DATA 572: Supervised Learning Tutorial - Introduction to Python

2023W2

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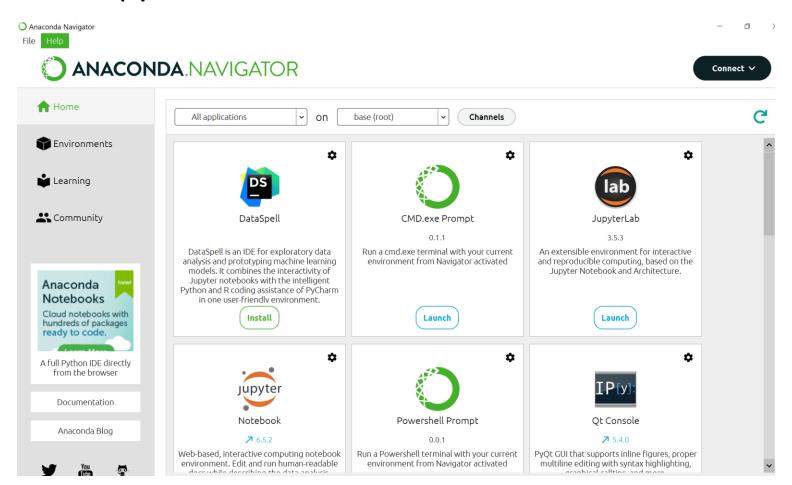
Installation

- To run the labs in this course, you will need two things:
 - 1. An installation of *Python3*, which is the specific version of Python used in the labs.
 - 2. Access to *Jupyter*, a very popular Python interface that runs code through a file called a *notebook*.

You can download and install Python3 by following the instructions available at anaconda.com.

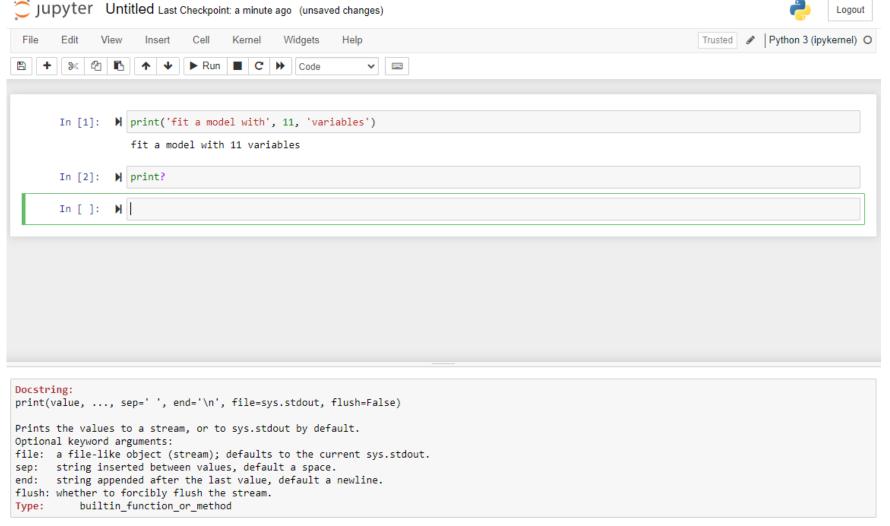
Installation

The Jupyter Notebook is also installed with Anaconda.



https://docs.python.org/3/tutorial/

Like most programming languages, Python uses functions to perform operations. To run a function called fun, we type fun(input1, input2), where the inputs (or arguments) input1 and input2 tell Python how to run the function. A function can have any number of inputs.



Adding two integers

```
In [3]: № 3 + 5
Out[3]: 8
```

 In Python, textual data is handled using strings. For instance, "hello" and 'hello' are strings.

- A string is actually a type of sequence: this is a generic term for an ordered list. The three most important types of sequences are lists, tuples, and strings.
- Example: Join together the numbers 3, 4, and 5, and to save them as a *list* named x.

```
In [7]: M x = [3, 4, 5]
x
Out[7]: [3, 4, 5]
```

Note that we used the brackets [] to construct this list.

```
In [7]: M = [3, 4, 5]

Out[7]: [3, 4, 5]

In [8]: M = [4, 9, 7]

X = [4, 9, 7]

X = [4, 9, 7]
```

• In Python, lists hold arbitrary objects, and are added using *concatenation*. Much of Python's data-specific functionality comes from other packages, notably *numpy* and *pandas*.

- A package is a collection of modules that are not necessarily included in the base Python distribution.
- The *numpy library*, or *package* is an abbreviation for *numerical Python*.
- To access numpy, we must first import it.

```
In [9]: ▶ import numpy as np
```

 In numpy, an array is a generic term for a multidimensional set of numbers. We use the np.array() function to define x and y, which are onedimensional arrays, i.e., vectors.

 The syntax np.array() indicates that the function being called is part of the numpy package, which we have abbreviated as np.

 In numpy, matrices are typically represented as two-dimensional arrays, and vectors as one-dimensional arrays. We can create a twodimensional array as follows.

 The object x has several attributes, or associated objects. To access an attribute of x, we type x.attribute, where we replace attribute with the name of the attribute. For instance, we can access the ndim attribute of x as follows.

 The output indicates that x is a twodimensional array. Similarly, x.dtype is the data type attribute of the object x. This indicates that x is comprised of 32-bit integers:

 This is because we created x by passing in exclusively integers to the np.array() function.

• If we had passed in any decimals, then we would have obtained an array of *floating point* numbers.

```
In [15]:  np.array([[1, 2], [3.0, 4]]).dtype
Out[15]: dtype('float64')
```

 We can create a floating point array by passing a dtype argument into np.array().dtype

• The array x is two-dimensional. We can find out the number of rows and columns by looking at its shape attribute.

In [18]: N x.shape

Out[18]: (2, 2)

A method is a function that is associated with an object. For instance, given an array method x, the expression x.sum() sums all of its elements, using the sum() method for arrays. The call x.sum() automatically provides x as the first argument to its sum() method.

 We could also sum the elements of x by passing in x as an argument to the np.sum() function.

• The *reshape()* method returns a new array with the same elements as *x*, but a different shape. We do this by passing in a tuple in our call to *reshape()*, in this case (2, 3). This tuple specifies that we would like to create a two-dimensional array with 2 rows and 3 columns.

- numpy arrays are specified as a sequence of rows. This is called row-major ordering, as opposed to column-major ordering.
- Python (and hence numpy) uses *O-based indexing*. This means that to access the top
 left element of x_reshape, we type in
 x_reshape[0,0].

```
In [22]: ► x_reshape[0, 0]
Out[22]: 1
```

 Modifying x_reshape also modified x because the two objects occupy the same space in memory.

```
In [23]:
          print('x before we modify x reshape:\n', x)
             print('x reshape before we modify x reshape:\n', x reshape)
             x reshape[0, 0] = 5
             print('x_reshape after we modify its top left element:\n',
             x reshape)
             print('x after we modify top left element of x_reshape:\n', x)
             x before we modify x reshape:
              [1 2 3 4 5 6]
             x reshape before we modify x reshape:
              [[1 2 3]
              [4 5 6]]
             x reshape after we modify its top left element:
              [[5 2 3]
              [4 5 6]]
             x after we modify top left element of x reshape:
              [5 2 3 4 5 6]
```

```
In [25]:  x_reshape.shape , x_reshape.ndim , x_reshape.T
    Out[25]: ((2, 3),
              array([5, 4],
                     [2, 5],
                     [3, 6]]))
In [26]:  ▶ np.sqrt(x)
   Out[26]: array([2.23606798, 1.41421356, 1.73205081, 2.
                                                                , 2.23606798,
                   2.44948974])
In [27]: ► x**2
   Out[27]: array([25, 4, 9, 16, 25, 36])
In [28]: ► x**0.5
   Out[28]: array([2.23606798, 1.41421356, 1.73205081, 2.
                                                               , 2.23606798,
                   2.44948974])
```

• Generate random data: *np.random.normal()* function generates a vector of random normal variables.

```
np.random.normal?
      In [29]:
Docstring:
normal(loc=0.0, scale=1.0, size=None)
Draw random samples from a normal (Gaussian) distribution.
The probability density function of the normal distribution, first
derived by De Moivre and 200 years later by both Gauss and Laplace
independently [2], is often called the bell curve because of
its characteristic shape (see the example below).
The normal distributions occurs often in nature. For example, it
describes the commonly occurring distribution of samples influenced
by a large number of tiny, random disturbances, each with its own
```

 By default, this function will generate random normal variable(s) with mean (*loc*) 0 and standard deviation (*scale*) 1; furthermore, a single random variable will be generated unless the argument to *size* is changed.

```
x = np.random.normal(size=50)
In [31]:
   Out[31]: array([ 1.016527 , 0.29630926, 0.96134982,
                                                          0.31023133, 0.52177376,
                   -0.91979143, -0.92215166, -1.14418682, -0.37095437, -1.09116511,
                   -0.29749639, -0.35213256, 0.3245657,
                                                          1.0522413 , 0.43113357,
                    0.48673004, -0.68270534, -0.19172473,
                                                          0.3020319 , 0.18085425,
                   -1.21985079, 0.29017371, -0.07931557,
                                                          0.84370524, -0.75146806,
                   -1.06603679, -0.28243178, -0.73068695, -1.9812266, 0.47570591,
                    0.51699622, -0.37154697, 1.02783947, 0.82409953, 2.16533796,
                    0.77911666, -0.13074178, 0.20675057, 0.16049944, -0.43860181,
                    1.8335012 , -1.20391787, -0.86482857, 0.09080541, -0.52948043,
                    0.33561294, 0.34369016, -0.72590738, -0.14139 , 1.38267245)
```

 We create an array y by adding an independent N(50, 1) random variable to each element of x.

 The np.corrcoef() function computes the correlation matrix between x and y. The offdiagonal elements give the correlation between x and y.

 In order to ensure that our code provides exactly the same results each time it is run, we can set a random seed using the np.random.default rng() function. This function takes an arbitrary, user-specified integer argument. If we set a random seed before generating random data, then rerunning our code will yield the same results.

 The np.mean(), np.var(), and np.std() functions can be used to compute the mean, variance, and standard deviation of arrays.

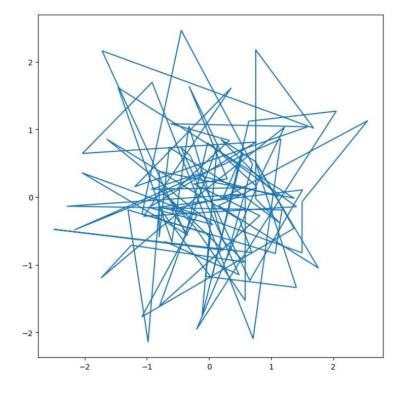
 The np.mean(), np.var(), and np.std() functions can also be applied to the rows and columns of a matrix. To see this, we construct a 10×3 matrix of N(0, 1) random variables, and consider computing its row sums.

• Since arrays are row-major ordered, the first axis, i.e., axis=0, refers to its rows. We pass this argument into the mean() method for the object X.

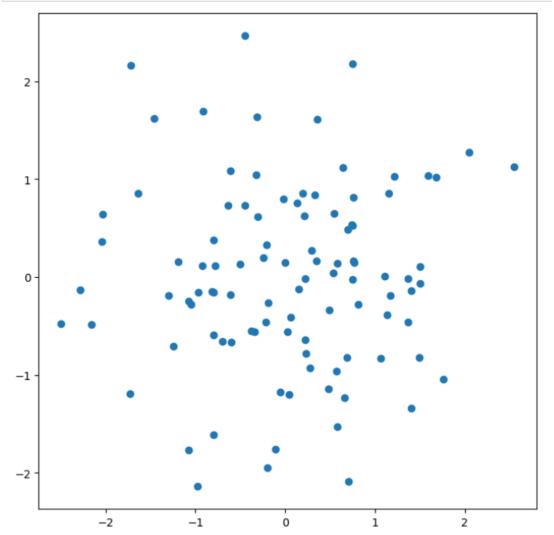
 In Python, common practice is to use the library matplotlib for graphics. However, since Python was not written with data analysis in mind, the notion of plotting is not intrinsic to the language. We will use the subplots() function from matplotlib.pyplot to create a figure and the axes onto which we plot our data.

matplotlib.org/stable/gallery/

- We begin by importing the subplots() function from matplotlib.
- The function returns a tuple of length two: a figure object as well as the relevant axes object.
- We will typically pass figsize as a keyword argument. Having created our axes, we attempt our first plot using its plot() method. To learn more about it, type ax.plot?



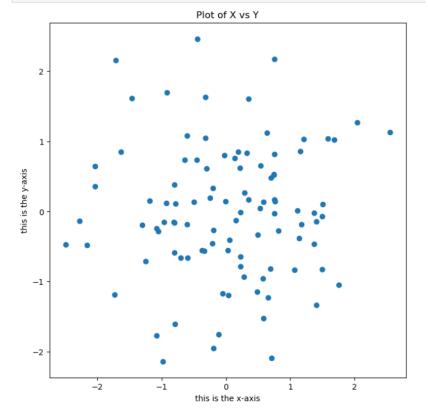
```
In [44]:  | fig , ax = subplots(figsize=(8, 8))
ax.plot(x, y, 'o');
```



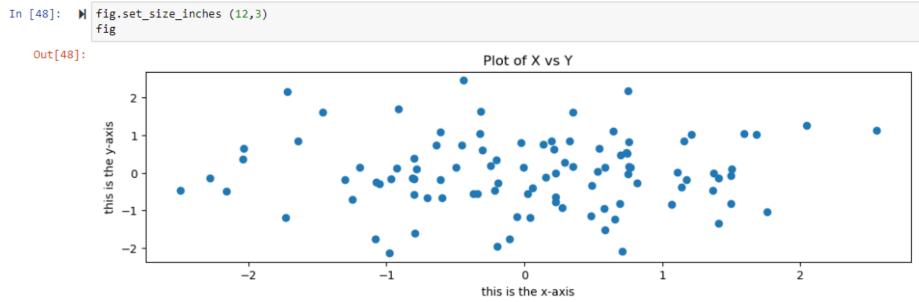
• To label our plot, we make use of the $set_xlabel()$, $set_ylabel()$, and $set_title()$ methods of ax.

In [47]: M fig. ax = subplots(figsize=(8, 8)) ax.scatter(x, y, marker='o')

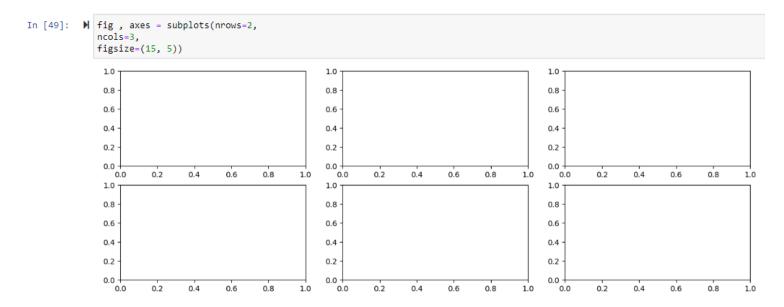




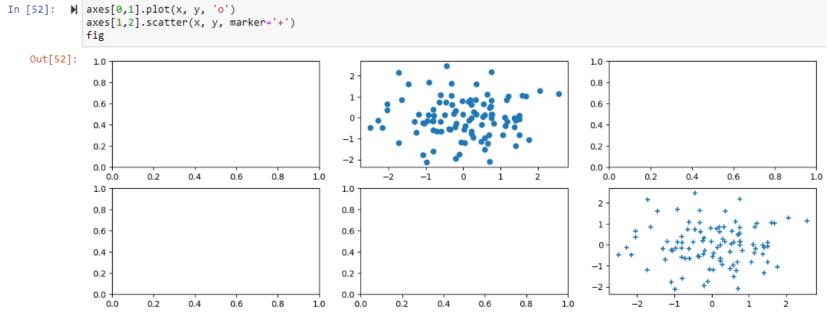
• Having access to the figure object *fig* itself means that we can go in and change some aspects and then redisplay it. Here, we change the size from (8, 8) to (12, 3).



• If we want to create several plots within a figure. This can be achieved by passing additional arguments to *subplots()*. Below, we create a 2 × 3 grid of plots in a figure of size determined by the *figsize* argument.



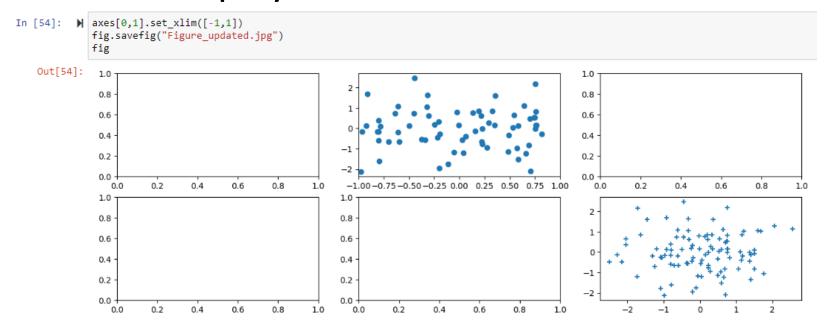
 We now produce a scatter plot with 'o' in the second column of the first row and a scatter plot with '+' in the third column of the second row.



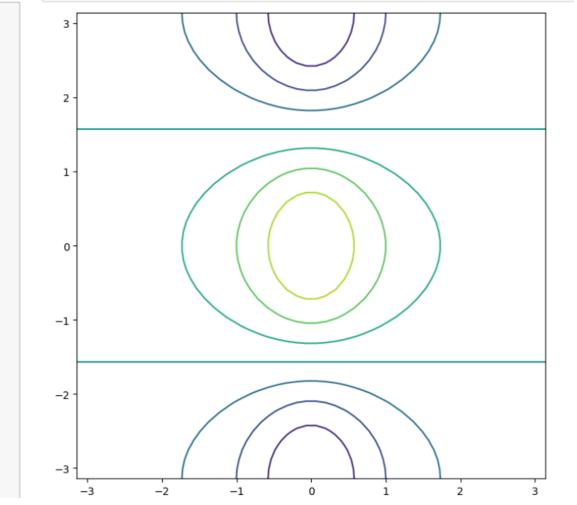
To save the output of fig, we call its savefig()
method. The argument dpi is the dots per
inch, used to determine how large the figure
will be in pixels.

```
In [53]: M fig.savefig("Figure.png", dpi=400)
fig.savefig("Figure.pdf", dpi=200);
```

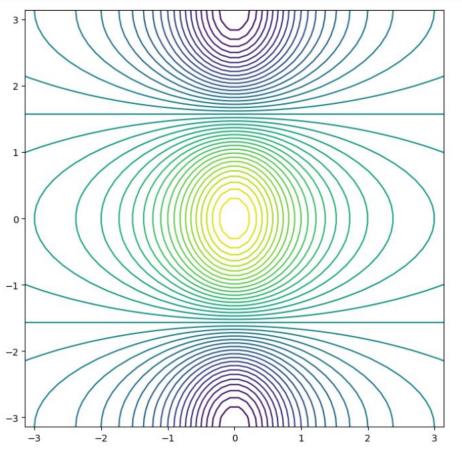
 We can continue to modify fig using step-bystep updates; for example, we can modify the range of the x-axis, re-save the figure, and even re-display it.



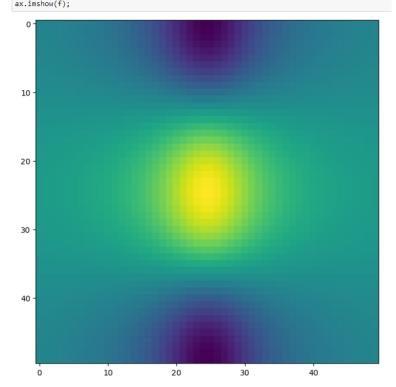
- The ax.contour() method produces a contour plot in order to represent three-dimensional data, similar to a topographical map. It takes three arguments:
 - A vector of x values (the first dimension),
- A vector of y values (the second dimension),
- A matrix whose elements correspond to the z value (the third dimension) for each pair of (x,y) coordinates.



 We can increase the resolution by adding more levels to the image. In [56]: M fig , ax = subplots(figsize=(8, 8)) ax.contour(x, y, f, levels=45);



• The ax.imshow() method is similar to ax.contour(), except that it produces a color-coded plot whose colors depend on the z value. This is known as a heatmap, and is sometimes used to plot temperature in weather forecasts. In [57]: M fig., ax = subplots(figsize-(8, 8))



Sequences and Slice Notation

• The function *np.linspace()* can be used to create a sequence of numbers.

• The function *np.arange()* returns a sequence of numbers spaced out by *step*. If *step* is not specified, then a default value of 1 is used.

```
In [54]:  seq2 = np.arange(0, 10)
    seq2
Out[54]: array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
```

Sequences and Slice Notation

- Slice notation is used to index sequences such as lists, tuples and arrays.
- Suppose we want to retrieve the fourth through sixth (inclusive) entries of a string. We obtain a slice of the string using the indexing notation [3:6].

```
In [55]:  | "hello world"[3:6]
Out[55]: 'lo '
```

which is the same as "hello world"[slice(3,6)]

Indexing Data

Create a two-dimensional numpy array:

 Typing A[1,2] retrieves the element corresponding to the second row and third column. (Python indexes from 0.)

```
In [58]: ► A[1,2]
Out[58]: 6
```

Indexing Rows, Columns, and Submatrices

 To select multiple rows at a time, we can pass in a list specifying our selection. For instance, [1,3] will retrieve the second and fourth rows:

 To select the first and third columns, we pass in [0,2] as the second argument in the square brackets. In this case we need to supply the first argument: which selects all rows.

Indexing Rows, Columns, and Submatrices

 Suppose we want to select the submatrix made up of the second and fourth rows as well as the first and third columns.

```
In [64]: ► A[[1,3],[0,2]]
Out[64]: array([4, 14])
```

 One easy way to do this is as follows. We first create a submatrix by subsetting the rows of A, and then on the fly we make a further submatrix by subsetting its columns.

Indexing Rows, Columns, and Submatrices

The convenience function np.ix_()
allows us to extract a submatrix
using lists, by creating an
intermediate mesh object.

```
idx = np.ix_([1,3],[0,2,3])
A[idx]
array([[ 4, 6, 7],
```

[12, 14, 15]])

 Alternatively, we can subset matrices efficiently using slices. The slice 1:4:2 captures the second and fourth items of a sequence, while the slice 0:3:2 captures the first and third items (the third element in a slice sequence is the step size).

Boolean Indexing

Boolean is a type that equals either *True* or *False* (also represented as Boolean 1 and 0, respectively).
 keep_rows = np.zeros(A.shape[0], bool)

```
keep_rows = np.zeros(A.snape[0], bool)
keep_rows
```

```
array([False, False, False, False])
```

• We now set two of the elements to *True*.

```
keep_rows[[1,3]] = True
keep_rows
array([False, True, False, True])
```

Boolean Indexing

Note that the elements of keep_rows, when viewed as integers, are the same as the values of np.array([0,1,0,1]). We can use == to verify their equality.

True

 The function np.all() has checked whether all entries of an array are True. A similar function, np.any(), can be used to check whether any entries of an array are True.)

Loading Data

- Data sets often contain different types of data, and may have names associated with the rows or columns.
- For these reasons, they typically are best accommodated using a data frame. We can think of a data frame as a sequence of arrays of identical length; these are the columns. Entries in data frame the different arrays can be combined to form a row.
- The pandas library can be used to create and work with data frame objects.

- Before attempting to load a data set, we must make sure that *Python* knows where to find the file containing it.
- If the file is in the same location as this notebook file, then we are all set. Otherwise, the command os.chdir() can be used to change directory. (You will need to call import os before calling os.chdir().)

```
In [66]: import pandas as pd
Auto = pd.read_csv('Auto.csv')
Auto
```

Out[66]:

	mpg	cylinders	displacement	horsepower	weight	acceleration	year	origin	name
0	18.0	8	307.0	130	3504	12.0	70	1	chevrolet chevelle malibu
1	15.0	8	350.0	165	3693	11.5	70	1	buick skylark 320
2	18.0	8	318.0	150	3436	11.0	70	1	plymouth satellite
3	16.0	8	304.0	150	3433	12.0	70	1	amc rebel sst
4	17.0	8	302.0	140	3449	10.5	70	1	ford torino
392	27.0	4	140.0	86	2790	15.6	82	1	ford mustang gl
393	44.0	4	97.0	52	2130	24.6	82	2	vw pickup
394	32.0	4	135.0	84	2295	11.6	82	1	dodge rampage
395	28.0	4	120.0	79	2625	18.6	82	1	ford ranger
396	31.0	4	119.0	82	2720	19.4	82	1	chevy s-10

397 rows × 9 columns

 The whitespace-delimited version of Auto.csv, called Auto.data can be read in using pd.read_csv():

```
In [68]: Auto = pd.read_csv('Auto.data', delim_whitespace=True)
```

 We now take a look at the column of Auto corresponding to the variable horsepower:

```
In [69]:
          ► Auto['horsepower']
   Out[69]: 0
                     130.0
                     165.0
                     150.0
                     150.0
                     140.0
                     86.00
              392
                     52.00
              393
              394
                     84.00
              395
                     79.00
              396
                     82.00
              Name: horsepower, Length: 397, dtype: object
```

 We see that the dtype of this column is object. It turns out that all values of the horsepower column were interpreted as strings when reading in the data. The culprit is the value ? which is being used to encode missing values

```
In [70]: | np.unique(Auto['horsepower'])
   Out[70]: array(['100.0', '102.0', '103.0', '105.0', '107.0', '108.0', '110.0',
                    '112.0', '113.0', '115.0', '116.0', '120.0', '122.0', '125.0',
                    '129.0', '130.0', '132.0', '133.0', '135.0', '137.0', '138.0',
                    '139.0', '140.0', '142.0', '145.0', '148.0', '149.0', '150.0',
                    '152.0', '153.0', '155.0', '158.0', '160.0', '165.0', '167.0',
                    '170.0', '175.0', '180.0', '190.0', '193.0', '198.0', '200.0',
                    '208.0', '210.0', '215.0', '220.0', '225.0', '230.0', '46.00',
                    '48.00', '49.00', '52.00', '53.00', '54.00', '58.00', '60.00',
                    '61.00', '62.00', '63.00', '64.00', '65.00', '66.00', '67.00',
                    '68.00', '69.00', '70.00', '71.00', '72.00', '74.00', '75.00',
                    '76.00', '77.00', '78.00', '79.00', '80.00', '81.00', '82.00',
                    '83.00', '84.00', '85.00', '86.00', '87.00', '88.00', '89.00',
                    '90.00', '91.00', '92.00', '93.00', '94.00', '95.00', '96.00',
                    '97.00', '98.00', '?'], dtype=object)
                                                                                 56
```

To fix the problem, we must provide
 pd.read_csv() with an argument called
 na_values. Now, each instance of ? (missing
 values) in the file is replaced with the value
 np.nan, which means not a number:

The Auto.shape attribute tells us that the data has 397 observations, or rows, and nine variables, or columns.

Out[72]: (397, 9)

There are various ways to deal with missing data.
In this case, since only five of the rows contain
missing observations, we choose to use the
Auto.dropna() method to simply remove these
rows.

Auto_new = Auto.dropna()
Auto_new.shape

(392, 9)

We can use Auto.columns to check the variable names.

 The first argument to the [] method is always applied to the rows of the array. Similarly, passing in a slice to the [] method creates a data frame whose rows are determined by the slice:

```
Auto[:3]
                       displacement
           cylinders
                                       horsepower
                                                    weight
     mpg
    18.0
                               307.0
                                            130.0
                                                    3504.0
    15.0
                               350.0
                                            165.0
                                                    3693.0
    18.0
                               318.0
                                            150.0
                                                    3436.0
```

• An array of Booleans can be used to subset the rows:

```
idx_80 = Auto['year'] > 80
Auto[idx_80]
```

 If we pass in a list of strings to the [] method, then we obtain a data frame containing the corresponding set of columns.

```
Auto[['mpg', 'horsepower']]
                 horsepower
        mpg
0
        18.0
                 130.0
        15.0
                 165.0
        18.0
                 150.0
        16.0
                 150.0
        17.0
                 140.0
392
        27.0
                 86.0
        44.0
                 52.0
393
394
        32.0
                 84.0
395
        28.0
                 79.0
        31.0
                 82.0
396
392 rows x 2 columns
```

 Since we did not specify an index column when we loaded our data frame, the rows are labeled using integers 0 to 396.

 We can use the set_index() method to re-name the rows using the contents of Auto['name']. The index has been set to name.

```
Auto_re = Auto.set_index('name')
Auto re
                                    cylinders
                                                displacement
                              mpg
                     name
chevrolet chevelle malibu
                             18.0
                                                        307.0
         buick skylark 32
                             15.0
                                                        350.0
       plymouth satellite
                             18.0
                                                        318.0
            amc rebel sst
                             16.0
                                                        304.0
Auto_re.columns
Index(['mpg', 'cylinders', 'displacement', 'horsepower',
       'weight', 'acceleration', 'year', 'origin'],
      dtype='object')
```

We can access rows of the data frame by name using the loc[]
 method of Auto: [rows = ['amc rebel sst', 'ford torino']

Auto_re.loc[rows]

```
mpg cylinders displacement horsepower ...
name
amc rebel sst 16.0 8 304.0 150.0 ...
ford torino 17.0 8 302.0 140.0 ...
```

 As an alternative to using the index name, we could retrieve rows and columns of Auto using the iloc[] method:

```
Auto_re.iloc[[3,4]]

Auto_re.iloc[:,[0,2,3]]

Auto_re.iloc[[3,4],[0,2,3]]
```

For Loops

```
total = 0
for value in [3,2,19]:
    total += value
print('Total is: {0}'.format(total))
```

Total is: 24

• The indented code beneath the line with the *for* statement is run for each value in the sequence specified in the *for* statement. The loop ends either when the cell ends or when code is indented at the same level as the original *for* statement. Loops can be nested by additional indentation.

```
total = 0
for value in [2,3,19]:
    for weight in [3, 2, 1]:
        total += value * weight
print('Total is: {0}'.format(total))
```

For Loops

 To compute the average value of a random variable that takes on possible values 2, 3 or 19 with probability 0.2, 0.3, 0.5 respectively we would compute the weighted sum. Tasks such as this can often be accomplished using the zip() function that loops over a sequence of tuples.

Weighted average is: 10.8

String Formatting

- Inserting the value of something into a string is a common task.
- For example, we may want to loop over the columns of a data frame and print the percent missing in each column.
- Let's create a data frame D with columns in which 20% of the entries are missing, i.e., set to np.nan. We'll create the values in D from a normal distribution with mean 0 and variance 1 using rng.standard_normal() and then overwrite some random entries using rng.choice().

String Formatting

	food	bar	pickle	snack	popcorn
0	0.345584	0.821618	0.330437	-1.303157	NaN
1	NaN	-0.536953	0.581118	0.364572	0.294132
2	NaN	0.546713	NaN	-0.162910	-0.482119

```
Column "food" has 16.54% missing values
Column "bar" has 25.98% missing values
Column "pickle" has 29.13% missing values
Column "snack" has 21.26% missing values
Column "popcorn" has 22.83% missing values
```

 We can use the ax.plot() or ax.scatter() functions to display the quantitative variables. However, Python does not know to look in the Auto data set for those variables.

```
fig, ax = subplots(figsize=(8, 8))
ax.plot(horsepower, mpg, 'o');
```

```
NameError: name 'horsepower' is not defined
```

 We can address this by accessing the columns directly:

```
fig, ax = subplots(figsize=(8, 8))
ax.plot(Auto['horsepower'], Auto['mpg'], 'o');
```

 Alternatively, we can use the plot() method with the call Auto.plot(). Using this method, the variables can be accessed by name.

```
ax = Auto.plot.scatter('horsepower', 'mpg');
ax.set_title('Horsepower vs. MPG')
```

• The columns of a data frame can be accessed as attributes: try typing in *Auto.horsepower*.

- Auto.cylinders.dtype reveals that it is being treated as a quantitative variable. However, since there is only a small number of possible values for this variable, we may wish to treat it as qualitative.
- Below, we replace the cylinders column with a categorical version of Auto.cylinders.

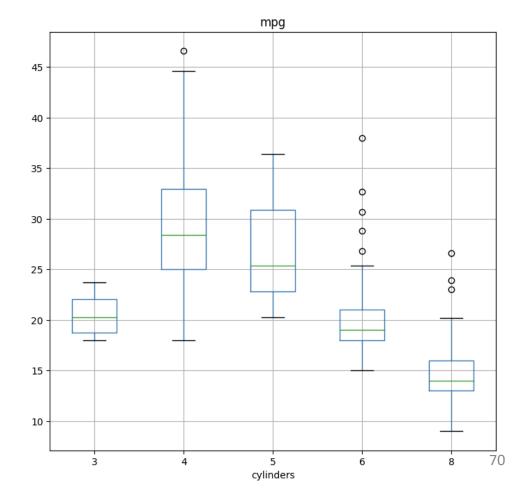
```
Auto.cylinders = pd.Series(Auto.cylinders, dtype='category')
Auto.cylinders.dtype
```

CategoricalDtype(categories=[3, 4, 5, 6, 8], ordered=False)

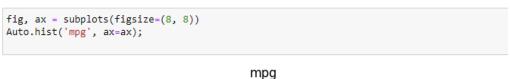
 Now that cylinders is qualitative, we can display it using the boxplot() method.

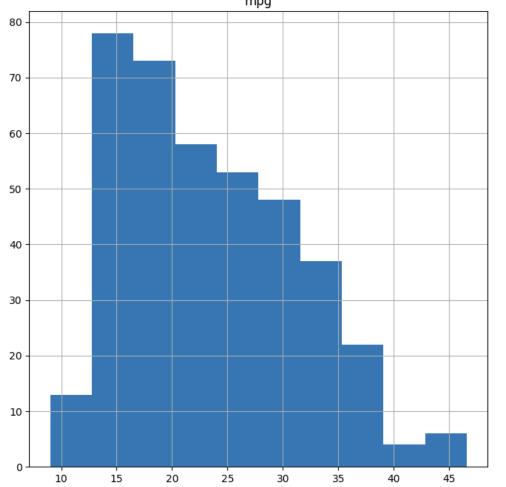


Boxplot grouped by cylinders



The hist()
 method can be
 used to plot a
 histogram.





We can use the pd.plotting.scat ter matrix() function to create a scatterplot matrix to visualize all of the pairwise relationships between the columns in a data frame.

pd.plotting.scatter_matrix(Auto); orsep**olispl**acement. 400 200 200 100 5000 yeaacceleration ັ້ວ 7**g** origin

displacementrsepower

용요 유유 Acceleration year

origin

72

 The describe() method produces a numerical summary of each column in a data frame.

Auto[['mpg', 'we	eight']].de	scribe()
	mpg	weight	
count	392.000000	392.000000	
mean	23.445918	2977.584184	
std	7.805007	849.402560	
min	9.000000	1613.000000	
25%	17.000000	2225.250000	
50%	22.750000	2803.500000	
75%	29.000000	3614.750000	
max	46.600000	5140.000000	