

## Department of CSE Data Science, ATME College of Engineering

<b>Subject with Code</b>	<b>Exploratory Data Analysis BDS613B</b>
<b>Sem</b>	<b>VI</b>

### Module 1

**Introduction to Python and NumPy:** Getting Started in IPython and Jupyter, Enhanced Interactive Features, The Basics of NumPy Arrays, Sorted Arrays, Structured Data: NumPy's Structured Arrays

#### Getting Started in IPython and Jupyter

There are many options for development environments for Python, and I'm often asked which one I use in my own work. My answer sometimes surprises people: my preferred environment is IPython plus a text editor (in my case, Emacs or Atom depending on my mood). IPython (short for *Interactive Python*) was started in 2001 by Fernando Perez as an enhanced Python interpreter, and has since grown into a project aiming to provide, in Perez's words, "Tools for the entire life cycle of research computing." If Python is the engine of our data science task, you might think of IPython as the interactive control panel.

As well as being a useful interactive interface to Python, IPython also provides a number of useful syntactic additions to the language; we'll cover the most useful of these additions here. In addition, IPython is closely tied with the Jupyter project, which provides a browser-based notebook that is useful for development, collaboration, sharing, and even publication of data science results. The IPython notebook is actually a special case of the broader Jupyter notebook structure, which encompasses notebooks for Julia, R, and other programming languages. As an example of the usefulness of the notebook format, look no further than the page you are reading: the entire manuscript for this book was composed as a set of IPython notebooks.

IPython is about using Python effectively for interactive scientific and data-intensive computing. This chapter will start by stepping through some of the IPython features that are useful to the practice of data science, focusing especially on the syntax it offers beyond the standard features of Python. Next, we will go into a bit more depth on some of the more useful "magic commands" that can speed-up common tasks in creating and using data science code. Finally, we will touch on some of the features of the notebook that make it useful in understanding data and sharing results.

# Shell or Notebook?

There are two primary means of using IPython that we'll discuss in this chapter: the IPython shell and the IPython notebook. The bulk of the material in this chapter is relevant to both, and the examples will switch between them depending on what is most convenient. In the few sections that are relevant to just one or the other, we will explicitly state that fact. Before we start, some words on how to launch the IPython shell and IPython notebook.

## Launching the IPython Shell

This chapter, like most of this book, is not designed to be absorbed passively. I recommend that as you read through it, you follow along and experiment with the tools and syntax we cover: the muscle-memory you build through doing this will be far more useful than the simple act of reading about it. Start by launching the IPython interpreter by typing `ipython` on the command-line; alternatively, if you've installed a distribution like Anaconda or EPD, there may be a launcher specific to your system.

Once you do this, you should see a prompt like the following:

```
IPython 4.0.1 -- An enhanced Interactive Python.

?      -> Introduction and overview of IPython's features.

%quickref -> Quick reference.

help    -> Python's own help system.

object? -> Details about 'object', use 'object??' for extra details.

In [1]:
```

With that, you're ready to follow along.

## Launching the Jupyter Notebook

The Jupyter notebook is a browser-based graphical interface to the IPython shell, and builds on it a rich set of dynamic display capabilities. As well as executing Python/IPython statements, the notebook allows the user to include formatted text, static and dynamic visualizations, mathematical

equations, JavaScript widgets, and much more. Furthermore, these documents can be saved in a way that lets other people open them and execute the code on their own systems.

Though the IPython notebook is viewed and edited through your web browser window, it must connect to a running Python process in order to execute code. This process (known as a "kernel") can be started by running the following command in your system shell:

```
$ jupyter notebook
```

This command will launch a local web server that will be visible to your browser. It immediately spits out a log showing what it is doing; that log will look something like this:

```
$ jupyter notebook
[NotebookApp] Serving notebooks from local directory: /Users/jakevdp/PythonDataScienceHandbook
[NotebookApp] 0 active kernels
[NotebookApp] The IPython Notebook is running at: http://localhost:8888/
[NotebookApp] Use Control-C to stop this server and shut down all kernels (twice to skip confirmation).
```

Upon issuing the command, your default browser should automatically open and navigate to the listed local URL; the exact address will depend on your system. If the browser does not open automatically, you can open a window and manually open this address (<http://localhost:8888/> in this example).

If you read no other section in this chapter, read this one: I find the tools discussed here to be the most transformative contributions of IPython to my daily workflow.

When a technologically-minded person is asked to help a friend, family member, or colleague with a computer problem, most of the time it's less a matter of knowing the answer as much as knowing how to quickly find an unknown answer. In data science it's the same: searchable web resources such as online documentation, mailing-list threads, and StackOverflow answers contain a wealth of information, even (especially?) if it is a topic you've found yourself searching before. Being an effective practitioner of data science is less about memorizing the tool or command you should

use for every possible situation, and more about learning to effectively find the information you don't know, whether through a web search engine or another means.

One of the most useful functions of IPython/Jupyter is to shorten the gap between the user and the type of documentation and search that will help them do their work effectively. While web searches still play a role in answering complicated questions, an amazing amount of information can be found through IPython alone. Some examples of the questions IPython can help answer in a few keystrokes:

- How do I call this function? What arguments and options does it have?
- What does the source code of this Python object look like?
- What is in this package I imported? What attributes or methods does this object have?

Here we'll discuss IPython's tools to quickly access this information, namely the `?` character to explore documentation, the `??` characters to explore source code, and the Tab key for auto-completion.

## Accessing Documentation with `?`

The Python language and its data science ecosystem is built with the user in mind, and one big part of that is access to documentation. Every Python object contains the reference to a string, known as a *doc string*, which in most cases will contain a concise summary of the object and how to use it. Python has a built-in `help()` function that can access this information and prints the results. For example, to see the documentation of the built-in `len` function, you can do the following:

```
In [1]: help(len)
```

```
Help on built-in function len in module builtins:
```

```
len(...)
```

```
len(object) -> integer
```

```
Return the number of items of a sequence or mapping.
```

Depending on your interpreter, this information may be displayed as inline text, or in some separate pop-up window.

Because finding help on an object is so common and useful, IPython introduces the `?` character as a shorthand for accessing this documentation and other relevant information:

```
In [2]: len?

Type:      builtin_function_or_method

String form: <built-in function len>

Namespace: Python builtin

Docstring:

len(object) -> integer

Return the number of items of a sequence or mapping.
```

This notation works for just about anything, including object methods:

```
In [3]: L = [1, 2, 3]

In [4]: L.insert?

Type:      builtin_function_or_method

String form: <built-in method insert of list object at 0x1024b8ea8>

Docstring: L.insert(index, object) -- insert object before index
```

or even objects themselves, with the documentation from their type:

```
In [5]: L?

Type:      list

String form: [1, 2, 3]

Length:    3

Docstring:

list() -> new empty list
```

```
list(iterable) -> new list initialized from iterable's items
```

Importantly, this will even work for functions or other objects you create yourself! Here we'll define a small function with a docstring:

```
In [6]: def square(a):  
.....:     """Return the square of a."""  
.....:     return a ** 2  
.....:
```

Note that to create a docstring for our function, we simply placed a string literal in the first line. Because doc strings are usually multiple lines, by convention we used Python's triple-quote notation for multi-line strings.

Now we'll use the `?` mark to find this doc string:

```
In [7]: square?  
  
Type:      function  
  
String form: <function square at 0x103713cb0>  
  
Definition: square(a)  
  
Docstring: Return the square of a.
```

This quick access to documentation via docstrings is one reason you should get in the habit of always adding such inline documentation to the code you write!

## Accessing Source Code with `??`

Because the Python language is so easily readable, another level of insight can usually be gained by reading the source code of the object you're curious about. IPython provides a shortcut to the source code with the double question mark (`??`):

```
In [8]: square??  
  
Type:      function
```

String form: <function square at 0x103713cb0>

Definition: square(a)

Source:

```
def square(a):  
    "Return the square of a"  
    return a ** 2
```

For simple functions like this, the double question-mark can give quick insight into the under-the-hood details.

If you play with this much, you'll notice that sometimes the `??` suffix doesn't display any source code: this is generally because the object in question is not implemented in Python, but in C or some other compiled extension language. If this is the case, the `??` suffix gives the same output as the `?` suffix. You'll find this particularly with many of Python's built-in objects and types, for example `len` from above:

In [9]: len??

Type: builtin\_function\_or\_method

String form: <built-in function len>

Namespace: Python builtin

Docstring:

len(object) -> integer

Return the number of items of a sequence or mapping.

Using `?` and/or `??` gives a powerful and quick interface for finding information about what any Python function or module does.

## Exploring Modules with Tab-Completion

IPython's other useful interface is the use of the tab key for auto-completion and exploration of the contents of objects, modules, and name-spaces. In the examples that follow, we'll use `<TAB>` to indicate when the Tab key should be pressed.

## Tab-completion of object contents

Every Python object has various attributes and methods associated with it. Like with the `help` function discussed before, Python has a built-in `dir` function that returns a list of these, but the tab-completion interface is much easier to use in practice. To see a list of all available attributes of an object, you can type the name of the object followed by a period (".") character and the Tab key:

```
In [10]: L.<TAB>

L.append L.copy L.extend L.insert L.remove L.sort

L.clear L.count L.index L.pop L.reverse
```

To narrow-down the list, you can type the first character or several characters of the name, and the Tab key will find the matching attributes and methods:

```
In [10]: L.c<TAB>

L.clear L.copy L.count


In [10]: L.co<TAB>

L.copy L.count
```

If there is only a single option, pressing the Tab key will complete the line for you. For example, the following will instantly be replaced with `L.count`:

```
In [10]: L.cou<TAB>
```

Though Python has no strictly-enforced distinction between public/external attributes and private/internal attributes, by convention a preceding underscore is used to denote such methods.



For clarity, these private methods and special methods are omitted from the list by default, but it's possible to list them by explicitly typing the underscore:

```
In [10]: L._<TAB>
L.__add__      L.__gt__      L.__reduce__
L.__class__    L.__hash__    L.__reduce_ex__
```

For brevity, we've only shown the first couple lines of the output. Most of these are Python's special double-underscore methods (often nicknamed "dunder" methods).

## Tab completion when importing

Tab completion is also useful when importing objects from packages. Here we'll use it to find all possible imports in the `itertools` package that start with `co`:

```
In [10]: from itertools import co<TAB>
combinations      compress
combinations_with_replacement count
```

Similarly, you can use tab-completion to see which imports are available on your system (this will change depending on which third-party scripts and modules are visible to your Python session):

```
In [10]: import <TAB>
Display all 399 possibilities? (y or n)
Crypto      dis      py_compile
Cython      distutils pycldr
...         ...      ...
difflib     pwd      zmq

In [10]: import h<TAB>
hashlib     hmac     http
```

```
heapq      html      husl
```

(Note that for brevity, I did not print here all 399 importable packages and modules on my system.)

## Beyond tab completion: wildcard matching

Tab completion is useful if you know the first few characters of the object or attribute you're looking for, but is little help if you'd like to match characters at the middle or end of the word. For this use-case, IPython provides a means of wildcard matching for names using the `*` character.

For example, we can use this to list every object in the namespace that ends with `Warning`:

```
In [10]: *Warning?

BytesWarning      RuntimeWarning
DeprecationWarning  SyntaxWarning
FutureWarning     UnicodeWarning
ImportWarning     UserWarning
PendingDeprecationWarning  Warning
ResourceWarning
```

Notice that the `*` character matches any string, including the empty string.

Similarly, suppose we are looking for a string method that contains the word `find` somewhere in its name. We can search for it this way:

```
In [10]: str.*find*?

str.find
str.rfind
```

I find this type of flexible wildcard search can be very useful for finding a particular command when getting to know a new package or reacquainting myself with a familiar one.

## Keyboard Shortcuts in the IPython Shell

If you spend any amount of time on the computer, you've probably found a use for keyboard shortcuts in your workflow. Most familiar perhaps are the Cmd-C and Cmd-V (or Ctrl-C and Ctrl-V) for copying and pasting in a wide variety of programs and systems. Power-users tend to go even further: popular text editors like Emacs, Vim, and others provide users an incredible range of operations through intricate combinations of keystrokes.

The IPython shell doesn't go this far, but does provide a number of keyboard shortcuts for fast navigation while typing commands. These shortcuts are not in fact provided by IPython itself, but through its dependency on the GNU Readline library: as such, some of the following shortcuts may differ depending on your system configuration. Also, while some of these shortcuts do work in the browser-based notebook, this section is primarily about shortcuts in the IPython shell.

Once you get accustomed to these, they can be very useful for quickly performing certain commands without moving your hands from the "home" keyboard position. If you're an Emacs user or if you have experience with Linux-style shells, the following will be very familiar. We'll group these shortcuts into a few categories: *navigation shortcuts*, *text entry shortcuts*, *command history shortcuts*, and *miscellaneous shortcuts*.

### Navigation shortcuts

While the use of the left and right arrow keys to move backward and forward in the line is quite obvious, there are other options that don't require moving your hands from the "home" keyboard position:

Keystroke	Action
Ctrl-a	Move cursor to the beginning of the line

Keystroke	Action
Ctrl-e	Move cursor to the end of the line
Ctrl-b or the left arrow key	Move cursor back one character
Ctrl-f or the right arrow key	Move cursor forward one character

## Text Entry Shortcuts

While everyone is familiar with using the Backspace key to delete the previous character, reaching for the key often requires some minor finger gymnastics, and it only deletes a single character at a time. In IPython there are several shortcuts for removing some portion of the text you're typing. The most immediately useful of these are the commands to delete entire lines of text. You'll know these have become second-nature if you find yourself using a combination of Ctrl-b and Ctrl-d instead of reaching for Backspace to delete the previous character!

Keystroke	Action
Backspace key	Delete previous character in line
Ctrl-d	Delete next character in line
Ctrl-k	Cut text from cursor to end of line
Ctrl-u	Cut text from beginning of line to cursor
Ctrl-y	Yank (i.e. paste) text that was previously cut
Ctrl-t	Transpose (i.e., switch) previous two characters

## Command History Shortcuts

Perhaps the most impactful shortcuts discussed here are the ones IPython provides for navigating the command history. This command history goes beyond your current IPython session: your entire command history is stored in a SQLite database in your IPython profile directory. The most straightforward way to access these is with the up and down arrow keys to step through the history, but other options exist as well:

Keystroke	Action
Ctrl-p (or the up arrow key)	Access previous command in history
Ctrl-n (or the down arrow key)	Access next command in history
Ctrl-r	Reverse-search through command history

The reverse-search can be particularly useful. Recall that in the previous section we defined a function called `square`. Let's reverse-search our Python history from a new IPython shell and find this definition again. When you press Ctrl-r in the IPython terminal, you'll see the following prompt:

```
In [1]:  
(reverse-i-search)␣:
```

If you start typing characters at this prompt, IPython will auto-fill the most recent command, if any, that matches those characters:

```
In [1]:  
(reverse-i-search)␣sqa': square??
```

At any point, you can add more characters to refine the search, or press Ctrl-r again to search further for another command that matches the query. If you followed along in the previous section, pressing Ctrl-r twice more gives:

```
In [1]:  
(reverse-i-search)␣sqa': def square(a):
```

```
"""Return the square of a"""
```

```
return a ** 2
```

Once you have found the command you're looking for, press Return and the search will end. We can then use the retrieved command, and carry-on with our session:

```
In [1]: def square(a):
```

```
"""Return the square of a"""
```

```
return a ** 2
```

```
In [2]: square(2)
```

```
Out[2]: 4
```

Note that Ctrl-p/Ctrl-n or the up/down arrow keys can also be used to search through history, but only by matching characters at the beginning of the line. That is, if you type `def` and then press Ctrl-p, it would find the most recent command (if any) in your history that begins with the characters `def`.

## Miscellaneous Shortcuts

Finally, there are a few miscellaneous shortcuts that don't fit into any of the preceding categories, but are nevertheless useful to know:

### KeystrokeAction

Ctrl-l      Clear terminal screen

Ctrl-c      Interrupt current Python command

Ctrl-d      Exit IPython session

The Ctrl-c in particular can be useful when you inadvertently start a very long-running job.

While some of the shortcuts discussed here may seem a bit tedious at first, they quickly become automatic with practice. Once you develop that muscle memory,

## Python Magic Commands

The previous two sections showed how IPython lets you use and explore Python efficiently and interactively. Here we'll begin discussing some of the enhancements that IPython adds on top of the normal Python syntax. These are known in IPython as *magic commands*, and are prefixed by the `%` character. These magic commands are designed to succinctly solve various common problems in standard data analysis. Magic commands come in two flavors: *line magics*, which are denoted by a single `%` prefix and operate on a single line of input, and *cell magics*, which are denoted by a double `%%` prefix and operate on multiple lines of input. We'll demonstrate and discuss a few brief examples here, and come back to more focused discussion of several useful magic commands later in the chapter.

### Pasting Code Blocks: `%paste` and `%cpaste`

When working in the IPython interpreter, one common gotcha is that pasting multi-line code blocks can lead to unexpected errors, especially when indentation and interpreter markers are involved. A common case is that you find some example code on a website and want to paste it into your interpreter. Consider the following simple function:

```
>>> def donothing(x):  
...     return x
```

The code is formatted as it would appear in the Python interpreter, and if you copy and paste this directly into IPython you get an error:

```
In [2]: >>> def donothing(x):  
...:     ...     return x  
...:
```

```
File "<ipython-input-20-5a66c8964687>", line 2
```

```
...     return x
```

```
      ^
```

```
SyntaxError: invalid syntax
```

In the direct paste, the interpreter is confused by the additional prompt characters. But never fear—IPython's `%paste` magic function is designed to handle this exact type of multi-line, marked-up input:

```
In [3]: %paste
```

```
>>> def donothing(x):
```

```
...     return x
```

```
## -- End pasted text --
```

The `%paste` command both enters and executes the code, so now the function is ready to be used:

```
In [4]: donothing(10)
```

```
Out[4]: 10
```

A command with a similar intent is `%cpaste`, which opens up an interactive multiline prompt in which you can paste one or more chunks of code to be executed in a batch:

```
In [5]: %cpaste
```

```
Pasting code; enter '--' alone on the line to stop or use Ctrl-D.
```

```
:>>> def donothing(x):
```

```
:...     return x
```

```
:--
```

These magic commands, like others we'll see, make available functionality that would be difficult or impossible in a standard Python interpreter.



# Running External Code: `%run`

As you begin developing more extensive code, you will likely find yourself working in both IPython for interactive exploration, as well as a text editor to store code that you want to reuse. Rather than running this code in a new window, it can be convenient to run it within your IPython session. This can be done with the `%run` magic.

For example, imagine you've created a `myscript.py` file with the following contents:

```
#-----  
  
# file: myscript.py  
  
def square(x):  
    """square a number"""  
    return x ** 2  
  
for N in range(1, 4):  
    print(N, "squared is", square(N))
```

You can execute this from your IPython session as follows:

```
In [6]: %run myscript.py  
  
1 squared is 1  
2 squared is 4  
3 squared is 9
```

Note also that after you've run this script, any functions defined within it are available for use in your IPython session:

```
In [7]: square(5)  
  
Out[7]: 25
```

There are several options to fine-tune how your code is run; you can see the documentation in the normal way, by typing `%run?` in the IPython interpreter.

## Timing Code Execution: `%timeit`

Another example of a useful magic function is `%timeit`, which will automatically determine the execution time of the single-line Python statement that follows it. For example, we may want to check the performance of a list comprehension:

```
In [8]: %timeit L = [n ** 2 for n in range(1000)]
```

```
1000 loops, best of 3: 325 µs per loop
```

The benefit of `%timeit` is that for short commands it will automatically perform multiple runs in order to attain more robust results. For multi line statements, adding a second `%` sign will turn this into a cell magic that can handle multiple lines of input. For example, here's the equivalent construction with a `for`-loop:

```
In [9]: %%timeit
```

```
...: L = []
```

```
...: for n in range(1000):
```

```
...:     L.append(n ** 2)
```

```
...:
```

```
1000 loops, best of 3: 373 µs per loop
```

We can immediately see that list comprehensions are about 10% faster than the equivalent `for`-loop construction in this case.

## Help on Magic Functions: `?`, `%magic`, and `%lsmagic`

Like normal Python functions, IPython magic functions have docstrings, and this useful documentation can be accessed in the standard manner. So, for example, to read the documentation of the `%timeit` magic simply type this:

```
In [10]: %timeit?
```

Documentation for other functions can be accessed similarly. To access a general description of available magic functions, including some examples, you can type this:

```
In [11]: %magic
```

For a quick and simple list of all available magic functions, type this:

```
In [12]: %lsmagic
```

Finally, I'll mention that it is quite straightforward to define your own magic functions if you wish.

## Input and Output History

Previously we saw that the IPython shell allows you to access previous commands with the up and down arrow keys, or equivalently the Ctrl-p/Ctrl-n shortcuts. Additionally, in both the shell and the notebook, IPython exposes several ways to obtain the output of previous commands, as well as string versions of the commands themselves. We'll explore those here.

## IPython's `In` and `Out` Objects

By now I imagine you're quite familiar with the `In [1]:/Out[1]:` style prompts used by IPython. But it turns out that these are not just pretty decoration: they give a clue as to how you can access previous inputs and outputs in your current session. Imagine you start a session that looks like this:

```
In [1]: import math
```

```
In [2]: math.sin(2)
```

```
Out[2]: 0.9092974268256817
```

```
In [3]: math.cos(2)
```

```
Out[3]: -0.4161468365471424
```

We've imported the built-in `math` package, then computed the sine and the cosine of the number 2. These inputs and outputs are displayed in the shell with `In/Out` labels, but there's more—IPython actually creates some Python variables called `In` and `Out` that are automatically updated to reflect this history:

```
In [4]: print(In)

['', 'import math', 'math.sin(2)', 'math.cos(2)', 'print(In)']

In [5]: Out

Out[5]: {2: 0.9092974268256817, 3: -0.4161468365471424}
```

The `In` object is a list, which keeps track of the commands in order (the first item in the list is a place-holder so that `In[1]` can refer to the first command):

```
In [6]: print(In[1])

import math
```

The `Out` object is not a list but a dictionary mapping input numbers to their outputs (if any):

```
In [7]: print(Out[2])

0.9092974268256817
```

Note that not all operations have outputs: for example, `import` statements and `print` statements don't affect the output. The latter may be surprising, but makes sense if you consider that `print` is a function that returns `None`; for brevity, any command that returns `None` is not added to `Out`.

Where this can be useful is if you want to interact with past results. For example, let's check the sum of `sin(2) ** 2` and `cos(2) ** 2` using the previously-computed results:

```
In [8]: Out[2] ** 2 + Out[3] ** 2

Out[8]: 1.0
```

The result is `1.0` as we'd expect from the well-known trigonometric identity. In this case, using these previous results probably is not necessary, but it can become very handy if you execute a very expensive computation and want to reuse the result!

## Underscore Shortcuts and Previous Outputs

The standard Python shell contains just one simple shortcut for accessing previous output; the variable `_` (i.e., a single underscore) is kept updated with the previous output; this works in IPython as well:

```
In [9]: print(_)
```

```
1.0
```

But IPython takes this a bit further—you can use a double underscore to access the second-to-last output, and a triple underscore to access the third-to-last output (skipping any commands with no output):

```
In [10]: print(__)
```

```
-0.4161468365471424
```

```
In [11]: print(___)
```

```
0.9092974268256817
```

IPython stops there: more than three underscores starts to get a bit hard to count, and at that point it's easier to refer to the output by line number.

There is one more shortcut we should mention, however—a shorthand for `Out[X]` is `_X` (i.e., a single underscore followed by the line number):

```
In [12]: Out[2]
```

```
Out[12]: 0.9092974268256817
```

```
In [13]: _2
```

```
Out[13]: 0.9092974268256817
```

## Suppressing Output

Sometimes you might wish to suppress the output of a statement (this is perhaps most common with the plotting commands that we'll explore in [Introduction to Matplotlib](#)). Or maybe the command you're executing produces a result that you'd prefer not like to store in your output history, perhaps so that it can be deallocated when other references are removed. The easiest way to suppress the output of a command is to add a semicolon to the end of the line:

```
In [14]: math.sin(2) + math.cos(2);
```

Note that the result is computed silently, and the output is neither displayed on the screen or stored in the `Out` dictionary:

```
In [15]: 14 in Out
```

```
Out[15]: False
```

## Related Magic Commands

For accessing a batch of previous inputs at once, the `%history` magic command is very helpful. Here is how you can print the first four inputs:

```
In [16]: %history -n 1-4
```

```
1: import math
```

```
2: math.sin(2)
```

```
3: math.cos(2)
```

```
4: print(In)
```

As usual, you can type `%history?` for more information and a description of options available. Other similar magic commands are `%rerun` (which will re-execute some portion of the command history) and `%save` (which saves some set of the command history to a file).

## IPython and Shell Commands

When working interactively with the standard Python interpreter, one of the frustrations is the need to switch between multiple windows to access Python tools and system command-line tools. IPython bridges this gap, and gives you a syntax for executing shell commands directly from within the IPython terminal. The magic happens with the exclamation point: anything appearing after `!` on a line will be executed not by the Python kernel, but by the system command-line.

The following assumes you're on a Unix-like system, such as Linux or Mac OSX. Some of the examples that follow will fail on Windows, which uses a different type of shell by default (though with the 2016 announcement of native Bash shells on Windows, soon this may no longer be an issue!).

## Quick Introduction to the Shell

A full intro to using the shell/terminal/command-line is well beyond the scope of this chapter, but for the uninitiated we will offer a quick introduction here. The shell is a way to interact textually with your computer. Ever since the mid 1980s, when Microsoft and Apple introduced the first versions of their now ubiquitous graphical operating systems, most computer users have interacted with their operating system through familiar clicking of menus and drag-and-drop movements. But operating systems existed long before these graphical user interfaces, and were primarily controlled through sequences of text input: at the prompt, the user would type a command, and the computer would do what the user told it to. Those early prompt systems are the precursors of the shells and terminals that most active data scientists still use today.

Someone unfamiliar with the shell might ask why you would bother with this, when many results can be accomplished by simply clicking on icons and menus. A shell user might reply with another question: why hunt icons and click menus when you can accomplish things much more easily by typing? While it might sound like a typical tech preference impasse, when moving beyond basic

tasks it quickly becomes clear that the shell offers much more control of advanced tasks, though admittedly the learning curve can intimidate the average computer user.

As an example, here is a sample of a Linux/OSX shell session where a user explores, creates, and modifies directories and files on their system (osx:~ \$ is the prompt, and everything after the \$ sign is the typed command; text that is preceded by a # is meant just as description, rather than something you would actually type in):

```
osx:~ $ echo "hello world"      # echo is like Python's print function
```

```
hello world
```

```
osx:~ $ pwd                    # pwd = print working directory
```

```
/home/jake                    # this is the "path" that we're sitting in
```

```
osx:~ $ ls                     # ls = list working directory contents
```

```
notebooks  projects
```

```
osx:~ $ cd projects/          # cd = change directory
```

```
osx:projects $ pwd
```

```
/home/jake/projects
```

```
osx:projects $ ls
```

```
datasci_book  mpld3  myproject.txt
```

```
osx:projects $ mkdir myproject  # mkdir = make new directory
```

```
osx:projects $ cd myproject/
```



```
osx:myproject $ mv ../myproject.txt ./ # mv = move file. Here we're moving the
```

```
# file myproject.txt from one directory
```

```
# up (../) to the current directory (./)
```

```
osx:myproject $ ls
```

```
myproject.txt
```

Notice that all of this is just a compact way to do familiar operations (navigating a directory structure, creating a directory, moving a file, etc.) by typing commands rather than clicking icons and menus. Note that with just a few commands (`pwd`, `ls`, `cd`, `mkdir`, and `cp`) you can do many of the most common file operations. It's when you go beyond these basics that the shell approach becomes really powerful.

## Shell Commands in IPython

Any command that works at the command-line can be used in IPython by prefixing it with the `!` character. For example, the `ls`, `pwd`, and `echo` commands can be run as follows:

```
In [1]: !ls
```

```
myproject.txt
```

```
In [2]: !pwd
```

```
/home/jake/projects/myproject
```

```
In [3]: !echo "printing from the shell"
```

```
printing from the shell
```

## Passing Values to and from the Shell

Shell commands can not only be called from IPython, but can also be made to interact with the IPython namespace. For example, you can save the output of any shell command to a Python list using the assignment operator:

```
In [4]: contents = !ls
```

```
In [5]: print(contents)
```

```
['myproject.txt']
```

```
In [6]: directory = !pwd
```

```
In [7]: print(directory)
```

```
['/Users/jakevdp/notebooks/tmp/myproject']
```

Note that these results are not returned as lists, but as a special shell return type defined in IPython:

```
In [8]: type(directory)
```

```
IPython.utils.text.SList
```

This looks and acts a lot like a Python list, but has additional functionality, such as the `grep` and `fields` methods and the `s`, `n`, and `p` properties that allow you to search, filter, and display the results in convenient ways. For more information on these, you can use IPython's built-in help features.

Communication in the other direction—passing Python variables into the shell—is possible using the `{varname}` syntax:

```
In [9]: message = "hello from Python"
```

```
In [10]: !echo {message}
```

```
hello from Python
```

The curly braces contain the variable name, which is replaced by the variable's contents in the shell command.

# Shell-Related Magic Commands

If you play with IPython's shell commands for a while, you might notice that you cannot use `!cd` to navigate the filesystem:

```
In [11]: !pwd
/home/jake/projects/myproject

In [12]: !cd ..

In [13]: !pwd
/home/jake/projects/myproject
```

The reason is that shell commands in the notebook are executed in a temporary subshell. If you'd like to change the working directory in a more enduring way, you can use the `%cd` magic command:

```
In [14]: %cd ..
/home/jake/projects
```

In fact, by default you can even use this without the `%` sign:

```
In [15]: cd myproject
/home/jake/projects/myproject
```

This is known as an `automagic` function, and this behavior can be toggled with the `%automagic` magic function.

Besides `%cd`, other available shell-like magic functions are `%cat`, `%cp`, `%env`, `%ls`, `%man`, `%mkdir`, `%more`, `%mv`, `%pwd`, `%rm`, and `%rmdir`, any of which can be used without the `%` sign if `automagic` is on. This makes it so that you can almost treat the IPython prompt as if it's a normal shell:

```
In [16]: mkdir tmp
```

```
In [17]: ls
```

```
myproject.txt tmp/
```

```
In [18]: cp myproject.txt tmp/
```

```
In [19]: ls tmp
```

```
myproject.txt
```

```
In [20]: rm -r tmp
```

This access to the shell from within the same terminal window as your Python session means that there is a lot less switching back and forth between interpreter and shell as you write your Python code.

## Errors and Debugging

Code development and data analysis always require a bit of trial and error, and IPython contains tools to streamline this process. This section will briefly cover some options for controlling Python's exception reporting, followed by exploring tools for debugging errors in code.

### Controlling Exceptions: `%xmode`

Most of the time when a Python script fails, it will raise an Exception. When the interpreter hits one of these exceptions, information about the cause of the error can be found in the *traceback*, which can be accessed from within Python. With the `%xmode` magic function, IPython allows you to control the amount of information printed when the exception is raised. Consider the following code:

```
In [1]:
```

```
def func1(a, b):
```

```

    return a / b

def func2(x):
    a = x
    b = x - 1
    return func1(a, b)
In [2]:
func2(1)
-----
ZeroDivisionError                                Traceback (most recent call last)
<ipython-input-2-b2e110f6fc8f> in <module>()
----> 1 func2(1)

<ipython-input-1-d849e34d61fb> in func2(x)
      5 a = x
      6 b = x - 1
----> 7 return func1(a, b)

<ipython-input-1-d849e34d61fb> in func1(a, b)
      1 def func1(a, b):
----> 2 return a / b
      3
      4 def func2(x):
      5 a = x

```

**ZeroDivisionError**: division by zero

Calling `func2` results in an error, and reading the printed trace lets us see exactly what happened. By default, this trace includes several lines showing the context of each step that led to the error. Using the `%xmode` magic function (short for *Exception mode*), we can change what information is printed.

`%xmode` takes a single argument, the mode, and there are three possibilities: `Plain`, `Context`, and `Verbose`. The default is `Context`, and gives output like that just shown before. `Plain` is more compact and gives less information:

```

In [3]:
%xmode Plain
Exception reporting mode: Plain
In [4]:
func2(1)
Traceback (most recent call last):

File "<ipython-input-4-b2e110f6fc8f>", line 1, in <module>
    func2(1)

File "<ipython-input-1-d849e34d61fb>", line 7, in func2
    return func1(a, b)

```

```
File "<ipython-input-1-d849e34d61fb>", line 2, in func1
    return a / b
```

**ZeroDivisionError:** division by zero

The `Verbose` mode adds some extra information, including the arguments to any functions that are called:

In [5]:

```
%xmode Verbose
Exception reporting mode: Verbose
```

In [6]:

```
func2(1)
```

**ZeroDivisionError** Traceback (most recent call last)

```
<ipython-input-6-b2e110f6fc8f> in <module>()
```

```
----> 1 func2(1)
```

```
      global func2 = <function func2 at 0x103729320>
```

```
<ipython-input-1-d849e34d61fb> in func2(x=1)
```

```
5     a = x
```

```
6     b = x - 1
```

```
----> 7     return func1(a, b)
```

```
      global func1 = <function func1 at 0x1037294d0>
```

```
      a = 1
```

```
      b = 0
```

```
<ipython-input-1-d849e34d61fb> in func1(a=1, b=0)
```

```
1 def func1(a, b):
```

```
----> 2     return a / b
```

```
      a = 1
```

```
      b = 0
```

```
3
```

```
4 def func2(x):
```

```
5     a = x
```

**ZeroDivisionError:** division by zero

This extra information can help narrow-in on why the exception is being raised. So why not use the `Verbose` mode all the time? As code gets complicated, this kind of traceback can get extremely long. Depending on the context, sometimes the brevity of `Default` mode is easier to work with.

## Debugging: When Reading Tracebacks Is Not Enough

The standard Python tool for interactive debugging is `pdb`, the Python debugger. This debugger lets the user step through the code line by line in order to see what might be causing a more difficult error. The IPython-enhanced version of this is `ipdb`, the IPython debugger.

There are many ways to launch and use both these debuggers; we won't cover them fully here. Refer to the online documentation of these two utilities to learn more.

In IPython, perhaps the most convenient interface to debugging is the `%debug` magic command. If you call it after hitting an exception, it will automatically open an interactive debugging prompt at the point of the exception. The `ipdb` prompt lets you explore the current state of the stack, explore the available variables, and even run Python commands!

Let's look at the most recent exception, then do some basic tasks—print the values of `a` and `b`, and type `quit` to quit the debugging session:

In [7]:

```
%debug
> <ipython-input-1-d849e34d61fb>(2)func1()
  1 def func1(a, b):
----> 2     return a / b
      3

ipdb> print(a)
1
ipdb> print(b)
0
ipdb> quit
```

The interactive debugger allows much more than this, though—we can even step up and down through the stack and explore the values of variables there:

In [8]:

```
%debug
> <ipython-input-1-d849e34d61fb>(2)func1()
  1 def func1(a, b):
----> 2     return a / b
      3

ipdb> up
> <ipython-input-1-d849e34d61fb>(7)func2()
  5 a = x
  6 b = x - 1
----> 7     return func1(a, b)

ipdb> print(x)
```

```
1
ipdb> up
> <ipython-input-6-b2e110f6fc8f>(1)<module>()
----> 1 func2(1)
```

```
ipdb> down
> <ipython-input-1-d849e34d61fb>(7)func2()
   5   a = x
   6   b = x - 1
----> 7   return func1(a, b)
```

```
ipdb> quit
```

This allows you to quickly find out not only what caused the error, but what function calls led up to the error.

If you'd like the debugger to launch automatically whenever an exception is raised, you can use the `%pdb` magic function to turn on this automatic behavior:

In [9]:

```
%xmode Plain
%pdb on
func2(1)
Exception reporting mode: Plain
Automatic pdb calling has been turned ON
Traceback (most recent call last):

File "<ipython-input-9-569a67d2d312>", line 3, in <module>
    func2(1)

File "<ipython-input-1-d849e34d61fb>", line 7, in func2
    return func1(a, b)

File "<ipython-input-1-d849e34d61fb>", line 2, in func1
    return a / b
```

```
ZeroDivisionError: division by zero
> <ipython-input-1-d849e34d61fb>(2)func1()
   1 def func1(a, b):
----> 2     return a / b
   3
```

```
ipdb> print(b)
0
ipdb> quit
```

Finally, if you have a script that you'd like to run from the beginning in interactive mode, you can run it with the command `%run -d`, and use the `next` command to step through the lines of code interactively.



## Partial list of debugging commands

There are many more available commands for interactive debugging than we've listed here; the following table contains a description of some of the more common and useful ones:

Command	Description
<code>list</code>	Show the current location in the file
<code>h(elp)</code>	Show a list of commands, or find help on a specific command
<code>q(uit)</code>	Quit the debugger and the program
<code>c(ontinue)</code>	Quit the debugger, continue in the program
<code>n(ext)</code>	Go to the next step of the program
<code>&lt;enter&gt;</code>	Repeat the previous command
<code>p(rint)</code>	Print variables
<code>s(tep)</code>	Step into a subroutine
<code>r(eturn)</code>	Return out of a subroutine

## Profiling and Timing Code

In the process of developing code and creating data processing pipelines, there are often trade-offs you can make between various implementations. Early in developing your algorithm, it can be counterproductive to worry about such things. As Donald Knuth famously quipped, "We should forget about small efficiencies, say about 97% of the time: premature optimization is the root of all evil."

But once you have your code working, it can be useful to dig into its efficiency a bit. Sometimes it's useful to check the execution time of a given command or set of commands; other times it's useful to dig into a multiline process and determine where the bottleneck lies in some complicated

series of operations. IPython provides access to a wide array of functionality for this kind of timing and profiling of code. Here we'll discuss the following IPython magic commands:

- `%time`: Time the execution of a single statement
- `%timeit`: Time repeated execution of a single statement for more accuracy
- `%prun`: Run code with the profiler
- `%lprun`: Run code with the line-by-line profiler
- `%memit`: Measure the memory use of a single statement
- `%mprun`: Run code with the line-by-line memory profiler

The last four commands are not bundled with IPython—you'll need to get the `line_profiler` and `memory_profiler` extensions, which we will discuss in the following sections.

## Timing Code Snippets: `%timeit` and `%time`

We saw the `%timeit` line-magic and `%%timeit` cell-magic in the introduction to magic functions in [IPython Magic Commands](#); it can be used to time the repeated execution of snippets of code:

In [1]:

```
%timeit sum(range(100))
100000 loops, best of 3: 1.54 µs per loop
```

Note that because this operation is so fast, `%timeit` automatically does a large number of repetitions. For slower commands, `%timeit` will automatically adjust and perform fewer repetitions:

In [2]:

```
%%timeit
total = 0
for i in range(1000):
    for j in range(1000):
        total += i * (-1) ** j
1 loops, best of 3: 407 ms per loop
```

Sometimes repeating an operation is not the best option. For example, if we have a list that we'd like to sort, we might be misled by a repeated operation. Sorting a pre-sorted list is much faster than sorting an unsorted list, so the repetition will skew the result:

In [3]:

```
import random
L = [random.random() for i in range(100000)]
%timeit L.sort()
```

100 loops, best of 3: 1.9 ms per loop

For this, the `%time` magic function may be a better choice. It also is a good choice for longer-running commands, when short, system-related delays are unlikely to affect the result. Let's time the sorting of an unsorted and a presorted list:

In [4]:

```
import random
L = [random.random() for i in range(100000)]
print("sorting an unsorted list:")
%time L.sort()
sorting an unsorted list:
CPU times: user 40.6 ms, sys: 896 µs, total: 41.5 ms
Wall time: 41.5 ms
```

In [5]:

```
print("sorting an already sorted list:")
%time L.sort()
sorting an already sorted list:
CPU times: user 8.18 ms, sys: 10 µs, total: 8.19 ms
Wall time: 8.24 ms
```

Notice how much faster the presorted list is to sort, but notice also how much longer the timing takes with `%time` versus `%timeit`, even for the presorted list! This is a result of the fact that `%timeit` does some clever things under the hood to prevent system calls from interfering with the timing. For example, it prevents cleanup of unused Python objects (known as *garbage collection*) which might otherwise affect the timing. For this reason, `%timeit` results are usually noticeably faster than `%time` results.

For `%time` as with `%timeit`, using the double-percent-sign cell magic syntax allows timing of multiline scripts:

In [6]:

```
%%time
total = 0
for i in range(1000):
    for j in range(1000):
        total += i * (-1) ** j
CPU times: user 504 ms, sys: 979 µs, total: 505 ms
Wall time: 505 ms
```

For more information on `%time` and `%timeit`, as well as their available options, use the IPython help functionality (i.e., type `%time?` at the IPython prompt).

## Profiling Full Scripts: `%prun`

A program is made of many single statements, and sometimes timing these statements in context is more important than timing them on their own. Python contains a built-in code profiler (which you can read about in the Python documentation), but IPython offers a much more convenient way to use this profiler, in the form of the magic function `%prun`.

By way of example, we'll define a simple function that does some calculations:

In [7]:

```
def sum_of_lists(N):
    total = 0
    for i in range(5):
        L = [j ^ (j >> i) for j in range(N)]
        total += sum(L)
    return total
```

Now we can call `%prun` with a function call to see the profiled results:

In [8]:

```
%prun sum_of_lists(1000000)
```

In the notebook, the output is printed to the pager, and looks something like this:

14 function calls in 0.714 seconds

Ordered by: internal time

ncalls	tottime	percall	cumtime	percall	filename:lineno(function)
5	0.599	0.120	0.599	0.120	<ipython-input-19>:4(<listcomp>)
5	0.064	0.013	0.064	0.013	{built-in method sum}
1	0.036	0.036	0.699	0.699	<ipython-input-19>:1(sum_of_lists)
1	0.014	0.014	0.714	0.714	<string>:1(<module>)
1	0.000	0.000	0.714	0.714	{built-in method exec}

The result is a table that indicates, in order of total time on each function call, where the execution is spending the most time. In this case, the bulk of execution time is in the list comprehension

inside `sum_of_lists`. From here, we could start thinking about what changes we might make to improve the performance in the algorithm.

For more information on `%prun`, as well as its available options, use the IPython help functionality (i.e., type `%prun?` at the IPython prompt).

## Line-By-Line Profiling with `%lprun`

The function-by-function profiling of `%prun` is useful, but sometimes it's more convenient to have a line-by-line profile report. This is not built into Python or IPython, but there is a `line_profiler` package available for installation that can do this. Start by using Python's packaging tool, `pip`, to install the `line_profiler` package:

```
$ pip install line_profiler
```

Next, you can use IPython to load the `line_profiler` IPython extension, offered as part of this package:

In [9]:

```
%load_ext line_profiler
```

Now the `%lprun` command will do a line-by-line profiling of any function—in this case, we need to tell it explicitly which functions we're interested in profiling:

In [10]:

```
%lprun -f sum_of_lists sum_of_lists(5000)
```

As before, the notebook sends the result to the pager, but it looks something like this:

```
Timer unit: 1e-06 s
```

```
Total time: 0.009382 s
```

```
File: <ipython-input-19-fa2be176cc3e>
```

```
Function: sum_of_lists at line 1
```

Line #	Hits	Time	Per Hit	% Time	Line Contents
--------	------	------	---------	--------	---------------

=====					
1	def sum_of_lists(N):				
2	1	2	2.0	0.0	total = 0
3	6	8	1.3	0.1	for i in range(5):
4	5	9001	1800.2	95.9	L = [j ^ (j >> i) for j in range(N)]
5	5	371	74.2	4.0	total += sum(L)
6	1	0	0.0	0.0	return total

The information at the top gives us the key to reading the results: the time is reported in microseconds and we can see where the program is spending the most time. At this point, we may be able to use this information to modify aspects of the script and make it perform better for our desired use case.

For more information on `%lprun`, as well as its available options, use the IPython help functionality (i.e., type `%lprun?` at the IPython prompt).

## Profiling Memory Use: `%memit` and `%mprun`

Another aspect of profiling is the amount of memory an operation uses. This can be evaluated with another IPython extension, the `memory_profiler`. As with the `line_profiler`, we start by pip-installing the extension:

```
$ pip install memory_profiler
```

Then we can use IPython to load the extension:

In [12]:

```
%load_ext memory_profiler
```

The memory profiler extension contains two useful magic functions: the `%memit` magic (which offers a memory-measuring equivalent of `%timeit`) and the `%mprun` function (which offers a memory-measuring equivalent of `%lprun`). The `%memit` function can be used rather simply:

In [13]:

```
%memit sum_of_lists(1000000)
peak memory: 100.08 MiB, increment: 61.36 MiB
```

We see that this function uses about 100 MB of memory.

For a line-by-line description of memory use, we can use the `%mprun` magic. Unfortunately, this magic works only for functions defined in separate modules rather than the notebook itself, so we'll start by using the `%%file` magic to create a simple module called `mprun_demo.py`, which contains our `sum_of_lists` function, with one addition that will make our memory profiling results more clear:

In [14]:

```
%%file mprun_demo.py
def sum_of_lists(N):
    total = 0
    for i in range(5):
        L = [j ^ (j >> i) for j in range(N)]
        total += sum(L)
        del L # remove reference to L
    return total
```

Overwriting `mprun_demo.py`

We can now import the new version of this function and run the memory line profiler:

In [15]:

```
from mprun_demo import sum_of_lists
%mprun -f sum_of_lists sum_of_lists(1000000)
```

The result, printed to the pager, gives us a summary of the memory use of the function, and looks something like this:

Filename: `./mprun_demo.py`

Line #	Mem usage	Increment	Line Contents
=====			
4	71.9 MiB	0.0 MiB	<code>L = [j ^ (j &gt;&gt; i) for j in range(N)]</code>

Filename: `./mprun_demo.py`

Line #	Mem usage	Increment	Line Contents
=====			
1	39.0 MiB	0.0 MiB	def sum_of_lists(N):
2	39.0 MiB	0.0 MiB	total = 0
3	46.5 MiB	7.5 MiB	for i in range(5):
4	71.9 MiB	25.4 MiB	L = [j ^ (j >> i) for j in range(N)]
5	71.9 MiB	0.0 MiB	total += sum(L)
6	46.5 MiB	-25.4 MiB	del L # remove reference to L
7	39.1 MiB	-7.4 MiB	return total

Here the `Increment` column tells us how much each line affects the total memory budget: observe that when we create and delete the list `L`, we are adding about 25 MB of memory usage. This is on top of the background memory usage from the Python interpreter itself.

For more information on `%memit` and `%mprun`, as well as their available options, use the IPython help functionality (i.e., type `%memit?` at the IPython prompt).

## Introduction to NumPy

This chapter will cover NumPy in detail. NumPy (short for *Numerical Python*) provides an efficient interface to store and operate on dense data buffers. In some ways, NumPy arrays are like Python's built-in `list` type, but NumPy arrays provide much more efficient storage and data operations as the arrays grow larger in size. NumPy arrays form the core of nearly the entire ecosystem of data science tools in Python, so time spent learning to use NumPy effectively will be valuable no matter what aspect of data science interests you.

If you followed the advice outlined in the Preface and installed the Anaconda stack, you already have NumPy installed and ready to go. If you're more the do-it-yourself type, you can go to <http://www.numpy.org/> and follow the installation instructions found there. Once you do, you can import NumPy and double-check the version:

In [1]:



```
import numpy
numpy.__version__
Out[1]:
```

```
'1.11.1'
```

For the pieces of the package discussed here, I'd recommend NumPy version 1.8 or later. By convention, you'll find that most people in the SciPy/PyData world will import NumPy using `np` as an alias:

```
In [2]:
```

```
import numpy as np
```

Throughout this chapter, and indeed the rest of the book, you'll find that this is the way we will import and use NumPy.

## Reminder about Built In Documentation

As you read through this chapter, don't forget that IPython gives you the ability to quickly explore the contents of a package (by using the tab-completion feature), as well as the documentation of various functions

For example, to display all the contents of the numpy namespace, you can type this:

```
In [3]: np.<TAB>
```

And to display NumPy's built-in documentation, you can use this:

```
In [4]: np?
```

## Understanding Data Types in Python

Effective data-driven science and computation requires understanding how data is stored and manipulated. This section outlines and contrasts how arrays of data are handled in the Python language itself, and how NumPy improves on this. Understanding this difference is fundamental to understanding much of the material throughout the rest of the book.

Users of Python are often drawn-in by its ease of use, one piece of which is dynamic typing. While a statically-typed language like C or Java requires each variable to be explicitly declared, a

dynamically-typed language like Python skips this specification. For example, in C you might specify a particular operation as follows:

```
/* C code */  
  
int result = 0;  
  
for(int i=0; i<100; i++){  
    result += i;  
}
```

While in Python the equivalent operation could be written this way:

```
# Python code  
  
result = 0  
  
for i in range(100):  
    result += i
```

Notice the main difference: in C, the data types of each variable are explicitly declared, while in Python the types are dynamically inferred. This means, for example, that we can assign any kind of data to any variable:

```
# Python code  
  
x = 4  
  
x = "four"
```

Here we've switched the contents of `x` from an integer to a string. The same thing in C would lead (depending on compiler settings) to a compilation error or other unintended consequences:

```
/* C code */  
  
int x = 4;  
  
x = "four"; // FAILS
```

This sort of flexibility is one piece that makes Python and other dynamically-typed languages convenient and easy to use. Understanding *how* this works is an important piece of learning to analyze data efficiently and effectively with Python. But what this type-flexibility also points to is the fact that Python variables are more than just their value; they also contain extra information about the type of the value. We'll explore this more in the sections that follow.

## A Python Integer Is More Than Just an Integer

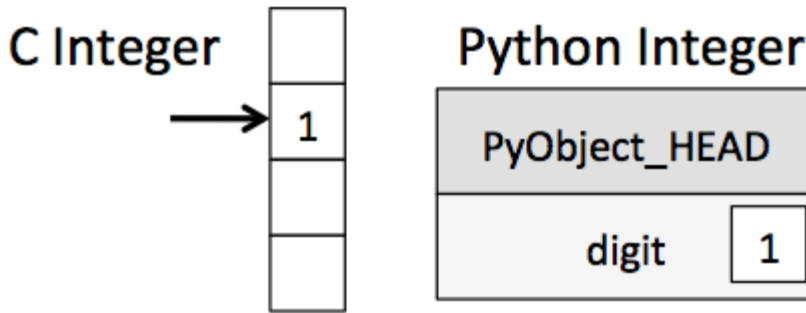
The standard Python implementation is written in C. This means that every Python object is simply a cleverly-disguised C structure, which contains not only its value, but other information as well. For example, when we define an integer in Python, such as `x = 10000`, `x` is not just a "raw" integer. It's actually a pointer to a compound C structure, which contains several values. Looking through the Python 3.4 source code, we find that the integer (long) type definition effectively looks like this (once the C macros are expanded):

```
struct _longobject {  
  
    long ob_refcnt;  
  
    PyTypeObject *ob_type;  
  
    size_t ob_size;  
  
    long ob_digit[1];  
  
};
```

A single integer in Python 3.4 actually contains four pieces:

- `ob_refcnt`, a reference count that helps Python silently handle memory allocation and deallocation
- `ob_type`, which encodes the type of the variable
- `ob_size`, which specifies the size of the following data members
- `ob_digit`, which contains the actual integer value that we expect the Python variable to represent.

This means that there is some overhead in storing an integer in Python as compared to an integer in a compiled language like C, as illustrated in the following figure:



Here `PyObject_HEAD` is the part of the structure containing the reference count, type code, and other pieces mentioned before.

Notice the difference here: a C integer is essentially a label for a position in memory whose bytes encode an integer value. A Python integer is a pointer to a position in memory containing all the Python object information, including the bytes that contain the integer value. This extra information in the Python integer structure is what allows Python to be coded so freely and dynamically. All this additional information in Python types comes at a cost, however, which becomes especially apparent in structures that combine many of these objects.

## A Python List Is More Than Just a List

Let's consider now what happens when we use a Python data structure that holds many Python objects. The standard mutable multi-element container in Python is the list. We can create a list of integers as follows:

In [1]:

```
L = list(range(10))  
L
```

Out[1]:

```
[0, 1, 2, 3, 4, 5, 6, 7, 8, 9]
```

In [2]:

```
type(L[0])
```

Out[2]:

```
int
```

Or, similarly, a list of strings:

In [3]:

```
L2 = [str(c) for c in L]
```

```
L2
```

```
Out[3]:
```

```
['0', '1', '2', '3', '4', '5', '6', '7', '8', '9']
```

```
In [4]:
```

```
type(L2[0])
```

```
Out[4]:
```

```
str
```

Because of Python's dynamic typing, we can even create heterogeneous lists:

```
In [5]:
```

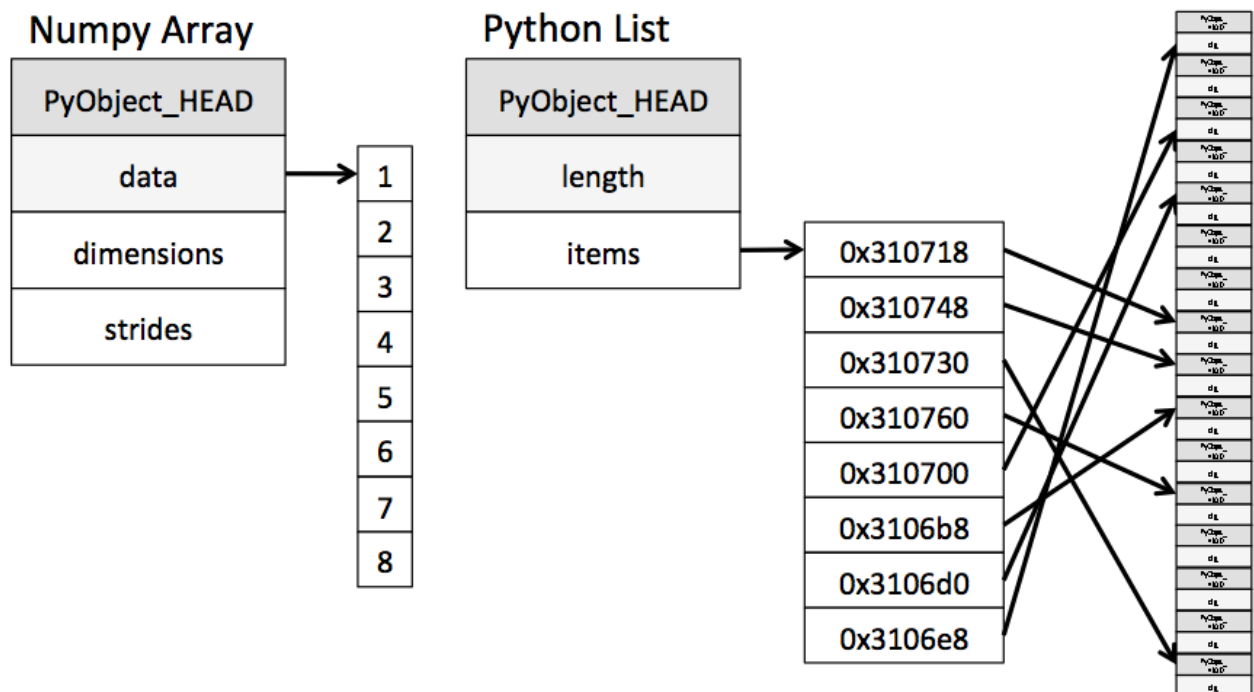
```
L3 = [True, "2", 3.0, 4]
```

```
[type(item) for item in L3]
```

```
Out[5]:
```

```
[bool, str, float, int]
```

But this flexibility comes at a cost: to allow these flexible types, each item in the list must contain its own type info, reference count, and other information—that is, each item is a complete Python object. In the special case that all variables are of the same type, much of this information is redundant: it can be much more efficient to store data in a fixed-type array. The difference between a dynamic-type list and a fixed-type (NumPy-style) array is illustrated in the following figure:



At the implementation level, the array essentially contains a single pointer to one contiguous block of data. The Python list, on the other hand, contains a pointer to a block of pointers, each of which in turn points to a full Python object like the Python integer we saw earlier. Again, the advantage of the list is flexibility: because each list element is a full structure containing both data and type information, the list can be filled with data of any desired type. Fixed-type NumPy-style arrays lack this flexibility, but are much more efficient for storing and manipulating data.

## Fixed-Type Arrays in Python

Python offers several different options for storing data in efficient, fixed-type data buffers. The built-in `array` module (available since Python 3.3) can be used to create dense arrays of a uniform type:

In [6]:

```
import array
L = list(range(10))
A = array.array('i', L)
A
```

Out[6]:

```
array('i', [0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
```

Here `'i'` is a type code indicating the contents are integers.

Much more useful, however, is the `ndarray` object of the NumPy package. While Python's `array` object provides efficient storage of array-based data, NumPy adds to this efficient *operations* on that data. We will explore these operations in later sections; here we'll demonstrate several ways of creating a NumPy array.

We'll start with the standard NumPy import, under the alias `np`:

In [7]:

```
import numpy as np
```

## Creating Arrays from Python Lists

First, we can use `np.array` to create arrays from Python lists:

In [8]:

```
# integer array:  
np.array([1, 4, 2, 5, 3])
```

Out[8]:

```
array([1, 4, 2, 5, 3])
```

Remember that unlike Python lists, NumPy is constrained to arrays that all contain the same type. If types do not match, NumPy will upcast if possible (here, integers are up-cast to floating point):

In [9]:

```
np.array([3.14, 4, 2, 3])
```

Out[9]:

```
array([ 3.14,  4. ,  2. ,  3. ])
```

If we want to explicitly set the data type of the resulting array, we can use the `dtype` keyword:

In [10]:

```
np.array([1, 2, 3, 4], dtype='float32')
```

Out[10]:

```
array([ 1.,  2.,  3.,  4.], dtype=float32)
```

Finally, unlike Python lists, NumPy arrays can explicitly be multi-dimensional; here's one way of initializing a multidimensional array using a list of lists:

In [11]:

```
# nested lists result in multi-dimensional arrays
```

```
np.array([range(i, i + 3) for i in [2, 4, 6]])
```

Out[11]:

```
array([[2, 3, 4],  
       [4, 5, 6],  
       [6, 7, 8]])
```

The inner lists are treated as rows of the resulting two-dimensional array.

## Creating Arrays from Scratch

Especially for larger arrays, it is more efficient to create arrays from scratch using routines built into NumPy. Here are several examples:

In [12]:

```
# Create a length-10 integer array filled with zeros
```

```
np.zeros(10, dtype=int)
```

Out[12]:

```
array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0])
```

In [13]:

```
# Create a 3x5 floating-point array filled with ones  
np.ones((3, 5), dtype=float)
```

Out[13]:

```
array([[ 1.,  1.,  1.,  1.,  1.],  
       [ 1.,  1.,  1.,  1.,  1.],  
       [ 1.,  1.,  1.,  1.,  1.]])
```

In [14]:

```
# Create a 3x5 array filled with 3.14  
np.full((3, 5), 3.14)
```

Out[14]:

```
array([[ 3.14,  3.14,  3.14,  3.14,  3.14],  
       [ 3.14,  3.14,  3.14,  3.14,  3.14],  
       [ 3.14,  3.14,  3.14,  3.14,  3.14]])
```

In [15]:

```
# Create an array filled with a linear sequence  
# Starting at 0, ending at 20, stepping by 2  
# (this is similar to the built-in range() function)  
np.arange(0, 20, 2)
```

Out[15]:

```
array([ 0,  2,  4,  6,  8, 10, 12, 14, 16, 18])
```

In [16]:

```
# Create an array of five values evenly spaced between 0 and 1  
np.linspace(0, 1, 5)
```

Out[16]:

```
array([ 0. , 0.25, 0.5 , 0.75,  1.  ])
```

In [17]:

```
# Create a 3x3 array of uniformly distributed  
# random values between 0 and 1  
np.random.random((3, 3))
```

Out[17]:

```
array([[ 0.99844933,  0.52183819,  0.22421193],  
       [ 0.08007488,  0.45429293,  0.20941444],  
       [ 0.14360941,  0.96910973,  0.946117  ]])
```

In [18]:

```
# Create a 3x3 array of normally distributed random values  
# with mean 0 and standard deviation 1  
np.random.normal(0, 1, (3, 3))
```

Out[18]:

```
array([[ 1.51772646,  0.39614948, -0.10634696],  
       [ 0.25671348,  0.00732722,  0.37783601],  
       [ 0.68446945,  0.15926039, -0.70744073]])
```

In [19]:



```
# Create a 3x3 array of random integers in the interval [0, 10)  
np.random.randint(0, 10, (3, 3))
```

Out[19]:

```
array([[2, 3, 4],  
       [5, 7, 8],  
       [0, 5, 0]])
```

In [20]:

```
# Create a 3x3 identity matrix  
np.eye(3)
```

Out[20]:

```
array([[ 1.,  0.,  0.],  
       [ 0.,  1.,  0.],  
       [ 0.,  0.,  1.]])
```

In [21]:

```
# Create an uninitialized array of three integers  
# The values will be whatever happens to already exist at that memory location  
np.empty(3)
```

Out[21]:

```
array([ 1.,  1.,  1.])
```

## NumPy Standard Data Types

NumPy arrays contain values of a single type, so it is important to have detailed knowledge of those types and their limitations. Because NumPy is built in C, the types will be familiar to users of C, Fortran, and other related languages.

The standard NumPy data types are listed in the following table. Note that when constructing an array, they can be specified using a string:

```
np.zeros(10, dtype='int16')
```

Or using the associated NumPy object:

```
np.zeros(10, dtype=np.int16)
```

### Data type Description

bool_	Boolean (True or False) stored as a byte
-------	--

## Data type Description

<code>int</code>	Default integer type (same as C <code>long</code> ; normally either <code>int64</code> or <code>int32</code> )
<code>intc</code>	Identical to C <code>int</code> (normally <code>int32</code> or <code>int64</code> )
<code>intp</code>	Integer used for indexing (same as C <code>ssize_t</code> ; normally either <code>int32</code> or <code>int64</code> )
<code>int8</code>	Byte (-128 to 127)
<code>int16</code>	Integer (-32768 to 32767)
<code>int32</code>	Integer (-2147483648 to 2147483647)
<code>int64</code>	Integer (-9223372036854775808 to 9223372036854775807)
<code>uint8</code>	Unsigned integer (0 to 255)
<code>uint16</code>	Unsigned integer (0 to 65535)
<code>uint32</code>	Unsigned integer (0 to 4294967295)
<code>uint64</code>	Unsigned integer (0 to 18446744073709551615)
<code>float</code>	Shorthand for <code>float64</code> .
<code>float16</code>	Half precision float: sign bit, 5 bits exponent, 10 bits mantissa
<code>float32</code>	Single precision float: sign bit, 8 bits exponent, 23 bits mantissa
<code>float64</code>	Double precision float: sign bit, 11 bits exponent, 52 bits mantissa
<code>complex</code>	Shorthand for <code>complex128</code> .
<code>complex64</code>	Complex number, represented by two 32-bit floats
<code>complex128</code>	Complex number, represented by two 64-bit floats

# The Basics of NumPy Arrays

Data manipulation in Python is nearly synonymous with NumPy array manipulation: even newer tools like Pandas are built around the NumPy array. This section will present several examples of using NumPy array manipulation to access data and subarrays, and to split, reshape, and join the arrays. While the types of operations shown here may seem a bit dry and pedantic, they comprise the building blocks of many other examples used throughout the book. Get to know them well!

We'll cover a few categories of basic array manipulations here:

- *Attributes of arrays*: Determining the size, shape, memory consumption, and data types of arrays
- *Indexing of arrays*: Getting and setting the value of individual array elements
- *Slicing of arrays*: Getting and setting smaller subarrays within a larger array
- *Reshaping of arrays*: Changing the shape of a given array
- *Joining and splitting of arrays*: Combining multiple arrays into one, and splitting one array into many

## NumPy Array Attributes

First let's discuss some useful array attributes. We'll start by defining three random arrays, a one-dimensional, two-dimensional, and three-dimensional array. We'll use NumPy's random number generator, which we will *seed* with a set value in order to ensure that the same random arrays are generated each time this code is run:

In [1]:

```
import numpy as np
np.random.seed(0) # seed for reproducibility

x1 = np.random.randint(10, size=6) # One-dimensional array
x2 = np.random.randint(10, size=(3, 4)) # Two-dimensional array
x3 = np.random.randint(10, size=(3, 4, 5)) # Three-dimensional array
```

Each array has attributes `ndim` (the number of dimensions), `shape` (the size of each dimension), and `size` (the total size of the array):

In [2]:

```
print("x3 ndim: ", x3.ndim)
print("x3 shape:", x3.shape)
print("x3 size: ", x3.size)
```

```
x3.ndim: 3
x3.shape: (3, 4, 5)
x3.size: 60
```

Another useful attribute is the `dtype`, the data type of the array (which we discussed previously in [Understanding Data Types in Python](#)):

In [3]:

```
print("dtype:", x3.dtype)
dtype: int64
```

Other attributes include `itemsize`, which lists the size (in bytes) of each array element, and `nbytes`, which lists the total size (in bytes) of the array:

In [4]:

```
print("itemsize:", x3.itemsize, "bytes")
print("nbytes:", x3.nbytes, "bytes")
itemsize: 8 bytes
nbytes: 480 bytes
```

In general, we expect that `nbytes` is equal to `itemsize` times `size`.

## Array Indexing: Accessing Single Elements

If you are familiar with Python's standard list indexing, indexing in NumPy will feel quite familiar. In a one-dimensional array, the  $i$ th value (counting from zero) can be accessed by specifying the desired index in square brackets, just as with Python lists:

In [5]:

```
x1
Out[5]:
array([5, 0, 3, 3, 7, 9])
```

In [6]:

```
x1[0]
Out[6]:
```

```
5
```

In [7]:

```
x1[4]
Out[7]:
```

```
7
```

To index from the end of the array, you can use negative indices:

In [8]:

```
x1[-1]
```

```
Out[8]:
```

```
9
```

```
In [9]:
```

```
x1[-2]
```

```
Out[9]:
```

```
7
```

In a multi-dimensional array, items can be accessed using a comma-separated tuple of indices:

```
In [10]:
```

```
x2
```

```
Out[10]:
```

```
array([[3, 5, 2, 4],  
       [7, 6, 8, 8],  
       [1, 6, 7, 7]])
```

```
In [11]:
```

```
x2[0, 0]
```

```
Out[11]:
```

```
3
```

```
In [12]:
```

```
x2[2, 0]
```

```
Out[12]:
```

```
1
```

```
In [13]:
```

```
x2[2, -1]
```

```
Out[13]:
```

```
7
```

Values can also be modified using any of the above index notation:

```
In [14]:
```

```
x2[0, 0] = 12
```

```
x2
```

```
Out[14]:
```

```
array([[12, 5, 2, 4],  
       [ 7, 6, 8, 8],  
       [ 1, 6, 7, 7]])
```

Keep in mind that, unlike Python lists, NumPy arrays have a fixed type. This means, for example, that if you attempt to insert a floating-point value to an integer array, the value will be silently truncated. Don't be caught unaware by this behavior!

In [15]:

```
x1[0] = 3.14159 # this will be truncated!  
x1
```

Out[15]:

```
array([3, 0, 3, 3, 7, 9])
```

## Array Slicing: Accessing Subarrays

Just as we can use square brackets to access individual array elements, we can also use them to access subarrays with the *slice* notation, marked by the colon (:) character. The NumPy slicing syntax follows that of the standard Python list; to access a slice of an array `x`, use this:

```
x[start:stop:step]
```

If any of these are unspecified, they default to the values `start=0`, `stop=size of dimension`, `step=1`. We'll take a look at accessing sub-arrays in one dimension and in multiple dimensions.

### One-dimensional subarrays

In [16]:

```
x = np.arange(10)  
x
```

Out[16]:

```
array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
```

In [17]:

```
x[:5] # first five elements
```

Out[17]:

```
array([0, 1, 2, 3, 4])
```

In [18]:

```
x[5:] # elements after index 5
```

Out[18]:

```
array([5, 6, 7, 8, 9])
```

In [19]:

```
x[4:7] # middle sub-array
```

Out[19]:

```
array([4, 5, 6])
```

In [20]:

```
x[::2] # every other element
```

Out[20]:

```
array([0, 2, 4, 6, 8])
```

In [21]:

```
x[1::2] # every other element, starting at index 1
```

Out[21]:

```
array([1, 3, 5, 7, 9])
```

A potentially confusing case is when the `step` value is negative. In this case, the defaults for `start` and `stop` are swapped. This becomes a convenient way to reverse an array:

In [22]:

```
x[::-1] # all elements, reversed
```

Out[22]:

```
array([9, 8, 7, 6, 5, 4, 3, 2, 1, 0])
```

In [23]:

```
x[5::-2] # reversed every other from index 5
```

Out[23]:

```
array([5, 3, 1])
```

## Multi-dimensional subarrays

Multi-dimensional slices work in the same way, with multiple slices separated by commas. For example:

In [24]:

```
x2
```

Out[24]:

```
array([[12, 5, 2, 4],  
       [ 7, 6, 8, 8],  
       [ 1, 6, 7, 7]])
```

In [25]:

```
x2[:,2, :3] # two rows, three columns
```

Out[25]:

```
array([[12, 5, 2],  
       [ 7, 6, 8]])
```

In [26]:

```
x2[:,3, ::2] # all rows, every other column
```

Out[26]:

```
array([[12, 2],  
       [ 7, 8],  
       [ 1, 7]])
```

Finally, subarray dimensions can even be reversed together:

In [27]:

```
x2[::-1, ::-1]
```

Out[27]:

```
array([[ 7,  7,  6,  1],
       [ 8,  8,  6,  7],
       [ 4,  2,  5, 12]])
```

## Accessing array rows and columns

One commonly needed routine is accessing of single rows or columns of an array. This can be done by combining indexing and slicing, using an empty slice marked by a single colon (:):

In [28]:

```
print(x2[:, 0]) # first column of x2
[12  7  1]
```

In [29]:

```
print(x2[0, :]) # first row of x2
[12  5  2  4]
```

In the case of row access, the empty slice can be omitted for a more compact syntax:

In [30]:

```
print(x2[0]) # equivalent to x2[0, :]
[12  5  2  4]
```

## Subarrays as no-copy views

One important—and extremely useful—thing to know about array slices is that they return *views* rather than *copies* of the array data. This is one area in which NumPy array slicing differs from Python list slicing: in lists, slices will be copies. Consider our two-dimensional array from before:

In [31]:

```
print(x2)
[[12  5  2  4]
 [ 7  6  8  8]
 [ 1  6  7  7]]
```

Let's extract a  $2 \times 2$  subarray from this:

In [32]:

```
x2_sub = x2[:2, :2]
print(x2_sub)
[[12  5]
 [ 7  6]]
```



Now if we modify this subarray, we'll see that the original array is changed! Observe:

In [33]:

```
x2_sub[0, 0] = 99
print(x2_sub)
[[99 5]
 [ 7 6]]
```

In [34]:

```
print(x2)
[[99 5 2 4]
 [ 7 6 8 8]
 [ 1 6 7 7]]
```

This default behavior is actually quite useful: it means that when we work with large datasets, we can access and process pieces of these datasets without the need to copy the underlying data buffer.

## Creating copies of arrays

Despite the nice features of array views, it is sometimes useful to instead explicitly copy the data within an array or a subarray. This can be most easily done with the `copy()` method:

In [35]:

```
x2_sub_copy = x2[:,2, :2].copy()
print(x2_sub_copy)
[[99 5]
 [ 7 6]]
```

If we now modify this subarray, the original array is not touched:

In [36]:

```
x2_sub_copy[0, 0] = 42
print(x2_sub_copy)
[[42 5]
 [ 7 6]]
```

In [37]:

```
print(x2)
[[99 5 2 4]
 [ 7 6 8 8]
 [ 1 6 7 7]]
```

## Reshaping of Arrays

Another useful type of operation is reshaping of arrays. The most flexible way of doing this is with the `reshape` method. For example, if you want to put the numbers 1 through 9 in a  $3 \times 3 \times 3$  grid, you can do the following:

In [38]:

```
grid = np.arange(1, 10).reshape((3, 3))
print(grid)
[[1 2 3]
 [4 5 6]
 [7 8 9]]
```

Note that for this to work, the size of the initial array must match the size of the reshaped array. Where possible, the `reshape` method will use a no-copy view of the initial array, but with non-contiguous memory buffers this is not always the case.

Another common reshaping pattern is the conversion of a one-dimensional array into a two-dimensional row or column matrix. This can be done with the `reshape` method, or more easily done by making use of the `newaxis` keyword within a slice operation:

In [39]:

```
x = np.array([1, 2, 3])

# row vector via reshape
x.reshape((1, 3))
```

Out[39]:

```
array([[1, 2, 3]])
```

In [40]:

```
# row vector via newaxis
x[np.newaxis, :]
```

Out[40]:

```
array([[1, 2, 3]])
```

In [41]:

```
# column vector via reshape
x.reshape((3, 1))
```

Out[41]:

```
array([[1],
       [2],
       [3]])
```

In [42]:

```
# column vector via newaxis
x[:, np.newaxis]
```

Out[42]:

```
array([[1],
```

```
[2],  
[3]])
```

We will see this type of transformation often throughout the remainder of the book.

## Array Concatenation and Splitting

All of the preceding routines worked on single arrays. It's also possible to combine multiple arrays into one, and to conversely split a single array into multiple arrays. We'll take a look at those operations here.

### Concatenation of arrays

Concatenation, or joining of two arrays in NumPy, is primarily accomplished using the routines `np.concatenate`, `np.vstack`, and `np.hstack`. `np.concatenate` takes a tuple or list of arrays as its first argument, as we can see here:

In [43]:

```
x = np.array([1, 2, 3])  
y = np.array([3, 2, 1])  
np.concatenate([x, y])
```

Out[43]:

```
array([1, 2, 3, 3, 2, 1])
```

You can also concatenate more than two arrays at once:

In [44]:

```
z = [99, 99, 99]  
print(np.concatenate([x, y, z]))  
[ 1  2  3  3  2  1 99 99 99]
```

It can also be used for two-dimensional arrays:

In [45]:

```
grid = np.array([[1, 2, 3],  
                 [4, 5, 6]])
```

In [46]:

```
# concatenate along the first axis  
np.concatenate([grid, grid])
```

Out[46]:

```
array([[1, 2, 3],  
       [4, 5, 6],  
       [1, 2, 3],  
       [4, 5, 6]])
```

```
[4, 5, 6]])
```

In [47]:

```
# concatenate along the second axis (zero-indexed)
np.concatenate([grid, grid], axis=1)
```

Out[47]:

```
array([[1, 2, 3, 1, 2, 3],
       [4, 5, 6, 4, 5, 6]])
```

For working with arrays of mixed dimensions, it can be clearer to use the `np.vstack` (vertical stack) and `np.hstack` (horizontal stack) functions:

In [48]:

```
x = np.array([1, 2, 3])
grid = np.array([[9, 8, 7],
                 [6, 5, 4]])
```

```
# vertically stack the arrays
np.vstack([x, grid])
```

Out[48]:

```
array([[1, 2, 3],
       [9, 8, 7],
       [6, 5, 4]])
```

In [49]:

```
# horizontally stack the arrays
y = np.array([[99],
              [99]])
np.hstack([grid, y])
```

Out[49]:

```
array([[ 9,  8,  7, 99],
       [ 6,  5,  4, 99]])
```

Similarly, `np.dstack` will stack arrays along the third axis.

## Splitting of arrays

The opposite of concatenation is splitting, which is implemented by the functions `np.split`, `np.hsplit`, and `np.vsplit`. For each of these, we can pass a list of indices giving the split points:

In [50]:

```
x = [1, 2, 3, 99, 99, 3, 2, 1]
x1, x2, x3 = np.split(x, [3, 5])
print(x1, x2, x3)
[1 2 3] [99 99] [3 2 1]
```

Notice that  $N$  split-points, leads to  $N + 1$  subarrays. The related functions `np.hsplit` and `np.vsplit` are similar:

In [51]:

```
grid = np.arange(16).reshape((4, 4))  
grid
```

Out[51]:

```
array([[ 0,  1,  2,  3],  
       [ 4,  5,  6,  7],  
       [ 8,  9, 10, 11],  
       [12, 13, 14, 15]])
```

In [52]:

```
upper, lower = np.vsplit(grid, [2])  
print(upper)  
print(lower)  
[[0 1 2 3]  
 [4 5 6 7]]  
[[ 8  9 10 11]  
 [12 13 14 15]]
```

In [53]:

```
left, right = np.hsplit(grid, [2])  
print(left)  
print(right)  
[[ 0  1]  
 [ 4  5]  
 [ 8  9]  
 [12 13]]  
[[ 2  3]  
 [ 6  7]  
 [10 11]  
 [14 15]]
```

Similarly, `np.dsplit` will split arrays along the third axis.

## Computation on NumPy Arrays: Universal Functions

Up until now, we have been discussing some of the basic nuts and bolts of NumPy; in the next few sections, we will dive into the reasons that NumPy is so important in the Python data science world. Namely, it provides an easy and flexible interface to optimized computation with arrays of data.

Computation on NumPy arrays can be very fast, or it can be very slow. The key to making it fast is to use *vectorized* operations, generally implemented through NumPy's *universal functions* (ufuncs). This section motivates the need for NumPy's ufuncs, which can be used to make repeated calculations on array elements much more efficient. It then introduces many of the most common and useful arithmetic ufuncs available in the NumPy package.

# The Slowness of Loops

Python's default implementation (known as CPython) does some operations very slowly. This is in part due to the dynamic, interpreted nature of the language: the fact that types are flexible, so that sequences of operations cannot be compiled down to efficient machine code as in languages like C and Fortran. Recently there have been various attempts to address this weakness: well-known examples are the [PyPy](#) project, a just-in-time compiled implementation of Python; the [Cython](#) project, which converts Python code to compilable C code; and the [Numba](#) project, which converts snippets of Python code to fast LLVM bytecode. Each of these has its strengths and weaknesses, but it is safe to say that none of the three approaches has yet surpassed the reach and popularity of the standard CPython engine.

The relative sluggishness of Python generally manifests itself in situations where many small operations are being repeated – for instance looping over arrays to operate on each element. For example, imagine we have an array of values and we'd like to compute the reciprocal of each. A straightforward approach might look like this:

In [1]:

```
import numpy as np
np.random.seed(0)

def compute_reciprocals(values):
    output = np.empty(len(values))
    for i in range(len(values)):
        output[i] = 1.0 / values[i]
    return output

values = np.random.randint(1, 10, size=5)
compute_reciprocals(values)
```

Out[1]:

```
array([ 0.16666667, 1.        , 0.25       , 0.25       , 0.125      ])
```

This implementation probably feels fairly natural to someone from, say, a C or Java background. But if we measure the execution time of this code for a large input, we see that this operation is very slow, perhaps surprisingly so! We'll benchmark this with IPython's `%timeit` magic (discussed in [Profiling and Timing Code](#)):

In [2]:

```
big_array = np.random.randint(1, 100, size=1000000)
%timeit compute_reciprocals(big_array)
```

1 loop, best of 3: 2.91 s per loop

It takes several seconds to compute these million operations and to store the result! When even cell phones have processing speeds measured in Giga-FLOPS (i.e., billions of numerical operations per second), this seems almost absurdly slow. It turns out that the bottleneck here is not the operations themselves, but the type-checking and function dispatches that CPython must do at each cycle of the loop. Each time the reciprocal is computed, Python first examines the object's type and does a dynamic lookup of the correct function to use for that type. If we were working in compiled code instead, this type specification would be known before the code executes and the result could be computed much more efficiently.

## Introducing UFuncs

For many types of operations, NumPy provides a convenient interface into just this kind of statically typed, compiled routine. This is known as a *vectorized* operation. This can be accomplished by simply performing an operation on the array, which will then be applied to each element. This vectorized approach is designed to push the loop into the compiled layer that underlies NumPy, leading to much faster execution.

Compare the results of the following two:

In [3]:

```
print(compute_reciprocals(values))
print(1.0 / values)
[ 0.16666667  1.      0.25    0.25    0.125   ]
[ 0.16666667  1.      0.25    0.25    0.125   ]
```

Looking at the execution time for our big array, we see that it completes orders of magnitude faster than the Python loop:

In [4]:

```
%timeit (1.0 / big_array)
100 loops, best of 3: 4.6 ms per loop
```

Vectorized operations in NumPy are implemented via *ufuncs*, whose main purpose is to quickly execute repeated operations on values in NumPy arrays. Ufuncs are extremely flexible – before we saw an operation between a scalar and an array, but we can also operate between two arrays:

In [5]:

```
np.arange(5) / np.arange(1, 6)
```

Out[5]:

```
array([ 0.        , 0.5        , 0.66666667, 0.75        , 0.8        ])
```

And ufunc operations are not limited to one-dimensional arrays—they can also act on multi-dimensional arrays as well:

In [6]:

```
x = np.arange(9).reshape((3, 3))
2 ** x
```

Out[6]:

```
array([[ 1,  2,  4],
       [ 8, 16, 32],
       [64, 128, 256]])
```

Computations using vectorization through ufuncs are nearly always more efficient than their counterpart implemented using Python loops, especially as the arrays grow in size. Any time you see such a loop in a Python script, you should consider whether it can be replaced with a vectorized expression.

## Exploring NumPy's UFuncs

Ufuncs exist in two flavors: *unary ufuncs*, which operate on a single input, and *binary ufuncs*, which operate on two inputs. We'll see examples of both these types of functions here.

### Array arithmetic

NumPy's ufuncs feel very natural to use because they make use of Python's native arithmetic operators. The standard addition, subtraction, multiplication, and division can all be used:

In [7]:

```
x = np.arange(4)
print("x =", x)
print("x + 5 =", x + 5)
print("x - 5 =", x - 5)
print("x * 2 =", x * 2)
print("x / 2 =", x / 2)
print("x // 2 =", x // 2) #floor division
x = [0 1 2 3]
x + 5 = [5 6 7 8]
x - 5 = [-5 -4 -3 -2]
x * 2 = [0 2 4 6]
x / 2 = [ 0.  0.5  1.  1.5]
x // 2 = [0 0 1 1]
```



There is also a unary ufunc for negation, and a `**` operator for exponentiation, and a `%` operator for modulus:

In [8]:

```
print("-x = ", -x)
print("x ** 2 = ", x ** 2)
print("x % 2 = ", x % 2)
-x = [ 0 -1 -2 -3]
x ** 2 = [0 1 4 9]
x % 2 = [0 1 0 1]
```

In addition, these can be strung together however you wish, and the standard order of operations is respected:

In [9]:

```
-(0.5*x + 1) ** 2
```

Out[9]:

```
array([-1. , -2.25, -4. , -6.25])
```

Each of these arithmetic operations are simply convenient wrappers around specific functions built into NumPy; for example, the `+` operator is a wrapper for the `add` function:

In [10]:

```
np.add(x, 2)
```

Out[10]:

```
array([2, 3, 4, 5])
```

The following table lists the arithmetic operators implemented in NumPy:

Operator	Equivalent ufunc	Description
----------	------------------	-------------

<code>+</code>	<code>np.add</code>	Addition (e.g., <code>1 + 1 = 2</code> )
<code>-</code>	<code>np.subtract</code>	Subtraction (e.g., <code>3 - 2 = 1</code> )
<code>-</code>	<code>np.negative</code>	Unary negation (e.g., <code>-2</code> )
<code>*</code>	<code>np.multiply</code>	Multiplication (e.g., <code>2 * 3 = 6</code> )
<code>/</code>	<code>np.divide</code>	Division (e.g., <code>3 / 2 = 1.5</code> )

## OperatorEquivalent ufuncDescription

//	np.floor_divide	Floor division (e.g., <code>3 // 2 = 1</code> )
**	np.power	Exponentiation (e.g., <code>2 ** 3 = 8</code> )
%	np.mod	Modulus/remainder (e.g., <code>9 % 4 = 1</code> )

Additionally there are Boolean/bitwise operators; we will explore these in [Comparisons, Masks, and Boolean Logic](#).

## Absolute value

Just as NumPy understands Python's built-in arithmetic operators, it also understands Python's built-in absolute value function:

In [11]:

```
x = np.array([-2, -1, 0, 1, 2])
abs(x)
```

Out[11]:

```
array([2, 1, 0, 1, 2])
```

The corresponding NumPy ufunc is `np.absolute`, which is also available under the alias `np.abs`:

In [12]:

```
np.absolute(x)
```

Out[12]:

```
array([2, 1, 0, 1, 2])
```

In [13]:

```
np.abs(x)
```

Out[13]:

```
array([2, 1, 0, 1, 2])
```

This ufunc can also handle complex data, in which the absolute value returns the magnitude:

In [14]:

```
x = np.array([3 - 4j, 4 - 3j, 2 + 0j, 0 + 1j])
np.abs(x)
```

Out[14]:

```
array([ 5., 5., 2., 1.]
```

## Trigonometric functions

NumPy provides a large number of useful ufuncs, and some of the most useful for the data scientist are the trigonometric functions. We'll start by defining an array of angles:

In [15]:

```
theta = np.linspace(0, np.pi, 3)
```

Now we can compute some trigonometric functions on these values:

In [16]:

```
print("theta    = ", theta)
print("sin(theta) = ", np.sin(theta))
print("cos(theta) = ", np.cos(theta))
print("tan(theta) = ", np.tan(theta))
theta    = [ 0.          1.57079633  3.14159265]
sin(theta) = [ 0.00000000e+00  1.00000000e+00  1.22464680e-16]
cos(theta) = [ 1.00000000e+00  6.12323400e-17 -1.00000000e+00]
tan(theta) = [ 0.00000000e+00  1.63312394e+16 -1.22464680e-16]
```

The values are computed to within machine precision, which is why values that should be zero do not always hit exactly zero. Inverse trigonometric functions are also available:

In [17]:

```
x = [-1, 0, 1]
print("x        = ", x)
print("arcsin(x) = ", np.arcsin(x))
print("arccos(x) = ", np.arccos(x))
print("arctan(x) = ", np.arctan(x))
x      = [-1, 0, 1]
arcsin(x) = [-1.57079633  0.          1.57079633]
arccos(x) = [ 3.14159265  1.57079633  0.          ]
arctan(x) = [-0.78539816  0.          0.78539816]
```

## Exponents and logarithms

Another common type of operation available in a NumPy ufunc are the exponentials:

In [18]:

```
x = [1, 2, 3]
print("x      =", x)
print("e^x    =", np.exp(x))
print("2^x    =", np.exp2(x))
print("3^x    =", np.power(3, x))
x      = [1, 2, 3]
```

```
e^x = [ 2.71828183  7.3890561 20.08553692]
2^x = [ 2.  4.  8.]
3^x = [ 3  9 27]
```

The inverse of the exponentials, the logarithms, are also available. The basic `np.log` gives the natural logarithm; if you prefer to compute the base-2 logarithm or the base-10 logarithm, these are available as well:

In [19]:

```
x = [1, 2, 4, 10]
print("x =", x)
print("ln(x) =", np.log(x))
print("log2(x) =", np.log2(x))
print("log10(x) =", np.log10(x))
x = [1, 2, 4, 10]
ln(x) = [ 0.          0.69314718  1.38629436  2.30258509]
log2(x) = [ 0.          1.          2.          3.32192809]
log10(x) = [ 0.          0.30103   0.60205999  1.          ]
```

There are also some specialized versions that are useful for maintaining precision with very small input:

In [20]:

```
x = [0, 0.001, 0.01, 0.1]
print("exp(x) - 1 =", np.expm1(x))
print("log(1 + x) =", np.log1p(x))
exp(x) - 1 = [ 0.          0.0010005  0.01005017  0.10517092]
log(1 + x) = [ 0.          0.0009995  0.00995033  0.09531018]
```

When `x` is very small, these functions give more precise values than if the raw `np.log` or `np.exp` were to be used.

## Specialized ufuncs

NumPy has many more ufuncs available, including hyperbolic trig functions, bitwise arithmetic, comparison operators, conversions from radians to degrees, rounding and remainders, and much more. A look through the NumPy documentation reveals a lot of interesting functionality.

Another excellent source for more specialized and obscure ufuncs is the submodule `scipy.special`. If you want to compute some obscure mathematical function on your data, chances are it is implemented in `scipy.special`. There are far too many functions to list them all, but the following snippet shows a couple that might come up in a statistics context:

In [21]:

```
from scipy import special
```

In [22]:

```
# Gamma functions (generalized factorials) and related functions
x = [1, 5, 10]
print("gamma(x) =", special.gamma(x))
print("ln|gamma(x)| =", special.gammaln(x))
print("beta(x, 2) =", special.beta(x, 2))
gamma(x) = [ 1.00000000e+00  2.40000000e+01  3.62880000e+05]
ln|gamma(x)| = [ 0.          3.17805383 12.80182748]
beta(x, 2) = [ 0.5        0.03333333 0.00909091]
```

In [23]:

```
# Error function (integral of Gaussian)
# its complement, and its inverse
x = np.array([0, 0.3, 0.7, 1.0])
print("erf(x) =", special.erf(x))
print("erfc(x) =", special.erfc(x))
print("erfinv(x) =", special.erfinv(x))
erf(x) = [ 0.          0.32862676 0.67780119 0.84270079]
erfc(x) = [ 1.          0.67137324 0.32219881 0.15729921]
erfinv(x) = [ 0.          0.27246271 0.73286908      inf]
```

There are many, many more ufuncs available in both NumPy and `scipy.special`. Because the documentation of these packages is available online, a web search along the lines of "gamma function python" will generally find the relevant information.

## Advanced Ufunc Features

Many NumPy users make use of ufuncs without ever learning their full set of features. We'll outline a few specialized features of ufuncs here.

### Specifying output

For large calculations, it is sometimes useful to be able to specify the array where the result of the calculation will be stored. Rather than creating a temporary array, this can be used to write computation results directly to the memory location where you'd like them to be. For all ufuncs, this can be done using the `out` argument of the function:

In [24]:

```
x = np.arange(5)
y = np.empty(5)
np.multiply(x, 10, out=y)
print(y)
[ 0. 10. 20. 30. 40.]
```

This can even be used with array views. For example, we can write the results of a computation to every other element of a specified array:

In [25]:

```
y = np.zeros(10)
np.power(2, x, out=y[::2])
print(y)
[ 1.  0.  2.  0.  4.  0.  8.  0. 16.  0.]
```

If we had instead written `y[::2] = 2 ** x`, this would have resulted in the creation of a temporary array to hold the results of `2 ** x`, followed by a second operation copying those values into the `y` array. This doesn't make much of a difference for such a small computation, but for very large arrays the memory savings from careful use of the `out` argument can be significant.

## Aggregates

For binary ufuncs, there are some interesting aggregates that can be computed directly from the object. For example, if we'd like to *reduce* an array with a particular operation, we can use the `reduce` method of any ufunc. A reduce repeatedly applies a given operation to the elements of an array until only a single result remains.

For example, calling `reduce` on the `add` ufunc returns the sum of all elements in the array:

In [26]:

```
x = np.arange(1, 6)
np.add.reduce(x)
Out[26]:
```

15

Similarly, calling `reduce` on the `multiply` ufunc results in the product of all array elements:

In [27]:

```
np.multiply.reduce(x)
Out[27]:
```

120

If we'd like to store all the intermediate results of the computation, we can instead use `accumulate`:

In [28]:

```
np.add.accumulate(x)
Out[28]:
```

```
array([ 1,  3,  6, 10, 15])
```

In [29]:

```
np.multiply.accumulate(x)
```

Out[29]:

```
array([ 1,  2,  6, 24, 120])
```

Note that for these particular cases, there are dedicated NumPy functions to compute the results (`np.sum`, `np.prod`, `np.cumsum`, `np.cumprod`), which we'll explore in [Aggregations: Min, Max, and Everything In Between](#).

## Outer products

Finally, any ufunc can compute the output of all pairs of two different inputs using the `outer` method. This allows you, in one line, to do things like create a multiplication table:

In [30]:

```
x = np.arange(1, 6)
np.multiply.outer(x, x)
```

Out[30]:

```
array([[ 1,  2,  3,  4,  5],
       [ 2,  4,  6,  8, 10],
       [ 3,  6,  9, 12, 15],
       [ 4,  8, 12, 16, 20],
       [ 5, 10, 15, 20, 25]])
```

The `ufunc.at` and `ufunc.reduceat` methods, which we'll explore in [Fancy Indexing](#), are very helpful as well.

Another extremely useful feature of ufuncs is the ability to operate between arrays of different sizes and shapes, a set of operations known as *broadcasting*. This subject is important enough that we will devote a whole section to it (see [Computation on Arrays: Broadcasting](#)).

## Ufuncs: Learning More

More information on universal functions (including the full list of available functions) can be found on the [NumPy](#) and [SciPy](#) documentation websites.

# Aggregations: Min, Max, and Everything In Between

Often when faced with a large amount of data, a first step is to compute summary statistics for the data in question. Perhaps the most common summary statistics are the mean and standard deviation, which allow you to summarize the "typical" values in a dataset, but other aggregates are useful as well (the sum, product, median, minimum and maximum, quantiles, etc.).

NumPy has fast built-in aggregation functions for working on arrays; we'll discuss and demonstrate some of them here.

## Summing the Values in an Array

As a quick example, consider computing the sum of all values in an array. Python itself can do this using the built-in `sum` function:

In [1]:

```
import numpy as np
```

In [2]:

```
L = np.random.random(100)
sum(L)
```

Out[2]:

55.61209116604941

The syntax is quite similar to that of NumPy's `sum` function, and the result is the same in the simplest case:

In [3]:

```
np.sum(L)
```

Out[3]:

55.612091166049424

However, because it executes the operation in compiled code, NumPy's version of the operation is computed much more quickly:

In [4]:

```
big_array = np.random.rand(1000000)
%timeit sum(big_array)
```



```
%timeit np.sum(big_array)
10 loops, best of 3: 104 ms per loop
1000 loops, best of 3: 442 µs per loop
```

Be careful, though: the `sum` function and the `np.sum` function are not identical, which can sometimes lead to confusion! In particular, their optional arguments have different meanings, and `np.sum` is aware of multiple array dimensions, as we will see in the following section.

## Minimum and Maximum

Similarly, Python has built-in `min` and `max` functions, used to find the minimum value and maximum value of any given array:

In [5]:

```
min(big_array), max(big_array)
```

Out[5]:

```
(1.1717128136634614e-06, 0.9999976784968716)
```

NumPy's corresponding functions have similar syntax, and again operate much more quickly:

In [6]:

```
np.min(big_array), np.max(big_array)
```

Out[6]:

```
(1.1717128136634614e-06, 0.9999976784968716)
```

In [7]:

```
%timeit min(big_array)
%timeit np.min(big_array)
10 loops, best of 3: 82.3 ms per loop
1000 loops, best of 3: 497 µs per loop
```

For `min`, `max`, `sum`, and several other NumPy aggregates, a shorter syntax is to use methods of the array object itself:

In [8]:

```
print(big_array.min(), big_array.max(), big_array.sum())
1.17171281366e-06 0.999997678497 499911.628197
```

Whenever possible, make sure that you are using the NumPy version of these aggregates when operating on NumPy arrays!

## Multi dimensional aggregates

One common type of aggregation operation is an aggregate along a row or column. Say you have some data stored in a two-dimensional array:

In [9]:

```
M = np.random.random((3, 4))
print(M)
[[ 0.8967576  0.03783739  0.75952519  0.06682827]
 [ 0.8354065  0.99196818  0.19544769  0.43447084]
 [ 0.66859307  0.15038721  0.37911423  0.6687194 ]]
```

By default, each NumPy aggregation function will return the aggregate over the entire array:

In [10]:

```
M.sum()
Out[10]:
```

```
6.0850555667307118
```

Aggregation functions take an additional argument specifying the *axis* along which the aggregate is computed. For example, we can find the minimum value within each column by specifying `axis=0`:

In [11]:

```
M.min(axis=0)
Out[11]:
array([ 0.66859307,  0.03783739,  0.19544769,  0.06682827])
```

The function returns four values, corresponding to the four columns of numbers.

Similarly, we can find the maximum value within each row:

In [12]:

```
M.max(axis=1)
Out[12]:
array([ 0.8967576 ,  0.99196818,  0.6687194 ])
```

The way the axis is specified here can be confusing to users coming from other languages. The `axis` keyword specifies the *dimension of the array that will be collapsed*, rather than the dimension that will be returned. So specifying `axis=0` means that the first axis will be collapsed: for two-dimensional arrays, this means that values within each column will be aggregated.

## Other aggregation functions

NumPy provides many other aggregation functions, but we won't discuss them in detail here. Additionally, most aggregates have a NaN-safe counterpart that computes the result while ignoring missing values, which are marked by the special IEEE floating-point NaN value (for a fuller discussion of missing data, see [Handling Missing Data](#)). Some of these NaN-safe functions were not added until NumPy 1.8, so they will not be available in older NumPy versions.

The following table provides a list of useful aggregation functions available in NumPy:

Function Name	NaN-safe	Version	Description
---------------	----------	---------	-------------

<code>np.sum</code>	<code>np.nansum</code>		Compute sum of elements
<code>np.prod</code>	<code>np.nanprod</code>		Compute product of elements
<code>np.mean</code>	<code>np.nanmean</code>		Compute mean of elements
<code>np.std</code>	<code>np.nanstd</code>		Compute standard deviation
<code>np.var</code>	<code>np.nanvar</code>		Compute variance
<code>np.min</code>	<code>np.nanmin</code>		Find minimum value
<code>np.max</code>	<code>np.nanmax</code>		Find maximum value
<code>np.argmin</code>	<code>np.nanargmin</code>		Find index of minimum value
<code>np.argmax</code>	<code>np.nanargmax</code>		Find index of maximum value
<code>np.median</code>	<code>np.nanmedian</code>		Compute median of elements
<code>np.percentile</code>	<code>np.nanpercentile</code>		Compute rank-based statistics of elements
<code>np.any</code>	N/A		Evaluate whether any elements are true
<code>np.all</code>	N/A		Evaluate whether all elements are true

We will see these aggregates often throughout the rest of the book.

# Example: What is the Average Height of US Presidents?

Aggregates available in NumPy can be extremely useful for summarizing a set of values. As a simple example, let's consider the heights of all US presidents. This data is available in the file *president\_heights.csv*, which is a simple comma-separated list of labels and values:

In [13]:

```
!head -4 data/president_heights.csv
order,name,height(cm)
1,George Washington,189
2,John Adams,170
3,Thomas Jefferson,189
```

We'll use the Pandas package, which we'll explore more fully in [Chapter 3](#), to read the file and extract this information (note that the heights are measured in centimeters).

In [14]:

```
import pandas as pd
data = pd.read_csv('data/president_heights.csv')
heights = np.array(data['height(cm)'])
print(heights)
[189 170 189 163 183 171 185 168 173 183 173 173 175 178 183 193 178 173
 174 183 183 168 170 178 182 180 183 178 182 188 175 179 183 193 182 183
 177 185 188 188 182 185]
```

Now that we have this data array, we can compute a variety of summary statistics:

In [15]:

```
print("Mean height:    ", heights.mean())
print("Standard deviation:", heights.std())
print("Minimum height:  ", heights.min())
print("Maximum height:  ", heights.max())
Mean height:    179.738095238
Standard deviation: 6.93184344275
Minimum height:   163
Maximum height:   193
```

Note that in each case, the aggregation operation reduced the entire array to a single summarizing value, which gives us information about the distribution of values. We may also wish to compute quantiles:

In [16]:

```
print("25th percentile: ", np.percentile(heights, 25))
print("Median:           ", np.median(heights))
print("75th percentile:  ", np.percentile(heights, 75))
```

```
25th percentile: 174.25
Median:          182.0
75th percentile: 183.0
```

We see that the median height of US presidents is 182 cm, or just shy of six feet.

Of course, sometimes it's more useful to see a visual representation of this data, which we can accomplish using tools in Matplotlib (we'll discuss Matplotlib more fully in [Chapter 4](#)). For example, this code generates the following chart:

In [17]:

```
%matplotlib inline
import matplotlib.pyplot as plt
import seaborn; seaborn.set() # set plot style
```

In [18]:

```
plt.hist(heights)
plt.title('Height Distribution of US Presidents')
plt.xlabel('height (cm)')
plt.ylabel('number');
```

These aggregates are some of the fundamental pieces of exploratory data analysis that we'll explore in more depth in later chapters of the book.

## Computation on Arrays: Broadcasting

We saw in the previous section how NumPy's universal functions can be used to *vectorize* operations and thereby remove slow Python loops. Another means of vectorizing operations is to use NumPy's *broadcasting* functionality. Broadcasting is simply a set of rules for applying binary ufuncs (e.g., addition, subtraction, multiplication, etc.) on arrays of different sizes.

## Introducing Broadcasting

Recall that for arrays of the same size, binary operations are performed on an element-by-element basis:

In [1]:

```
import numpy as np
```

In [2]:

```
a = np.array([0, 1, 2])
b = np.array([5, 5, 5])
```

```
a + b
```

```
Out[2]:
```

```
array([5, 6, 7])
```

Broadcasting allows these types of binary operations to be performed on arrays of different sizes—for example, we can just as easily add a scalar (think of it as a zero-dimensional array) to an array:

```
In [3]:
```

```
a + 5
```

```
Out[3]:
```

```
array([5, 6, 7])
```

We can think of this as an operation that stretches or duplicates the value 5 into the array [5, 5, 5], and adds the results. The advantage of NumPy's broadcasting is that this duplication of values does not actually take place, but it is a useful mental model as we think about broadcasting.

We can similarly extend this to arrays of higher dimension. Observe the result when we add a one-dimensional array to a two-dimensional array:

```
In [4]:
```

```
M = np.ones((3, 3))
```

```
M
```

```
Out[4]:
```

```
array([[ 1.,  1.,  1.],
       [ 1.,  1.,  1.],
       [ 1.,  1.,  1.]])
```

```
In [5]:
```

```
M + a
```

```
Out[5]:
```

```
array([[ 1.,  2.,  3.],
       [ 1.,  2.,  3.],
       [ 1.,  2.,  3.]])
```

Here the one-dimensional array `a` is stretched, or broadcast across the second dimension in order to match the shape of `M`.

While these examples are relatively easy to understand, more complicated cases can involve broadcasting of both arrays. Consider the following example:

```
In [6]:
```

```
a = np.arange(3)
```

```
b = np.arange(3)[: , np.newaxis]
```

```
print(a)
print(b)
[0 1 2]
[[0]
 [1]
 [2]]
```

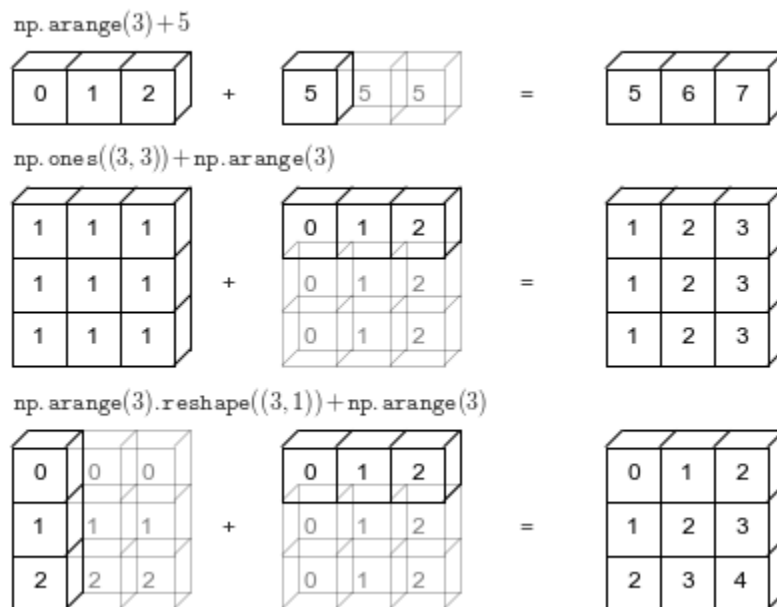
In [7]:

a + b

Out[7]:

```
array([[0, 1, 2],
       [1, 2, 3],
       [2, 3, 4]])
```

Just as before we stretched or broadcasted one value to match the shape of the other, here we've stretched *both* `a` and `b` to match a common shape, and the result is a two-dimensional array! The geometry of these examples is visualized in the following figure (Code to produce this plot can be found in the [appendix](#), and is adapted from source published in the [astroML](#) documentation. Used by permission).



The light boxes represent the broadcasted values: again, this extra memory is not actually allocated in the course of the operation, but it can be useful conceptually to imagine that it is.

## Rules of Broadcasting

Broadcasting in NumPy follows a strict set of rules to determine the interaction between the two arrays:

- Rule 1: If the two arrays differ in their number of dimensions, the shape of the one with fewer dimensions is *padded* with ones on its leading (left) side.
- Rule 2: If the shape of the two arrays does not match in any dimension, the array with shape equal to 1 in that dimension is stretched to match the other shape.
- Rule 3: If in any dimension the sizes disagree and neither is equal to 1, an error is raised.

To make these rules clear, let's consider a few examples in detail.

## Broadcasting example 1

Let's look at adding a two-dimensional array to a one-dimensional array:

In [8]:

```
M = np.ones((2, 3))
a = np.arange(3)
```

Let's consider an operation on these two arrays. The shape of the arrays are

- `M.shape = (2, 3)`
- `a.shape = (3,)`

We see by rule 1 that the array `a` has fewer dimensions, so we pad it on the left with ones:

- `M.shape -> (2, 3)`
- `a.shape -> (1, 3)`

By rule 2, we now see that the first dimension disagrees, so we stretch this dimension to match:

- `M.shape -> (2, 3)`
- `a.shape -> (2, 3)`

The shapes match, and we see that the final shape will be `(2, 3)`:

In [9]:

```
M + a
```



Out[9]:

```
array([[ 1.,  2.,  3.],  
       [ 1.,  2.,  3.]])
```

## Broadcasting example 2

Let's take a look at an example where both arrays need to be broadcast:

In [10]:

```
a = np.arange(3).reshape((3, 1))  
b = np.arange(3)
```

Again, we'll start by writing out the shape of the arrays:

- `a.shape = (3, 1)`
- `b.shape = (3,)`

Rule 1 says we must pad the shape of `b` with ones:

- `a.shape -> (3, 1)`
- `b.shape -> (1, 3)`

And rule 2 tells us that we upgrade each of these ones to match the corresponding size of the other array:

- `a.shape -> (3, 3)`
- `b.shape -> (3, 3)`

Because the result matches, these shapes are compatible. We can see this here:

In [11]:

```
a + b
```

Out[11]:

```
array([[0, 1, 2],  
       [1, 2, 3],  
       [2, 3, 4]])
```

## Broadcasting example 3

Now let's take a look at an example in which the two arrays are not compatible:

In [12]:

```
M = np.ones((3, 2))  
a = np.arange(3)
```

This is just a slightly different situation than in the first example: the matrix `M` is transposed. How does this affect the calculation? The shape of the arrays are

- `M.shape = (3, 2)`
- `a.shape = (3,)`

Again, rule 1 tells us that we must pad the shape of `a` with ones:

- `M.shape -> (3, 2)`
- `a.shape -> (1, 3)`

By rule 2, the first dimension of `a` is stretched to match that of `M`:

- `M.shape -> (3, 2)`
- `a.shape -> (3, 3)`

Now we hit rule 3—the final shapes do not match, so these two arrays are incompatible, as we can observe by attempting this operation:

In [13]:

```
M + a
```

```
-----  
ValueError                                Traceback (most recent call last)  
<ipython-input-13-9e16e9f98da6> in <module>()  
----> 1 M + a
```

**ValueError:** operands could not be broadcast together with shapes (3,2) (3,)

Note the potential confusion here: you could imagine making `a` and `M` compatible by, say, padding `a`'s shape with ones on the right rather than the left. But this is not how the broadcasting rules work! That sort of flexibility might be useful in some cases, but it would lead to potential areas of ambiguity. If right-side padding is what you'd like, you can do this explicitly by reshaping the array (we'll use the `np.newaxis` keyword introduced in [The Basics of NumPy Arrays](#)):

In [14]:

```
a[:, np.newaxis].shape
```

Out[14]:

```
(3, 1)
```

```
In [15]:
```

```
M + a[:, np.newaxis]
```

```
Out[15]:
```

```
array([[ 1.,  1.],  
       [ 2.,  2.],  
       [ 3.,  3.]])
```

Also note that while we've been focusing on the `+` operator here, these broadcasting rules apply to *any* binary ufunc. For example, here is the `logaddexp(a, b)` function, which computes `log(exp(a) + exp(b))` with more precision than the naive approach:

```
In [16]:
```

```
np.logaddexp(M, a[:, np.newaxis])
```

```
Out[16]:
```

```
array([[ 1.31326169,  1.31326169],  
       [ 1.69314718,  1.69314718],  
       [ 2.31326169,  2.31326169]])
```

## Broadcasting in Practice

Broadcasting operations form the core of many examples we'll see throughout this book. We'll now take a look at a couple simple examples of where they can be useful.

### Centering an array

In the previous section, we saw that ufuncs allow a NumPy user to remove the need to explicitly write slow Python loops. Broadcasting extends this ability. One commonly seen example is when centering an array of data. Imagine you have an array of 10 observations, each of which consists of 3 values. Using the standard convention, we'll store this in a  $10 \times 3$  array:

```
In [17]:
```

```
X = np.random.random((10, 3))
```

We can compute the mean of each feature using the `mean` aggregate across the first dimension:

```
In [18]:
```

```
Xmean = X.mean(0)  
Xmean
```

```
Out[18]:
```

```
array([ 0.53514715,  0.66567217,  0.44385899])
```

And now we can center the `X` array by subtracting the mean (this is a broadcasting operation):

In [19]:

```
X_centered = X - Xmean
```

To double-check that we've done this correctly, we can check that the centered array has near zero mean:

In [20]:

```
X_centered.mean(0)
```

Out[20]:

```
array([ 2.22044605e-17, -7.77156117e-17, -1.66533454e-17])
```

To within machine precision, the mean is now zero.

## Plotting a two-dimensional function

One place that broadcasting is very useful is in displaying images based on two-dimensional functions. If we want to define a function  $z=f(x,y)$ , broadcasting can be used to compute the function across the grid:

In [21]:

```
# x and y have 50 steps from 0 to 5
```

```
x = np.linspace(0, 5, 50)
```

```
y = np.linspace(0, 5, 50)[: , np.newaxis]
```

```
z = np.sin(x) ** 10 + np.cos(10 + y * x) * np.cos(x)
```

In [22]:

```
%matplotlib inline
```

```
import matplotlib.pyplot as plt
```

In [23]:

```
plt.imshow(z, origin='lower', extent=[0, 5, 0, 5],
```

```
        cmap='viridis')
```

```
plt.colorbar();
```

The result is a compelling visualization of the two-dimensional function.

# Comparisons, Masks, and Boolean Logic

This section covers the use of Boolean masks to examine and manipulate values within NumPy arrays. Masking comes up when you want to extract, modify, count, or otherwise manipulate values in an array based on some criterion: for example, you might wish to count all values greater than a certain value, or perhaps remove all outliers that are above some threshold. In NumPy, Boolean masking is often the most efficient way to accomplish these types of tasks.

## Example: Counting Rainy Days

Imagine you have a series of data that represents the amount of precipitation each day for a year in a given city. For example, here we'll load the daily rainfall statistics for the city of Seattle in 2014, using Pandas.

In [1]:

```
import numpy as np
import pandas as pd

# use pandas to extract rainfall inches as a NumPy array
rainfall = pd.read_csv('data/Seattle2014.csv')['PRCP'].values
inches = rainfall / 254.0 # 1/10mm -> inches
inches.shape
```

Out[1]:

(365,)

The array contains 365 values, giving daily rainfall in inches from January 1 to December 31, 2014.

As a first quick visualization, let's look at the histogram of rainy days, which was generated using Matplotlib (we will explore this tool more fully in [Chapter 4](#)):

In [2]:

```
%matplotlib inline
import matplotlib.pyplot as plt
import seaborn; seaborn.set() # set plot styles
```

In [3]:

```
plt.hist(inches, 40);
```

This histogram gives us a general idea of what the data looks like: despite its reputation, the vast majority of days in Seattle saw near zero measured rainfall in 2014. But this doesn't do a good job of conveying some information we'd like to see: for example, how many rainy days were there in the year? What is the average precipitation on those rainy days? How many days were there with more than half an inch of rain?

## Digging into the data

One approach to this would be to answer these questions by hand: loop through the data, incrementing a counter each time we see values in some desired range. For reasons discussed throughout this chapter, such an approach is very inefficient, both from the standpoint of time writing code and time computing the result. We saw in [Computation on NumPy Arrays: Universal Functions](#) that NumPy's ufuncs can be used in place of loops to do fast element-wise arithmetic operations on arrays; in the same way, we can use other ufuncs to do element-wise *comparisons* over arrays, and we can then manipulate the results to answer the questions we have. We'll leave the data aside for right now, and discuss some general tools in NumPy to use *masking* to quickly answer these types of questions.

## Comparison Operators as ufuncs

In [Computation on NumPy Arrays: Universal Functions](#) we introduced ufuncs, and focused in particular on arithmetic operators. We saw that using `+`, `-`, `*`, `/`, and others on arrays leads to element-wise operations. NumPy also implements comparison operators such as `<` (less than) and `>` (greater than) as element-wise ufuncs. The result of these comparison operators is always an array with a Boolean data type. All six of the standard comparison operations are available:

In [4]:

```
x = np.array([1, 2, 3, 4, 5])
```

In [5]:

```
x < 3 # less than
```

Out[5]:

```
array([ True,  True, False, False, False], dtype=bool)
```

In [6]:

```
x > 3 # greater than
```

Out[6]:

```
array([False, False, False,  True,  True], dtype=bool)
```

In [7]:

```
x <= 3 # less than or equal
```

Out[7]:

```
array([ True,  True,  True, False, False], dtype=bool)
```

In [8]:

```
x >= 3 # greater than or equal
```

Out[8]:

```
array([False, False,  True,  True,  True], dtype=bool)
```

In [9]:

```
x != 3 # not equal
```

Out[9]:

```
array([ True,  True, False,  True,  True], dtype=bool)
```

In [10]:

```
x == 3 # equal
```

Out[10]:

```
array([False, False,  True, False, False], dtype=bool)
```

It is also possible to do an element-wise comparison of two arrays, and to include compound expressions:

In [11]:

```
(2 * x) == (x ** 2)
```

Out[11]:

```
array([False,  True, False, False, False], dtype=bool)
```

As in the case of arithmetic operators, the comparison operators are implemented as ufuncs in NumPy; for example, when you write `x < 3`, internally NumPy uses `np.less(x, 3)`. A summary of the comparison operators and their equivalent ufunc is shown here:

#### OperatorEquivalent ufuncOperatorEquivalent ufunc

<code>==</code>	<code>np.equal</code>	<code>!=</code>	<code>np.not_equal</code>
<code>&lt;</code>	<code>np.less</code>	<code>&lt;=</code>	<code>np.less_equal</code>
<code>&gt;</code>	<code>np.greater</code>	<code>&gt;=</code>	<code>np.greater_equal</code>

Just as in the case of arithmetic ufuncs, these will work on arrays of any size and shape. Here is a two-dimensional example:

In [12]:

```
rng = np.random.RandomState(0)
x = rng.randint(10, size=(3, 4))
x
```

Out[12]:

```
array([[5, 0, 3, 3],
       [7, 9, 3, 5],
       [2, 4, 7, 6]])
```

In [13]:

```
x < 6
```

Out[13]:

```
array([[ True,  True,  True,  True],
       [False, False,  True,  True],
       [ True,  True, False, False]], dtype=bool)
```

In each case, the result is a Boolean array, and NumPy provides a number of straightforward patterns for working with these Boolean results.

## Working with Boolean Arrays

Given a Boolean array, there are a host of useful operations you can do. We'll work with `x`, the two-dimensional array we created earlier.

In [14]:

```
print(x)
[[5 0 3 3]
 [7 9 3 5]
 [2 4 7 6]]
```

## Counting entries

To count the number of `True` entries in a Boolean array, `np.count_nonzero` is useful:

In [15]:



```
# how many values less than 6?
```

```
np.count_nonzero(x < 6)
```

```
Out[15]:
```

```
8
```

We see that there are eight array entries that are less than 6. Another way to get at this information is to use `np.sum`; in this case, `False` is interpreted as 0, and `True` is interpreted as 1:

```
In [16]:
```

```
np.sum(x < 6)
```

```
Out[16]:
```

```
8
```

The benefit of `sum()` is that like with other NumPy aggregation functions, this summation can be done along rows or columns as well:

```
In [17]:
```

```
# how many values less than 6 in each row?
```

```
np.sum(x < 6, axis=1)
```

```
Out[17]:
```

```
array([4, 2, 2])
```

This counts the number of values less than 6 in each row of the matrix.

If we're interested in quickly checking whether any or all the values are true, we can use (you guessed it) `np.any` or `np.all`:

```
In [18]:
```

```
# are there any values greater than 8?
```

```
np.any(x > 8)
```

```
Out[18]:
```

```
True
```

```
In [19]:
```

```
# are there any values less than zero?
```

```
np.any(x < 0)
```

```
Out[19]:
```

```
False
```

```
In [20]:
```

```
# are all values less than 10?
```

```
np.all(x < 10)
```

```
Out[20]:
```

```
True
```

In [21]:

```
# are all values equal to 6?
```

```
np.all(x == 6)
```

Out[21]:

```
False
```

`np.all` and `np.any` can be used along particular axes as well. For example:

In [22]:

```
# are all values in each row less than 8?
```

```
np.all(x < 8, axis=1)
```

Out[22]:

```
array([ True, False,  True], dtype=bool)
```

Here all the elements in the first and third rows are less than 8, while this is not the case for the second row.

Finally, a quick warning: as mentioned in *Aggregations: Min, Max, and Everything In Between*, Python has built-in `sum()`, `any()`, and `all()` functions. These have a different syntax than the NumPy versions, and in particular will fail or produce unintended results when used on multidimensional arrays. Be sure that you are using `np.sum()`, `np.any()`, and `np.all()` for these examples!

## Boolean operators

We've already seen how we might count, say, all days with rain less than four inches, or all days with rain greater than two inches. But what if we want to know about all days with rain less than four inches and greater than one inch? This is accomplished through Python's *bitwise logic operators*, `&`, `|`, `^`, and `~`. Like with the standard arithmetic operators, NumPy overloads these as ufuncs which work element-wise on (usually Boolean) arrays.

For example, we can address this sort of compound question as follows:

In [23]:

```
np.sum((inches > 0.5) & (inches < 1))
```

Out[23]:

So we see that there are 29 days with rainfall between 0.5 and 1.0 inches.

Note that the parentheses here are important—because of operator precedence rules, with parentheses removed this expression would be evaluated as follows, which results in an error:

```
inches > (0.5 & inches) < 1
```

Using the equivalence of  $A \text{ AND } B$  and  $\text{NOT} (\text{NOT } A \text{ OR NOT } B)$  (which you may remember if you've taken an introductory logic course), we can compute the same result in a different manner:

In [24]:

```
np.sum(~( (inches <= 0.5) | (inches >= 1) ))
```

Out[24]:

29

Combining comparison operators and Boolean operators on arrays can lead to a wide range of efficient logical operations.

The following table summarizes the bitwise Boolean operators and their equivalent ufuncs:

Operator	Equivalent ufunc	Operator	Equivalent ufunc
----------	------------------	----------	------------------

&	np.bitwise_and		np.bitwise_or
---	----------------	--	---------------

^	np.bitwise_xor	~	np.bitwise_not
---	----------------	---	----------------

Using these tools, we might start to answer the types of questions we have about our weather data.

Here are some examples of results we can compute when combining masking with aggregations:

In [25]:

```
print("Number days without rain: ", np.sum(inches == 0))
print("Number days with rain: ", np.sum(inches != 0))
print("Days with more than 0.5 inches:", np.sum(inches > 0.5))
print("Rainy days with < 0.2 inches :", np.sum((inches > 0) &
(inches < 0.2)))
Number days without rain: 215
Number days with rain: 150
Days with more than 0.5 inches: 37
Rainy days with < 0.2 inches : 75
```

# Boolean Arrays as Masks

In the preceding section we looked at aggregates computed directly on Boolean arrays. A more powerful pattern is to use Boolean arrays as masks, to select particular subsets of the data themselves. Returning to our `x` array from before, suppose we want an array of all values in the array that are less than, say, 5:

In [26]:

```
x
```

Out[26]:

```
array([[5, 0, 3, 3],
       [7, 9, 3, 5],
       [2, 4, 7, 6]])
```

We can obtain a Boolean array for this condition easily, as we've already seen:

In [27]:

```
x < 5
```

Out[27]:

```
array([[False,  True,  True,  True],
       [False, False,  True, False],
       [ True,  True, False, False]], dtype=bool)
```

Now to *select* these values from the array, we can simply index on this Boolean array; this is known as a *masking* operation:

In [28]:

```
x[x < 5]
```

Out[28]:

```
array([0, 3, 3, 3, 2, 4])
```

What is returned is a one-dimensional array filled with all the values that meet this condition; in other words, all the values in positions at which the mask array is `True`.

We are then free to operate on these values as we wish. For example, we can compute some relevant statistics on our Seattle rain data:

In [29]:

```
# construct a mask of all rainy days
rainy = (inches > 0)
```

```
# construct a mask of all summer days (June 21st is the 172nd day)
```

```

days = np.arange(365)
summer = (days > 172) & (days < 262)

print("Median precip on rainy days in 2014 (inches): ",
      np.median(inches[rainy]))
print("Median precip on summer days in 2014 (inches): ",
      np.median(inches[summer]))
print("Maximum precip on summer days in 2014 (inches): ",
      np.max(inches[summer]))
print("Median precip on non-summer rainy days (inches):",
      np.median(inches[rainy & ~summer]))
Median precip on rainy days in 2014 (inches): 0.194881889764
Median precip on summer days in 2014 (inches): 0.0
Maximum precip on summer days in 2014 (inches): 0.850393700787
Median precip on non-summer rainy days (inches): 0.200787401575

```

By combining Boolean operations, masking operations, and aggregates, we can very quickly answer these sorts of questions for our dataset.

## Aside: Using the Keywords and/or Versus the Operators &/|

One common point of confusion is the difference between the keywords `and` and `or` on one hand, and the operators `&` and `|` on the other hand. When would you use one versus the other?

The difference is this: `and` and `or` gauge the truth or falsehood of *entire object*, while `&` and `|` refer to *bits within each object*.

When you use `and` or `or`, it's equivalent to asking Python to treat the object as a single Boolean entity. In Python, all nonzero integers will evaluate as True. Thus:

In [30]:

```
bool(42), bool(0)
```

Out[30]:

```
(True, False)
```

In [31]:

```
bool(42 and 0)
```

Out[31]:

```
False
```

In [32]:

```
bool(42 or 0)
```

Out[32]:

True

When you use `&` and `|` on integers, the expression operates on the bits of the element, applying the *and* or the *or* to the individual bits making up the number:

In [33]:

```
bin(42)
```

Out[33]:

```
'0b101010'
```

In [34]:

```
bin(59)
```

Out[34]:

```
'0b111011'
```

In [35]:

```
bin(42 & 59)
```

Out[35]:

```
'0b101010'
```

In [36]:

```
bin(42 | 59)
```

Out[36]:

```
'0b111011'
```

Notice that the corresponding bits of the binary representation are compared in order to yield the result.

When you have an array of Boolean values in NumPy, this can be thought of as a string of bits where `1 = True` and `0 = False`, and the result of `&` and `|` operates similarly to above:

In [37]:

```
A = np.array([1, 0, 1, 0, 1, 0], dtype=bool)
```

```
B = np.array([1, 1, 1, 0, 1, 1], dtype=bool)
```

```
A | B
```

Out[37]:

```
array([ True,  True,  True, False,  True,  True], dtype=bool)
```

Using `or` on these arrays will try to evaluate the truth or falsehood of the entire array object, which is not a well-defined value:

In [38]:

```
A or B
```

-----  
ValueError

Traceback (most recent call last)

```
<ipython-input-38-5d8e4f2e21c0> in <module>()
----> 1 A or B
```

**ValueError:** The truth value of an array with more than one element is ambiguous. Use a.any() or a.all()

Similarly, when doing a Boolean expression on a given array, you should use `|` or `&` rather than `or` or `and`:

In [39]:

```
x = np.arange(10)
(x > 4) & (x < 8)
```

Out[39]:

```
array([False, False, False, False, False,  True,  True,  True, False, False], dtype=bool)
```

Trying to evaluate the truth or falsehood of the entire array will give the same **ValueError** we saw previously:

In [40]:

```
(x > 4) and (x < 8)
```

```
-----
ValueError                                Traceback (most recent call last)
<ipython-input-40-3d24f1ffd63d> in <module>()
----> 1 (x > 4) and (x < 8)
```

**ValueError:** The truth value of an array with more than one element is ambiguous. Use a.any() or a.all()

So remember this: `and` and `or` perform a single Boolean evaluation on an entire object, while `&` and `|` perform multiple Boolean evaluations on the content (the individual bits or bytes) of an object. For Boolean NumPy arrays, the latter is nearly always the desired operation.

## Fancy Indexing

In the previous sections, we saw how to access and modify portions of arrays using simple indices (e.g., `arr[0]`), slices (e.g., `arr[:5]`), and Boolean masks (e.g., `arr[arr > 0]`). In this section, we'll look at another style of array indexing, known as *fancy indexing*. Fancy indexing is like the simple indexing we've already seen, but we pass arrays of indices in place of single scalars. This allows us to very quickly access and modify complicated subsets of an array's values.

## Exploring Fancy Indexing

Fancy indexing is conceptually simple: it means passing an array of indices to access multiple array elements at once. For example, consider the following array:

In [1]:

```
import numpy as np
rand = np.random.RandomState(42)

x = rand.randint(100, size=10)
print(x)
[51 92 14 71 60 20 82 86 74 74]
```

Suppose we want to access three different elements. We could do it like this:

In [2]:

```
[x[3], x[7], x[2]]
Out[2]:
[71, 86, 14]
```

Alternatively, we can pass a single list or array of indices to obtain the same result:

In [3]:

```
ind = [3, 7, 4]
x[ind]
Out[3]:
array([71, 86, 60])
```

When using fancy indexing, the shape of the result reflects the shape of the *index arrays* rather than the shape of the *array being indexed*:

In [4]:

```
ind = np.array([[3, 7],
                [4, 5]])
x[ind]
Out[4]:
array([[71, 86],
       [60, 20]])
```

Fancy indexing also works in multiple dimensions. Consider the following array:

In [5]:

```
X = np.arange(12).reshape((3, 4))
X
Out[5]:
array([[ 0,  1,  2,  3],
       [ 4,  5,  6,  7],
       [ 8,  9, 10, 11]])
```

Like with standard indexing, the first index refers to the row, and the second to the column:



In [6]:

```
row = np.array([0, 1, 2])
col = np.array([2, 1, 3])
X[row, col]
```

Out[6]:

```
array([ 2,  5, 11])
```

Notice that the first value in the result is `X[0, 2]`, the second is `X[1, 1]`, and the third is `X[2, 3]`. The pairing of indices in fancy indexing follows all the broadcasting rules that were mentioned in *Computation on Arrays: Broadcasting*. So, for example, if we combine a column vector and a row vector within the indices, we get a two-dimensional result:

In [7]:

```
X[row[:, np.newaxis], col]
```

Out[7]:

```
array([[ 2,  1,  3],
       [ 6,  5,  7],
       [10,  9, 11]])
```

Here, each row value is matched with each column vector, exactly as we saw in broadcasting of arithmetic operations. For example:

In [8]:

```
row[:, np.newaxis] * col
```

Out[8]:

```
array([[0, 0, 0],
       [2, 1, 3],
       [4, 2, 6]])
```

It is always important to remember with fancy indexing that the return value reflects the *broadcasted shape of the indices*, rather than the shape of the array being indexed.

## Combined Indexing

For even more powerful operations, fancy indexing can be combined with the other indexing schemes we've seen:

In [9]:

```
print(X)
[[ 0  1  2  3]
 [ 4  5  6  7]
 [ 8  9 10 11]]
```

We can combine fancy and simple indices:

In [10]:

```
X[2, [2, 0, 1]]
```

Out[10]:

```
array([10, 8, 9])
```

We can also combine fancy indexing with slicing:

In [11]:

```
X[1:, [2, 0, 1]]
```

Out[11]:

```
array([[ 6, 4, 5],  
       [10, 8, 9]])
```

And we can combine fancy indexing with masking:

In [12]:

```
mask = np.array([1, 0, 1, 0], dtype=bool)  
X[row[:, np.newaxis], mask]
```

Out[12]:

```
array([[ 0, 2],  
       [ 4, 6],  
       [ 8, 10]])
```

All of these indexing options combined lead to a very flexible set of operations for accessing and modifying array values.

## Example: Selecting Random Points

One common use of fancy indexing is the selection of subsets of rows from a matrix. For example, we might have an  $N \times D$  matrix representing  $N$  points in  $D$  dimensions, such as the following points drawn from a two-dimensional normal distribution:

In [13]:

```
mean = [0, 0]  
cov = [[1, 2],  
       [2, 5]]  
X = rand.multivariate_normal(mean, cov, 100)  
X.shape
```

Out[13]:

```
(100, 2)
```

Using the plotting tools we will discuss in [Introduction to Matplotlib](#), we can visualize these points as a scatter-plot:

In [14]:

```
%matplotlib inline
import matplotlib.pyplot as plt
import seaborn; seaborn.set() #for plot styling

plt.scatter(X[:, 0], X[:, 1]);
```

Let's use fancy indexing to select 20 random points. We'll do this by first choosing 20 random indices with no repeats, and use these indices to select a portion of the original array:

In [15]:

```
indices = np.random.choice(X.shape[0], 20, replace=False)
indices
```

Out[15]:

```
array([93, 45, 73, 81, 50, 10, 98, 94,  4, 64, 65, 89, 47, 84, 82, 80, 25,
       90, 63, 20])
```

In [16]:

```
selection = X[indices] #fancy indexing here
selection.shape
```

Out[16]:

```
(20, 2)
```

Now to see which points were selected, let's over-plot large circles at the locations of the selected points:

In [17]:

```
plt.scatter(X[:, 0], X[:, 1], alpha=0.3)
plt.scatter(selection[:, 0], selection[:, 1],
            facecolor='none', s=200);
```

This sort of strategy is often used to quickly partition datasets, as is often needed in train/test splitting for validation of statistical models (see [Hyperparameters and Model Validation](#)), and in sampling approaches to answering statistical questions.

## Modifying Values with Fancy Indexing

Just as fancy indexing can be used to access parts of an array, it can also be used to modify parts of an array. For example, imagine we have an array of indices and we'd like to set the corresponding items in an array to some value:

In [18]:

```
x = np.arange(10)
i = np.array([2, 1, 8, 4])
x[i] = 99
print(x)
[ 0 99 99  3 99  5  6  7 99  9]
```

We can use any assignment-type operator for this. For example:

In [19]:

```
x[i] -= 10
print(x)
[ 0 89 89  3 89  5  6  7 89  9]
```

Notice, though, that repeated indices with these operations can cause some potentially unexpected results. Consider the following:

In [20]:

```
x = np.zeros(10)
x[[0, 0]] = [4, 6]
print(x)
[ 6.  0.  0.  0.  0.  0.  0.  0.  0.  0.]
```

Where did the 4 go? The result of this operation is to first assign `x[0] = 4`, followed by `x[0] = 6`. The result, of course, is that `x[0]` contains the value 6.

Fair enough, but consider this operation:

In [21]:

```
i = [2, 3, 3, 4, 4, 4]
x[i] += 1
x
```

Out[21]:

```
array([ 6.,  0.,  1.,  1.,  1.,  0.,  0.,  0.,  0.,  0.])
```

You might expect that `x[3]` would contain the value 2, and `x[4]` would contain the value 3, as this is how many times each index is repeated. Why is this not the case? Conceptually, this is because `x[i] += 1` is meant as a shorthand of `x[i] = x[i] + 1`. `x[i] + 1` is evaluated, and then the result is assigned to the indices in `x`. With this in mind, it is not the augmentation that happens multiple times, but the assignment, which leads to the rather nonintuitive results.

So what if you want the other behavior where the operation is repeated? For this, you can use the `at()` method of ufuncs (available since NumPy 1.8), and do the following:

In [22]:

```
x = np.zeros(10)
np.add.at(x, i, 1)
print(x)
[ 0.  0.  1.  2.  3.  0.  0.  0.  0.  0.]
```

The `at()` method does an in-place application of the given operator at the specified indices (here, `i`) with the specified value (here, 1). Another method that is similar in spirit is the `reduceat()` method of ufuncs, which you can read about in the NumPy documentation.

## Example: Binning Data

You can use these ideas to efficiently bin data to create a histogram by hand. For example, imagine we have 1,000 values and would like to quickly find where they fall within an array of bins. We could compute it using `ufunc.at` like this:

In [23]:

```
np.random.seed(42)
x = np.random.randn(100)

# compute a histogram by hand
bins = np.linspace(-5, 5, 20)
counts = np.zeros_like(bins)

# find the appropriate bin for each x
i = np.searchsorted(bins, x)

# add 1 to each of these bins
np.add.at(counts, i, 1)
```

The counts now reflect the number of points within each bin—in other words, a histogram:

In [24]:

```
# plot the results
plt.plot(bins, counts, linestyle='steps');
```

Of course, it would be silly to have to do this each time you want to plot a histogram. This is why Matplotlib provides the `plt.hist()` routine, which does the same in a single line:

```
plt.hist(x, bins, histtype='step');
```

This function will create a nearly identical plot to the one seen here. To compute the binning, `matplotlib` uses the `np.histogram` function, which does a very similar computation to what we did before. Let's compare the two here:

In [25]:

```
print("NumPy routine:")
%timeit counts, edges = np.histogram(x, bins)

print("Custom routine:")
%timeit np.add.at(counts, np.searchsorted(bins, x), 1)
NumPy routine:
10000 loops, best of 3: 97.6 µs per loop
Custom routine:
10000 loops, best of 3: 19.5 µs per loop
```

Our own one-line algorithm is several times faster than the optimized algorithm in NumPy! How can this be? If you dig into the `np.histogram` source code (you can do this in IPython by typing `np.histogram??`), you'll see that it's quite a bit more involved than the simple search-and-count that we've done; this is because NumPy's algorithm is more flexible, and particularly is designed for better performance when the number of data points becomes large:

In [26]:

```
x = np.random.randn(1000000)
print("NumPy routine:")
%timeit counts, edges = np.histogram(x, bins)

print("Custom routine:")
%timeit np.add.at(counts, np.searchsorted(bins, x), 1)
NumPy routine:
10 loops, best of 3: 68.7 ms per loop
Custom routine:
10 loops, best of 3: 135 ms per loop
```

What this comparison shows is that algorithmic efficiency is almost never a simple question. An algorithm efficient for large datasets will not always be the best choice for small datasets, and vice versa (see [Big-O Notation](#)). But the advantage of coding this algorithm yourself is that with an understanding of these basic methods, you could use these building blocks to extend this to do some very interesting custom behaviors. The key to efficiently using Python in data-intensive applications is knowing about general convenience routines like `np.histogram` and when they're appropriate, but also knowing how to make use of lower-level functionality when you need more pointed behavior.

# Sorting Arrays

Up to this point we have been concerned mainly with tools to access and operate on array data with NumPy. This section covers algorithms related to sorting values in NumPy arrays. These algorithms are a favorite topic in introductory computer science courses: if you've ever taken one, you probably have had dreams (or, depending on your temperament, nightmares) about *insertion sorts*, *selection sorts*, *merge sorts*, *quick sorts*, *bubble sorts*, and many, many more. All are means of accomplishing a similar task: sorting the values in a list or array.

For example, a simple *selection sort* repeatedly finds the minimum value from a list, and makes swaps until the list is sorted. We can code this in just a few lines of Python:

In [1]:

```
import numpy as np
```

```
def selection_sort(x):  
    for i in range(len(x)):  
        swap = i + np.argmin(x[i:])  
        (x[i], x[swap]) = (x[swap], x[i])  
    return x
```

In [2]:

```
x = np.array([2, 1, 4, 3, 5])  
selection_sort(x)
```

Out[2]:

```
array([1, 2, 3, 4, 5])
```

As any first-year computer science major will tell you, the selection sort is useful for its simplicity, but is much too slow to be useful for larger arrays. For a list of  $N$  values, it requires  $N$  loops, each of which does on order  $\sim N$  comparisons to find the swap value. In terms of the "big-O" notation often used to characterize these algorithms (see [Big-O Notation](#)), selection sort averages  $O[N^2]$ : if you double the number of items in the list, the execution time will go up by about a factor of four.

Even selection sort, though, is much better than my all-time favorite sorting algorithms, the *bogosort*:

In [3]:

```
def bogosort(x):  
    while np.any(x[:-1] > x[1:]):  
        np.random.shuffle(x)
```

```
return x
```

In [4]:

```
x = np.array([2, 1, 4, 3, 5])
bogosort(x)
```

Out[4]:

```
array([1, 2, 3, 4, 5])
```

This silly sorting method relies on pure chance: it repeatedly applies a random shuffling of the array until the result happens to be sorted. With an average scaling of  $O[N \times N!]O[N \times N!]$ , (that's  $N$  times  $N$  factorial) this should—quite obviously—never be used for any real computation.

Fortunately, Python contains built-in sorting algorithms that are *much* more efficient than either of the simplistic algorithms just shown. We'll start by looking at the Python built-ins, and then take a look at the routines included in NumPy and optimized for NumPy arrays.

## Fast Sorting in NumPy: `np.sort` and `np.argsort`

Although Python has built-in `sort` and `sorted` functions to work with lists, we won't discuss them here because NumPy's `np.sort` function turns out to be much more efficient and useful for our purposes. By default `np.sort` uses an  $O[N \log N]O[N \log^{[f_0]} N]$ , *quicksort* algorithm, though *mergesort* and *heapsort* are also available. For most applications, the default quicksort is more than sufficient.

To return a sorted version of the array without modifying the input, you can use `np.sort`:

In [5]:

```
x = np.array([2, 1, 4, 3, 5])
np.sort(x)
```

Out[5]:

```
array([1, 2, 3, 4, 5])
```

If you prefer to sort the array in-place, you can instead use the `sort` method of arrays:

In [6]:

```
x.sort()
print(x)
[1 2 3 4 5]
```

A related function is `argsort`, which instead returns the *indices* of the sorted elements:

In [7]:

```
x = np.array([2, 1, 4, 3, 5])
```



```
i = np.argsort(x)
print(i)
[1 0 3 2 4]
```

The first element of this result gives the index of the smallest element, the second value gives the index of the second smallest, and so on. These indices can then be used (via fancy indexing) to construct the sorted array if desired:

In [8]:

```
x[i]
Out[8]:
array([1, 2, 3, 4, 5])
```

## Sorting along rows or columns

A useful feature of NumPy's sorting algorithms is the ability to sort along specific rows or columns of a multidimensional array using the `axis` argument. For example:

In [9]:

```
rand = np.random.RandomState(42)
X = rand.randint(0, 10, (4, 6))
print(X)
[[6 3 7 4 6 9]
 [2 6 7 4 3 7]
 [7 2 5 4 1 7]
 [5 1 4 0 9 5]]
```

In [10]:

```
# sort each column of X
np.sort(X, axis=0)
```

Out[10]:

```
array([[2, 1, 4, 0, 1, 5],
       [5, 2, 5, 4, 3, 7],
       [6, 3, 7, 4, 6, 7],
       [7, 6, 7, 4, 9, 9]])
```

In [11]:

```
# sort each row of X
np.sort(X, axis=1)
```

Out[11]:

```
array([[3, 4, 6, 6, 7, 9],
       [2, 3, 4, 6, 7, 7],
       [1, 2, 4, 5, 7, 7],
       [0, 1, 4, 5, 5, 9]])
```

Keep in mind that this treats each row or column as an independent array, and any relationships between the row or column values will be lost!

# Partial Sorts: Partitioning

Sometimes we're not interested in sorting the entire array, but simply want to find the  $k$  smallest values in the array. NumPy provides this in the `np.partition` function. `np.partition` takes an array and a number  $K$ ; the result is a new array with the smallest  $K$  values to the left of the partition, and the remaining values to the right, in arbitrary order:

In [12]:

```
x = np.array([7, 2, 3, 1, 6, 5, 4])
np.partition(x, 3)
```

Out[12]:

```
array([2, 1, 3, 4, 6, 5, 7])
```

Note that the first three values in the resulting array are the three smallest in the array, and the remaining array positions contain the remaining values. Within the two partitions, the elements have arbitrary order.

Similarly to sorting, we can partition along an arbitrary axis of a multidimensional array:

In [13]:

```
np.partition(X, 2, axis=1)
```

Out[13]:

```
array([[3, 4, 6, 7, 6, 9],
       [2, 3, 4, 7, 6, 7],
       [1, 2, 4, 5, 7, 7],
       [0, 1, 4, 5, 9, 5]])
```

The result is an array where the first two slots in each row contain the smallest values from that row, with the remaining values filling the remaining slots.

Finally, just as there is a `np.argsort` that computes indices of the sort, there is a `np.argpartition` that computes indices of the partition. We'll see this in action in the following section.

## Example: k-Nearest Neighbors

Let's quickly see how we might use this `argsort` function along multiple axes to find the nearest neighbors of each point in a set. We'll start by creating a random set of 10 points on a two-dimensional plane. Using the standard convention, we'll arrange these in a  $10 \times 2$  array:

In [14]:

```
X = rand.rand(10, 2)
```

To get an idea of how these points look, let's quickly scatter plot them:

In [15]:

```
%matplotlib inline
import matplotlib.pyplot as plt
import seaborn; seaborn.set() # Plot styling
plt.scatter(X[:, 0], X[:, 1], s=100);
```

Now we'll compute the distance between each pair of points. Recall that the squared-distance between two points is the sum of the squared differences in each dimension; using the efficient broadcasting (Computation on Arrays: Broadcasting) and aggregation (Aggregations: Min, Max, and Everything In Between) routines provided by NumPy we can compute the matrix of square distances in a single line of code:

In [16]:

```
dist_sq = np.sum((X[:, np.newaxis, :] - X[np.newaxis, :, :]) ** 2, axis=-1)
```

This operation has a lot packed into it, and it might be a bit confusing if you're unfamiliar with NumPy's broadcasting rules. When you come across code like this, it can be useful to break it down into its component steps:

In [17]:

```
# for each pair of points, compute differences in their coordinates
differences = X[:, np.newaxis, :] - X[np.newaxis, :, :]
differences.shape
```

Out[17]:

```
(10, 10, 2)
```

In [18]:

```
# square the coordinate differences
sq_differences = differences ** 2
sq_differences.shape
```

Out[18]:

```
(10, 10, 2)
```

In [19]:

```
# sum the coordinate differences to get the squared distance
dist_sq = sq_differences.sum(-1)
dist_sq.shape
```

Out[19]:

```
(10, 10)
```

Just to double-check what we are doing, we should see that the diagonal of this matrix (i.e., the set of distances between each point and itself) is all zero:

In [20]:

```
dist_sq.diagonal()
```

Out[20]:

```
array([ 0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.])
```

It checks out! With the pairwise square-distances converted, we can now use `np.argsort` to sort along each row. The leftmost columns will then give the indices of the nearest neighbors:

In [21]:

```
nearest = np.argsort(dist_sq, axis=1)
```

```
print(nearest)
```

```
[[0 3 9 7 1 4 2 5 6 8]
 [1 4 7 9 3 6 8 5 0 2]
 [2 1 4 6 3 0 8 9 7 5]
 [3 9 7 0 1 4 5 8 6 2]
 [4 1 8 5 6 7 9 3 0 2]
 [5 8 6 4 1 7 9 3 2 0]
 [6 8 5 4 1 7 9 3 2 0]
 [7 9 3 1 4 0 5 8 6 2]
 [8 5 6 4 1 7 9 3 2 0]
 [9 7 3 0 1 4 5 8 6 2]]
```

Notice that the first column gives the numbers 0 through 9 in order: this is due to the fact that each point's closest neighbor is itself, as we would expect.

By using a full sort here, we've actually done more work than we need to in this case. If we're simply interested in the nearest  $k$  neighbors, all we need is to partition each row so that the smallest  $k+1$  squared distances come first, with larger distances filling the remaining positions of the array. We can do this with the `np.argpartition` function:

In [22]:

```
K = 2
```

```
nearest_partition = np.argpartition(dist_sq, K + 1, axis=1)
```

In order to visualize this network of neighbors, let's quickly plot the points along with lines representing the connections from each point to its two nearest neighbors:

In [23]:

```
plt.scatter(X[:, 0], X[:, 1], s=100)
```

```
# draw lines from each point to its two nearest neighbors
```

```
K = 2
```

```

for i in range(X.shape[0]):
    for j in nearest_partition[i, :K+1]:
        # plot a line from X[i] to X[j]
        # use some zip magic to make it happen:
        plt.plot(*zip(X[j], X[i]), color='black')

```

Each point in the plot has lines drawn to its two nearest neighbors. At first glance, it might seem strange that some of the points have more than two lines coming out of them: this is due to the fact that if point A is one of the two nearest neighbors of point B, this does not necessarily imply that point B is one of the two nearest neighbors of point A.

Although the broadcasting and row-wise sorting of this approach might seem less straightforward than writing a loop, it turns out to be a very efficient way of operating on this data in Python. You might be tempted to do the same type of operation by manually looping through the data and sorting each set of neighbors individually, but this would almost certainly lead to a slower algorithm than the vectorized version we used. The beauty of this approach is that it's written in a way that's agnostic to the size of the input data: we could just as easily compute the neighbors among 100 or 1,000,000 points in any number of dimensions, and the code would look the same.

Finally, I'll note that when doing very large nearest neighbor searches, there are tree-based and/or approximate algorithms that can scale as  $O[N \log N]$  or better rather than the  $O[N^2]$  of the brute-force algorithm. One example of this is the KD-Tree, implemented in Scikit-learn.

## Aside: Big-O Notation

Big-O notation is a means of describing how the number of operations required for an algorithm scales as the input grows in size. To use it correctly is to dive deeply into the realm of computer science theory, and to carefully distinguish it from the related small-o notation, big- $\Theta$  notation, big- $\Omega$  notation, and probably many mutant hybrids thereof. While these distinctions add precision to statements about algorithmic scaling, outside computer science theory exams and the remarks of pedantic blog commenters, you'll rarely see such distinctions made in practice. Far more common in the data science world is a less rigid use of big-O notation: as a general (if imprecise) description of the scaling of an algorithm. With apologies to theorists and pedants, this is the interpretation we'll use throughout this book.

Big-O notation, in this loose sense, tells you how much time your algorithm will take as you increase the amount of data. If you have an  $O[N]O[N]$  (read "order  $NN$ ") algorithm that takes 1 second to operate on a list of length  $N=1,000$ , then you should expect it to take roughly 5 seconds for a list of length  $N=5,000$ . If you have an  $O[N^2]O[N^2]$  (read "order  $N$  squared") algorithm that takes 1 second for  $N=1000$ , then you should expect it to take about 25 seconds for  $N=5000$ .

For our purposes, the  $N$  will usually indicate some aspect of the size of the dataset (the number of points, the number of dimensions, etc.). When trying to analyze billions or trillions of samples, the difference between  $O[N]O[N]$  and  $O[N^2]O[N^2]$  can be far from trivial!

Notice that the big-O notation by itself tells you nothing about the actual wall-clock time of a computation, but only about its scaling as you change  $N$ . Generally, for example, an  $O[N]O[N]$  algorithm is considered to have better scaling than an  $O[N^2]O[N^2]$  algorithm, and for good reason. But for small datasets in particular, the algorithm with better scaling might not be faster. For example, in a given problem an  $O[N^2]O[N^2]$  algorithm might take 0.01 seconds, while a "better"  $O[N]O[N]$  algorithm might take 1 second. Scale up  $N$  by a factor of 1,000, though, and the  $O[N]O[N]$  algorithm will win out.

Even this loose version of Big-O notation can be very useful when comparing the performance of algorithms, and we'll use this notation throughout the book when talking about how algorithms scale.

## Structured Data: NumPy's Structured Arrays

While often our data can be well represented by a homogeneous array of values, sometimes this is not the case. This section demonstrates the use of NumPy's *structured arrays* and *record arrays*, which provide efficient storage for compound, heterogeneous data. While the patterns shown here are useful for simple operations, scenarios like this often lend themselves to the use of Pandas `Dataframes`.

In [1]:

```
import numpy as np
```

Imagine that we have several categories of data on a number of people (say, name, age, and weight), and we'd like to store these values for use in a Python program. It would be possible to store these in three separate arrays:

In [2]:

```
name = ['Alice', 'Bob', 'Cathy', 'Doug']
age = [25, 45, 37, 19]
weight = [55.0, 85.5, 68.0, 61.5]
```

But this is a bit clumsy. There's nothing here that tells us that the three arrays are related; it would be more natural if we could use a single structure to store all of this data. NumPy can handle this through structured arrays, which are arrays with compound data types.

Recall that previously we created a simple array using an expression like this:

In [3]:

```
x = np.zeros(4, dtype=int)
```

We can similarly create a structured array using a compound data type specification:

In [4]:

```
# Use a compound data type for structured arrays
data = np.zeros(4, dtype={'names':('name', 'age', 'weight'),
                          'formats':('U10', 'i4', 'f8')})
print(data.dtype)
[('name', '<U10'), ('age', '<i4'), ('weight', '<f8')]
```

Here 'U10' translates to "Unicode string of maximum length 10," 'i4' translates to "4-byte (i.e., 32 bit) integer," and 'f8' translates to "8-byte (i.e., 64 bit) float." We'll discuss other options for these type codes in the following section.

Now that we've created an empty container array, we can fill the array with our lists of values:

In [5]:

```
data['name'] = name
data['age'] = age
data['weight'] = weight
print(data)
[('Alice', 25, 55.0) ('Bob', 45, 85.5) ('Cathy', 37, 68.0)
 ('Doug', 19, 61.5)]
```

As we had hoped, the data is now arranged together in one convenient block of memory.

The handy thing with structured arrays is that you can now refer to values either by index or by name:

In [6]:

```
# Get all names
data['name']
```

Out[6]:

```
array(['Alice', 'Bob', 'Cathy', 'Doug'],  
      dtype='<U10')
```

In [7]:

```
# Get first row of data  
data[0]
```

Out[7]:

```
('Alice', 25, 55.0)
```

In [8]:

```
# Get the name from the last row  
data[-1]['name']
```

Out[8]:

```
'Doug'
```

Using Boolean masking, this even allows you to do some more sophisticated operations such as filtering on age:

In [9]:

```
# Get names where age is under 30  
data[data['age'] < 30]['name']
```

Out[9]:

```
array(['Alice', 'Doug'],  
      dtype='<U10')
```

Note that if you'd like to do any operations that are any more complicated than these, you should probably consider the Pandas package, covered in the next chapter. As we'll see, Pandas provides a `Dataframe` object, which is a structure built on NumPy arrays that offers a variety of useful data manipulation functionality similar to what we've shown here, as well as much, much more.

## Creating Structured Arrays

Structured array data types can be specified in a number of ways. Earlier, we saw the dictionary method:

In [10]:

```
np.dtype({'names':('name', 'age', 'weight'),  
         'formats':('U10', 'i4', 'f8')})
```

Out[10]:

```
dtype([('name', '<U10'), ('age', '<i4'), ('weight', '<f8')])
```

For clarity, numerical types can be specified using Python types or NumPy `dtypes` instead:



In [11]:

```
np.dtype({'names':('name', 'age', 'weight'),  
         'formats':((np.str_, 10), int, np.float32)})
```

Out[11]:

```
dtype([('name', '<U10'), ('age', '<i8'), ('weight', '<f4')])
```

A compound type can also be specified as a list of tuples:

In [12]:

```
np.dtype([('name', 'S10'), ('age', 'i4'), ('weight', 'f8')])
```

Out[12]:

```
dtype([('name', 'S10'), ('age', '<i4'), ('weight', '<f8')])
```

If the names of the types do not matter to you, you can specify the types alone in a comma-separated string:

In [13]:

```
np.dtype('S10,i4,f8')
```

Out[13]:

```
dtype([('f0', 'S10'), ('f1', '<i4'), ('f2', '<f8')])
```

The shortened string format codes may seem confusing, but they are built on simple principles. The first (optional) character is `<` or `>`, which means "little endian" or "big endian," respectively, and specifies the ordering convention for significant bits. The next character specifies the type of data: characters, bytes, ints, floating points, and so on (see the table below). The last character or characters represents the size of the object in bytes.

CharacterDescription		Example
'b'	Byte	<code>np.dtype('b')</code>
'i'	Signed integer	<code>np.dtype('i4') == np.int32</code>
'u'	Unsigned integer	<code>np.dtype('u1') == np.uint8</code>
'f'	Floating point	<code>np.dtype('f8') == np.int64</code>
'c'	Complex floating point	<code>np.dtype('c16') == np.complex128</code>

Character	Description	Example
'S', 'a'	String	<code>np.dtype('S5')</code>
'U'	Unicode string	<code>np.dtype('U') == np.str_</code>
'V'	Raw data (void)	<code>np.dtype('V') == np.void</code>

## More Advanced Compound Types

It is possible to define even more advanced compound types. For example, you can create a type where each element contains an array or matrix of values. Here, we'll create a data type with a `mat` component consisting of a  $3 \times 3 \times 3$  floating-point matrix:

In [14]:

```
tp = np.dtype([('id', 'i8'), ('mat', 'f8', (3, 3))])
X = np.zeros(1, dtype=tp)
print(X[0])
print(X['mat'][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.  0.]
 [ 0.  0.  0.]
```

Now each element in the `X` array consists of an `id` and a  $3 \times 3 \times 3$  matrix. Why would you use this rather than a simple multidimensional array, or perhaps a Python dictionary? The reason is that this NumPy `dtype` directly maps onto a C structure definition, so the buffer containing the array content can be accessed directly within an appropriately written C program. If you find yourself writing a Python interface to a legacy C or Fortran library that manipulates structured data, you'll probably find structured arrays quite useful!

## RecordArrays: Structured Arrays with a Twist

NumPy also provides the `np.recarray` class, which is almost identical to the structured arrays just described, but with one additional feature: fields can be accessed as attributes rather than as dictionary keys. Recall that we previously accessed the ages by writing:

In [15]:

```
data['age']
Out[15]:
```

```
array([25, 45, 37, 19], dtype=int32)
```

If we view our data as a record array instead, we can access this with slightly fewer keystrokes:

In [16]:

```
data_rec = data.view(np.recarray)
data_rec.age
```

Out[16]:

```
array([25, 45, 37, 19], dtype=int32)
```

The downside is that for record arrays, there is some extra overhead involved in accessing the fields, even when using the same syntax. We can see this here:

In [17]:

```
%timeit data['age']
%timeit data_rec['age']
%timeit data_rec.age
1000000 loops, best of 3: 241 ns per loop
100000 loops, best of 3: 4.61 µs per loop
100000 loops, best of 3: 7.27 µs per loop
```

Whether the more convenient notation is worth the additional overhead will depend on your own application.