplot:>

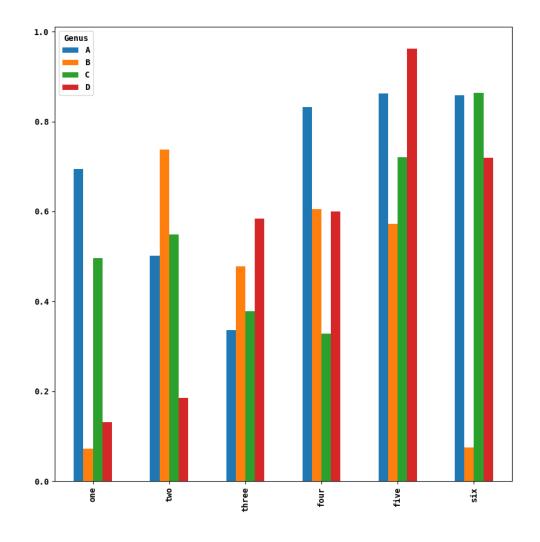
The options color='k' and alpha=0.7 set the color of the plots to black and use partial transparency on the filling.

With a DataFrame, bar plots group the values in each row together in a group in bars, side by side, for each value. See Figure 9-16:

```
In [64]: df = pd.DataFrame(np.random.rand(6, 4),index=['one', 'two', 'three', 'four', 'five'
df
```

Out[64]:	Genus	Α	В	С	D
	one	0.694788	0.072510	0.496193	0.131117
	two	0.502030	0.737705	0.549194	0.184443
	three	0.335535	0.477594	0.377460	0.584557
	four	0.831610	0.604899	0.328508	0.599296
	five	0.862400	0.572269	0.720580	0.962653
	six	0.858087	0.074653	0.864282	0.718909

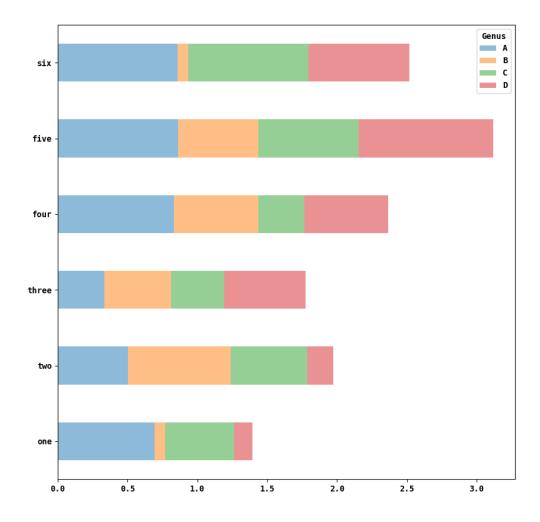
In [65]: df.plot.bar()



Out[65]: <AxesSubplot:>

Note that the name "Genus" on the DataFrame's columns is used to title the legend.

We create stacked bar plots from a DataFrame by passing stacked=True, resulting in the value in each row being stacked together (see Figure 9-17):



#### Out[67]: <AxesSubplot:>

A useful recipe for bar plots is to visualize a Series's value frequency using value\_counts: s.value\_counts().plot.bar().

Returning to the tipping dataset used earlier in the book, suppose we wanted to make a stacked bar plot showing the percentage of data points for each party size on each day. I load the data using read\_csv and make a cross-tabulation by day and party size:

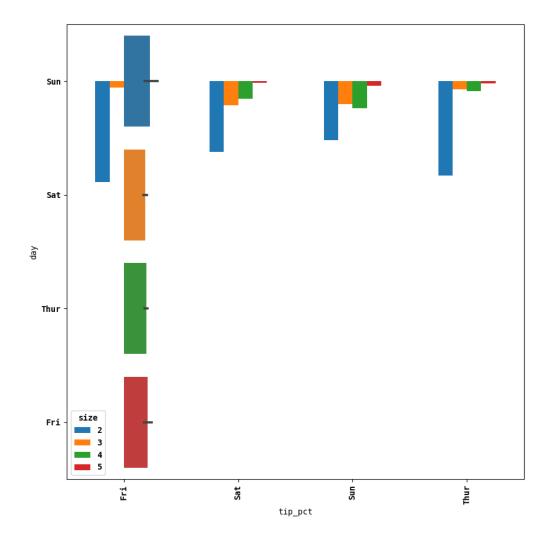
```
In [69]: tips = pd.read_csv('tips.csv')
    party_counts = pd.crosstab(tips['day'], tips['size'])
    party_counts
```

```
Out[69]:
          size 1
                  2
                      3
                          4 5 6
          day
           Fri
              1 16
                          1 0
                               0
          Sat
              2
                  53
                     18
                         13
          Sun
               0
                             3
                  39
                     15
                         18
         Thur 1 48
                          5 1 3
```

```
In [70]: party_counts = party_counts.loc[:, 2:5]
```

Then, normalize so that each row sums to 1 and make the plot (see Figure 9-18):

```
In [72]: party_pcts.plot.bar()
```



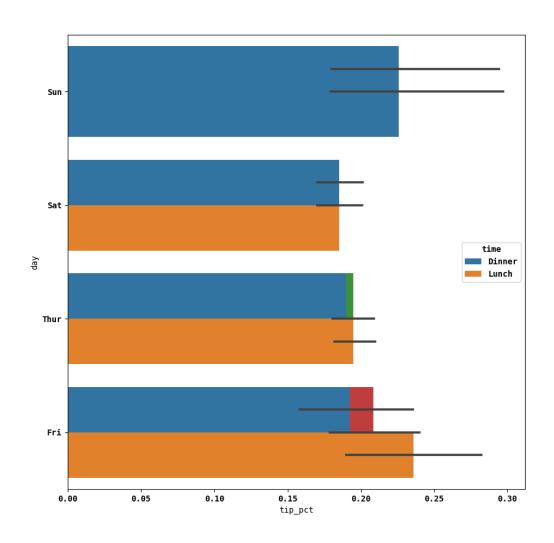
Out[72]: <AxesSubplot:xlabel='day'>

So you can see that party sizes appear to increase on the weekend in this dataset. With data that requires aggregation or summarization before making a plot, using the seaborn package can make things much simpler. Let's look now at the tipping percentage by day with seaborn (see Figure 9-19 for the resulting plot):

```
In [73]: import seaborn as sns
tips['tip_pct'] = tips['tip'] / (tips['total_bill'] - tips['tip'])
tips.head()
```

Out[73]: -		total_bill	tip	sex	smoker	day	time	size	tip_pct
	0	16.99	1.01	Female	No	Sun	Dinner	2	0.063204
	1	10.34	1.66	Male	No	Sun	Dinner	3	0.191244
	2	21.01	3.50	Male	No	Sun	Dinner	3	0.199886
	3	23.68	3.31	Male	No	Sun	Dinner	2	0.162494
	4	24.59	3.61	Female	No	Sun	Dinner	4	0.172069

```
In [76]: fig=plt.figure()
sns.barplot(x='tip_pct', y='day', data=tips, orient='h')
```



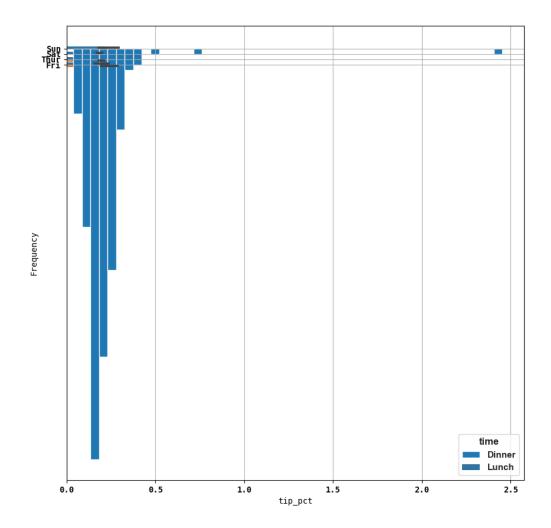
Out[76]: <AxesSubplot:xlabel='tip\_pct', ylabel='day'>

Plotting functions in seaborn take a data argument, which can be a pandas DataFrame. The other arguments refer to column names. Because there are multiple observations for each value in the day, the bars are the average value of tip\_pct. The black lines drawn on the bars

represent the 95% confidence interval (this can be configured through optional arguments).

seaborn.barplot has a hue option that enables us to split by an additional categorical value (Figure 9-20):

```
In [78]: fig=plt.figure()
sns.barplot(x='tip_pct', y='day', hue='time', data=tips, orient='h')
```



Out[78]: <AxesSubplot:xlabel='tip\_pct', ylabel='day'>

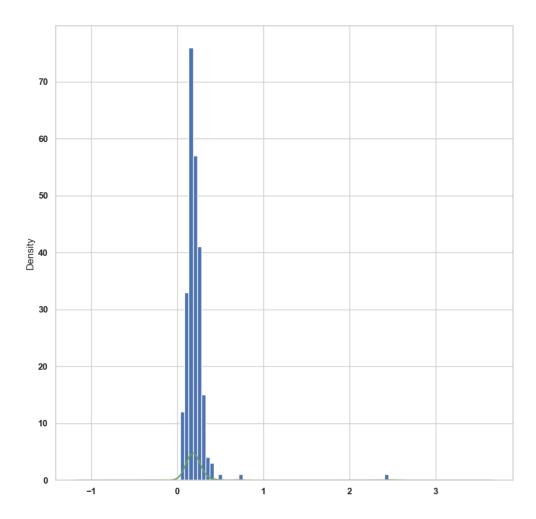
Notice that seaborn has automatically changed the aesthetics of plots: the default color palette, plot background, and grid line colors. You can switch between different plot appearances using seaborn.set:

```
In [79]: sns.set(style="whitegrid")
```

### **Histograms and Density Plots**

A histogram is a kind of bar plot that gives a discretized display of value frequency. The data points are split into discrete, evenly spaced bins, and the number of data points in each bin is plotted. Using the tipping data from before, we can make a histogram of tip percentages of the total bill using the plot.hist method on the Series (see Figure 9-21):

```
In [81]: fig=plt.figure()
tips['tip_pct'].plot.hist(bins=50)
```



Out[81]: <AxesSubplot:ylabel='Frequency'>

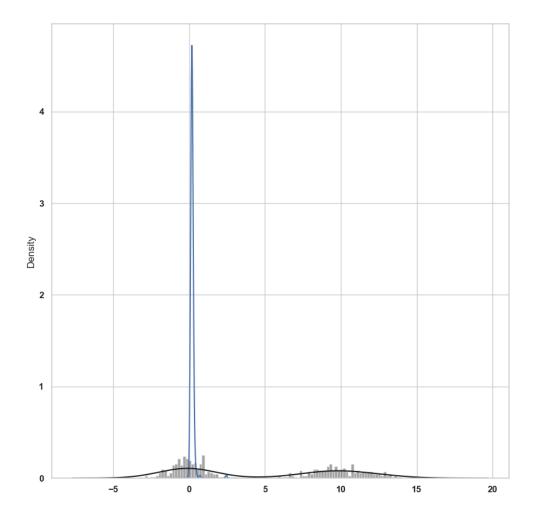
A related plot type is a density plot, which is formed by computing an estimate of a continuous probability distribution that might have generated the observed data. The usual procedure is to approximate this distribution as a mixture of "kernels"—that is, simpler distributions like the normal distribution. Thus, density plots are also known as kernel density

estimate (KDE) plots. Using plot.kde makes a density plot using the conventional mixture-of-normals estimate (see Figure 9-22):

```
In [84]: fig=plt.figure()
tips['tip_pct'].plot.density()
```

C:\Users\ankit19.gupta\OneDrive - Reliance Corporate IT Park Limited\Desktop\Practic e\_Code\Python\_Practice\Python\_For\_Data\_Analysis\myenv\lib\site-packages\ipykernel\_la uncher.py:1: RuntimeWarning: More than 20 figures have been opened. Figures created through the pyplot interface (`matplotlib.pyplot.figure`) are retained until explicitly closed and may consume too much memory. (To control this warning, see the rcPara m `figure.max\_open\_warning`).

"""Entry point for launching an IPython kernel.

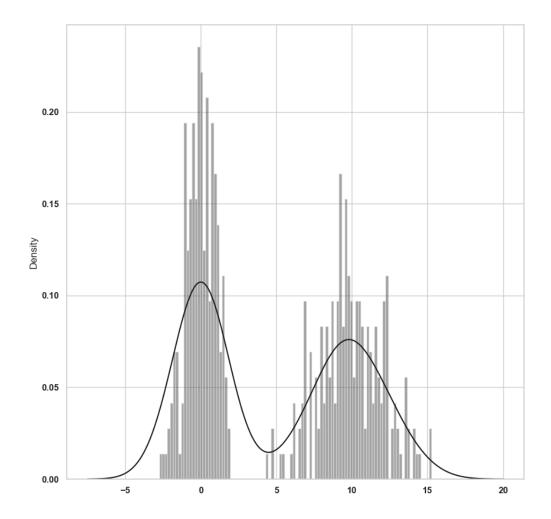


Out[84]: <AxesSubplot:ylabel='Density'>

Seaborn makes histograms and density plots even easier through its distplot method, which can plot both a histogram and a continuous density estimate simultaneously. As an example,

consider a bimodal distribution consisting of draws from two different standard normal distributions (see Figure 9-23)

```
In [86]: fig=plt.figure()
    comp1 = np.random.normal(0, 1, size=200)
    comp2 = np.random.normal(10, 2, size=200)
    values = pd.Series(np.concatenate([comp1, comp2]))
    sns.distplot(values, bins=100, color='k')
```



C:\Users\ankit19.gupta\OneDrive - Reliance Corporate IT Park Limited\Desktop\Practic e\_Code\Python\_Practice\Python\_For\_Data\_Analysis\myenv\lib\site-packages\seaborn\dist ributions.py:2619: FutureWarning: `distplot` is a deprecated function and will be re moved in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for h istograms).

warnings.warn(msg, FutureWarning)

Out[86]: <AxesSubplot:ylabel='Density'>

### **Scatter or Point Plots**

Point plots or scatter plots can be a useful way of examining the relationship between two one-dimensional data series. For example, here we load the macrodata dataset from the statsmodels project, select a few variables, then compute log differences:

```
In [87]: macro = pd.read_csv('macrodata.csv')
    data = macro[['cpi', 'm1', 'tbilrate', 'unemp']]
    trans_data = np.log(data).diff().dropna()
    trans_data[-5:]
Out[87]: cpi m1 tbilrate unemp
```

```
        cpi
        m1
        tbilrate
        unemp

        198
        -0.007904
        0.045361
        -0.396881
        0.105361

        199
        -0.021979
        0.066753
        -2.277267
        0.139762

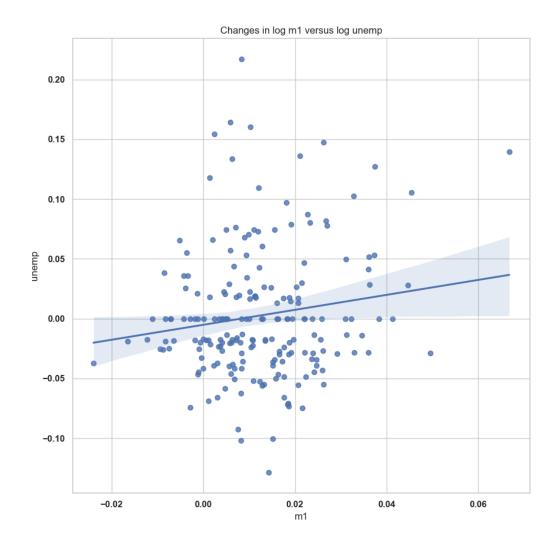
        200
        0.002340
        0.010286
        0.606136
        0.160343

        201
        0.008419
        0.037461
        -0.200671
        0.127339

        202
        0.008894
        0.012202
        -0.405465
        0.042560
```

We can then use seaborn's regplot method, which makes a scatter plot and fits a linear regression line (see Figure 9-24):

```
In [88]: fig=plt.figure()
    sns.regplot('m1', 'unemp', data=trans_data)
    plt.title('Changes in log %s versus log %s' % ('m1', 'unemp'))
```



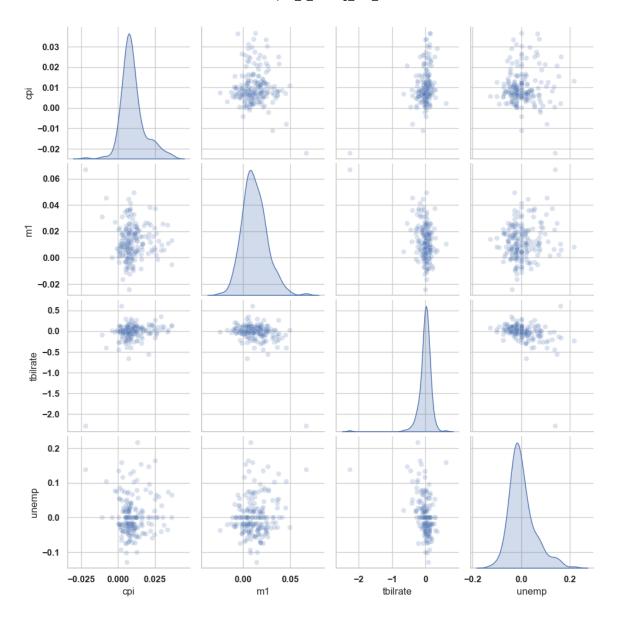
C:\Users\ankit19.gupta\OneDrive - Reliance Corporate IT Park Limited\Desktop\Practic e\_Code\Python\_Practice\Python\_For\_Data\_Analysis\myenv\lib\site-packages\seaborn\\_dec orators.py:43: FutureWarning: Pass the following variables as keyword args: x, y. Fr om version 0.12, the only valid positional argument will be `data`, and passing othe r arguments without an explicit keyword will result in an error or misinterpretatio n.

Out[88]: Text(0.5, 1.0, 'Changes in log m1 versus log unemp')

In exploratory data analysis it's helpful to be able to look at all the scatter plots among a group of variables; this is known as a pairs plot or scatter plot matrix. Making such a plot from scratch is a bit of work, so seaborn has a convenient pairplot function, which supports placing histograms or density estimates of each variable along the diagonal (see Figure 9-25 for the resulting plot):

```
In [90]: sns.pairplot(trans_data, diag_kind='kde', plot_kws={'alpha': 0.2})
```

FutureWarning



Out[90]: <seaborn.axisgrid.PairGrid at 0x188964e62b0>

You may notice the plot\_kws argument. This enables us to pass down configuration options to the individual plotting calls on the off-diagonal elements. Check out the seaborn.pairplot docstring for more granular configuration options

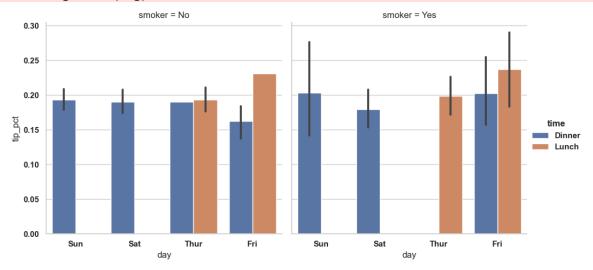
# **Facet Grids and Categorical Data**

What about datasets where we have additional grouping dimensions? One way to visualize data with many categorical variables is to use a facet grid. Seaborn has a useful built-in function factorplot that simplifies making many kinds of faceted plots (see Figure 9-26 for the resulting plot):

```
In [91]: sns.factorplot(x='day', y='tip_pct', hue='time', col='smoker',kind='bar', data=tips
```

C:\Users\ankit19.gupta\OneDrive - Reliance Corporate IT Park Limited\Desktop\Practic e\_Code\Python\_Practice\Python\_For\_Data\_Analysis\myenv\lib\site-packages\seaborn\cate gorical.py:3717: UserWarning: The `factorplot` function has been renamed to `catplot`. The original name will be removed in a future release. Please update your code. N ote that the default `kind` in `factorplot` (`'point'`) has changed `'strip'` in `catplot`.

warnings.warn(msg)

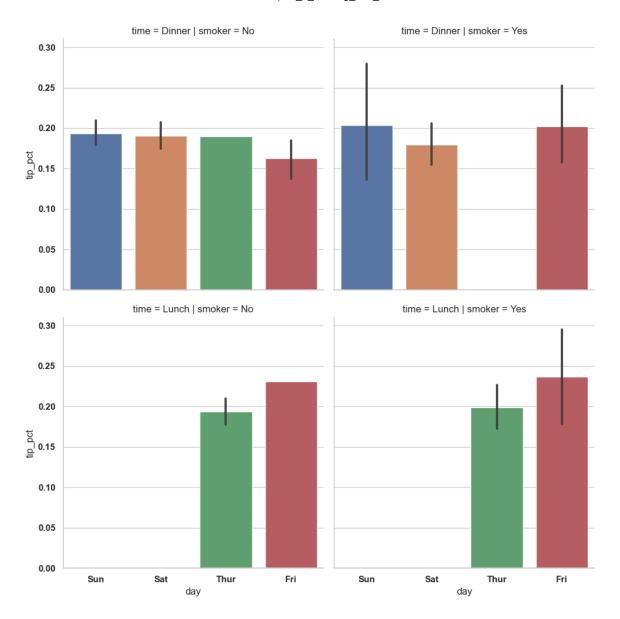


Out[91]: <seaborn.axisgrid.FacetGrid at 0x1888a37bef0>

Instead of grouping by 'time' by different bar colors within a facet, we can also expand the facet grid by adding one row per time value (Figure 9-27):

C:\Users\ankit19.gupta\OneDrive - Reliance Corporate IT Park Limited\Desktop\Practic e\_Code\Python\_Practice\Python\_For\_Data\_Analysis\myenv\lib\site-packages\seaborn\cate gorical.py:3717: UserWarning: The `factorplot` function has been renamed to `catplot`. The original name will be removed in a future release. Please update your code. N ote that the default `kind` in `factorplot` (`'point'`) has changed `'strip'` in `catplot`.

warnings.warn(msg)

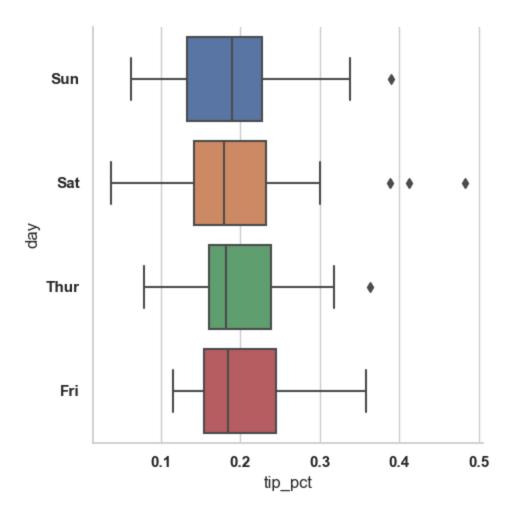


Out[92]: <seaborn.axisgrid.FacetGrid at 0x18889da5400>

factorplot supports other plot types that may be useful depending on what you are trying to display. For example, box plots (which show the median, quartiles, and outliers) can be an effective visualization type (Figure 9-28):

C:\Users\ankit19.gupta\OneDrive - Reliance Corporate IT Park Limited\Desktop\Practic e\_Code\Python\_Practice\Python\_For\_Data\_Analysis\myenv\lib\site-packages\seaborn\cate gorical.py:3717: UserWarning: The `factorplot` function has been renamed to `catplot`. The original name will be removed in a future release. Please update your code. N ote that the default `kind` in `factorplot` (`'point'`) has changed `'strip'` in `catplot`.

warnings.warn(msg)



Out[93]: <seaborn.axisgrid.FacetGrid at 0x18884cbb5f8>

You can create your own facet grid plots using the more general seaborn. Facet Grid class. See the seaborn documentation for more.

### 9.3 Other Python Visualization Tools

As is common with open source, there are a plethora of options for creating graphics in Python (too many to list). Since 2010, much development effort has been focused on creating interactive graphics for publication on the web. With tools like Bokeh and Plotly, it's now possible to specify dynamic, interactive graphics in Python that are destined for a web browser.

For creating static graphics for print or web, I recommend defaulting to matplotlib and addon libraries like pandas and seaborn for your needs. For other data visualiza- tion requirements, it may be useful to learn one of the other available tools out there. I encourage you to explore the ecosystem as it continues to involve and innovate into the future.

In [ ]: