PYTHON FOR DATA ANALYSIS

Content from Jose Portilla's Udemy course *Learning Python for Data Analysis and Visualization*https://www.udemy.com/learning-python-for-data-analysis-and-visualization/
Notes by Michael Brothers, available on http://github.com/mikebrothers/data-science/

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LEARNING PYTHON FOR DATA ANALYSIS & VISUALIZATION Udemy course by Jose Portilla (notes by Michael Brothers)

What's What:

Numpy – fundamental package for scientific computing, working with arrays

Pandas – create high-performance data structures, Series, Data Frames. incl built-in visualization, file reading tools

Matplotlib - data visualization package

Seaborn Libraries – heatmap plots et al

Beautiful Soup - a web-scraping tool

SciKit-Learn – machine learning library

Skills:

Importing data from a variety of formats: JSON, HTML, text, csv, Excel

Data Visualization – using Matplotlib and the Seaborn libraries

Portfolio – set up a portfolio of data projects on GitHub

Machine Learning – using SciKit Learn

Resources:

stock market analysis (access Yahoo finance using pandas datareader)

FDIC list of failed banks (pull data from html)

Kaggle Titanic data set

political election data set

http://www.data.gov (home of the US Government's open data)

http://AWS.amazon.com/public-data-sets/ (Amazon web services public data sets)

http://www.google.com/publicdata/directory

create personal accounts on GitHub and Kaggle

Appendix Materials:

Statistics – includes using SciPy to create distributions & solve statistics problems

SQL with Python – includes using SQLAlchemy to fully integrate SQL with Python to run SQL queries from a Python environment. Also performing basic SQL commands with Python and pandas.

Web Scraping with Python – using Python web requests and the Beautiful-Soup library to scrape the web for data

For Further Reading:

Numpy: http://docs.scipy.org/doc/numpy/reference/

Numpy Universal Functions (ufuncs): http://docs.scipy.org/doc/numpy/reference/ufuncs.html#available-ufuncs

Numpy supplemental materials: http://cs231n.github.io/python-numpy-tutorial/

Philosophy:

What's the difference between a Series, a DataFrame and an Array? (answers by Jose Portilla)

A Numpy Array is the basic data structure holding the data itself and allowing you to store and get elements from it.

A **Series** is built on top of an array, allowing you to label the data and index it formally, as well as do other pandas related Series operations.

A **DataFrame** is built on top of Series, and is essentially many series put together with different column names but sharing the same index.

Also, a 1-d numpy array is **not** a **list**. A list is a built-in data structure in regular Python, a numpy array is an object type only available once you've set up numpy. It is able to perform operations much faster than a list due to built-in optimizations.

Arrays are NumPy data types while Series and DataFrame are Pandas data types. They have different available methods and attributes.

```
NUMPY
```

```
import numpy as np
```

[0, 0, 0]])

do this for every new Jupyter notebook

```
Creating Arrays
my list1 = [1, 2, 3, 4]
my array1 = np.array(my list1) creates a 1-dimensional array from a list
my array1
array([1, 2, 3, 4])
my list2 = [11, 22, 33, 44]
my lists = [my list1, my list2]
my array2 = np.array(my lists) creates a multi-dimensional array from a list of lists
my array2
array([[ 1, 2, 3, 4],
      [11, 22, 33, 44]])
array 2d = (([1,2,3], [4,5,6])) creating from scratch requires <u>two</u> sets of parentheses!
my array2.shape describes the size & shape of the array (rows, columns)
(2L, 4L)
                      describes the data type of the array
my array2.dtype
dtype('int32')
Special Case Arrays
np.zeros(5)
                                                             called the "identity array"
                                             np.eve(5)
array([ 0., 0., 0., 0., 0.])
                                             array([[ 1., 0., 0.,
                                                                       0., 0.1,
                                                    [ 0., 1., 0.,
                                                                            0.],
                                                                       0.,
np.ones((4,4))
                                                    [ 0., 0., 1.,
                                                                       0.,
                                                                            0.],
array([[ 1., 1.,
                   1., 1.],
                                                    [ 0., 0., 0.,
                                                                     1., 0.],
       [ 1., 1.,
                   1.,
                        1.],
                                                     [ 0., 0., 0.,
                                                                       0.,
                                                                           1.11)
       [ 1., 1., 1., 1.],
       [ 1., 1., 1., 1.]])
                                             dtype ('float64') for the above arrays
np.empty(5)
                                             np.arange([start,] stop[, step])
np.empty((3,4))
                                             np.arange (5, 10, 2) uses a range
resemble zeros arrays
                                             array([5, 7, 9])
Using Arrays and Scalars
from future import division if running Python v2
arr1 = np.array([[1,2,3], [8,9,10]]) note the double parentheses/brackets
arr1
array([[ 1, 2, 3],
     [ 8, 9, 10]])
                               Multiplying arrays:
Adding arrays:
arr1+arr1
                               arr1*arr1
array([[ 2, 4, 6],
                               array([[ 1, 4, 9],
     [16, 18, 20]])
                                [ 64, 81, 100]])
Subtracting arrays:
                               <u>Dividing arrays:</u> (Float return)
                               arr1/arr1
arr1-arr1
array([[0, 0, 0],
```

array([[1., 1., 1.],

[1., 1., 1.]])

```
Arithmetic operations with scalars on arrays:
```

Indexing Arrays

```
Arrays are sequenced. They are modified in place by slice operations.
```

Note that the changes *also* occur in our original array.

Data is not copied, it's a view of the original array. This avoids memory problems.

```
arr_copy = arr.copy() To get a copy, you need to be explicit
arr_copy
array([99, 99, 99, 99, 99, 99, 6, 7, 8, 9, 10])
```

Indexing a 2D Array

```
format follows arr_2d[row][col] or arr_2d[row,col]
arr_2d[1] grab a row
array([20, 25, 30])
```

arr_2d[1][0] or arr_2d[1,0] grab an individual element
20

Slicing a 2D Array

Fancy Indexing

```
arr
array([[ 0., 10., 20., 30., 40.],
       [ 1., 11., 21., 31., 41.],
       [ 2., 12., 22.,
                            32.,
                                  42.11)
                      fancy indexing allows a selection of rows in any order using embedded brackets
arr[[2,1]]
array([[ 2., 12., 22., 32., 42.],
                                               (note that arr[2,1] returns 12.0)
       [ 1., 11., 21., 31., 41.]])
>>> a[0,3:5]
array([3,4])
                                1
                                    2
                                       3
                                              5
>>> a[4:,4:]
array([[44, 45],
                            10 | 11
                                   12
                                      13
                                          14
                                             15
       [54, 55]])
                            20
                               21
                                   22
                                      23
                                             25
                                          24
>>> a[:.2]
array([2,12,22,32,42,52])
                                   32
                            30
                               31
                                      33
                                          34
                                             35
>>> a[2::2,::2]
                                   42
                            40 41
                                      43
                                          44
                                             45
array([[20,22,24]
                                          54
                                             55
                                   52
                                      53
                            50 | 51
       [40,42,44]])
```

Source: http://www.scipy-lectures.org/ images/numpy indexing.png

Array Transposition

```
arr = np.arange(24).reshape((4,6)) create an array
arr
array([[ 0, 1, 2, 3, 4, 5],
       [ 6, 7, 8, 9, 10, 11],
       [12, 13, 14, 15, 16, 17],
       [18, 19, 20, 21, 22, 23]])
                                   transpose the array (this does NOT change the array in place)
arr.T
array([[ 0, 6, 12, 18],
       [ 1, 7, 13, 19],
       [ 2, 8, 14, 20],
       [3, 9, 15, 21],
       [ 4, 10, 16, 22],
       [ 5, 11, 17, 23]])
                                   take the dot product of these two arrays
np.dot(arr.T,arr)
array([[504, 540, 576, 612, 648, 684],
                                            504 = (0*0) + (6*6) + (12*12) + (18*18)
       [540, 580, 620, 660, 700, 740],
                                              540 = (0*1) + (6*7) + (12*13) + (18*19)
       [576, 620, 664, 708, 752, 796],
       [612, 660, 708, 756, 804, 852],
       [648, 700, 752, 804, 856, 908],
       [684, 740, 796, 852, 908, 964]])
```

See https://www.mathsisfun.com/algebra/matrix-multiplying.html for a simple explanation of dot products!

You can also transpose a 3D matrix:

```
arr3d = np.arange(18).reshape((3,3,2))
```

```
arr3d
                            arr3d.transpose((1,0,2))
array([[[ 0, 1],
                            array([[[ 0, 1],
       [ 2, 3],
                                    [ 6, 7],
       [4, 5]],
                                    [12, 13]],
      [[6, 7],
                                   [[2, 3],
       [8, 9],
                                    [8, 9],
       [10, 11]],
                                    [14, 15]],
      [[12, 13],
                                   [[4, 5],
       [14, 15],
                                    [10, 11],
       [16, 17]])
                                    [16, 17]])
```

If you need to get more specific use **swapaxes**:

Universal Array Functions

Binary Functions (require two arrays):

```
np.add (A, B) returns sum of matching values of two arrays np.maximum (A, B) returns maximum between matching values of two arrays
```

Random number generator:

```
np.random.randn(10) random array (normal distribution)
array([-0.10313268, 1.05811992, -1.98543659, -0.43591721, 0.03393424,
-1.15738081, -0.35316064, 1.12707714, -0.09061522, 0.28226307])
```

For full and extensive list of all universal functions

```
website = "http://docs.scipy.org/doc/numpy/reference/ufuncs.html#available-ufuncs"
import webbrowser
webbrowser.open(website) conveniently opens site from within Jupyter notebook!
```

Array Processing

```
import numpy as np
import matplotlib.pyplot as plt import the pyplot libraries from matplotlib
                                          which let us visualize the grids & meshes we'll be making
                                          this lets us see these visualizations in Jupyter notebooks
%matplotlib inline
```

```
Using matplotlib.pyplot for visualization
                                 creates a 1-d array with 1000 data points
points = np.arange(-5, 5, 0.01)
dx, dy=np.meshgrid (points, points) creates a grid (returns coordinate matrices from the vectors we give it)
           these are our rows:
dx
array([[-5. , -4.99, -4.98, ..., 4.97,
                                                    4.99],
                                            4.98,
       [-5., -4.99, -4.98, \ldots,
                                    4.97,
                                            4.98,
                                                    4.991,
       [-5., -4.99, -4.98, \ldots, 4.97,
                                            4.98,
                                                    4.991,
       [-5., -4.99, -4.98, \ldots, 4.97,
                                           4.98,
                                                    4.991,
       [-5., -4.99, -4.98, \ldots, 4.97, 4.98,
                                                    4.991,
       [-5., -4.99, -4.98, \ldots, 4.97, 4.98,
                                                    4.9911)
           these are our columns: (note that values increase downward)
dy
array([[-5., -5., -5., -5., -5., -5., -5.]),
       [-4.99, -4.99, -4.99, ..., -4.99, -4.99, -4.99]
       [-4.98, -4.98, -4.98, \ldots, -4.98, -4.98, -4.98],
       [4.97, 4.97, 4.97, \ldots, 4.97, 4.97, 4.97],
       [ 4.98, 4.98, 4.98, ..., 4.98, 4.98,
                                                    4.98],
       [4.99, 4.99, 4.99, \ldots, 4.99, 4.99,
                                                   4.9911)
z = (np.sin(dx) + np.sin(dy))
                                 this is just an evaluating function
array([[ 1.91784855e+00, 1.92063718e+00, 1.92332964e+00, ...,
         -8.07710558e-03, -5.48108704e-03, -2.78862876e-03],
                                               1.92611827e+00, ...,
       [ 1.92063718e+00,
                            1.92342581e+00,
         -5.28847682e-03, -2.69245827e-03, -5.85087534e-14],
       [ 1.92332964e+00,
                            1.92611827e+00, 1.92881072e+00, ...,
                            -5.63993297e-14, 2.69245827e-03],
         -2.59601854e-03,
                            -5.28847682e-03, -2.59601854e-03, ...,
       [-8.07710558e-03,
         -1.93400276e+00,
                            -1.93140674e+00, -1.92871428e+00],
       [-5.48108704e-03,
                            -2.69245827e-03, -5.63993297e-14, ...,
         -1.93140674e+00,
                            -1.92881072e+00, -1.92611827e+00],
       [ -2.78862876e-03, -5.85087534e-14,
                                                2.69245827e-03, ...,
         -1.92871428e+00, -1.92611827e+00, -1.92342581e+00]])
plt.imshow(z); plot the array (semicolon avoids extra Out line)
red/blue colored plot for evaluating function, y-axis from 1000-0, x-axis from 0-1000
plt.colorbar(); needs to be in same cell as plt.imshow(z)
adds vertical colorbar to right of plot (red 2.0 to blue -2.)
plt.title("Plot for sin(x)+sin(y)");
```

adds title above plot SEE PLOT IN SEPARATE FILE: PythonDataVisualizations.pdf

```
Using numpy.where
```

```
A = np.array([1,2,3,4])
B = np.array([100, 200, 300, 400])
condition = np.array([True, True, False, False]) a Boolean array
The slow way: Using a list comprehension
answer1 = [(A val if cond else B val) for A val, B val, cond in zip(A, B, condition)]
answer1
[1, 2, 300, 400] Problems include speed issues and multi-dimensional array issues
The numpy.where way:
answer2 = np.where(condition, A, B) follows(test, if true, if false)
answer2
array([ 1, 2, 300, 400])
Using numpy.where for 2D manipulation:
from numpy.random import randn
arr = randn(5, 5)
np.where (arr < 0,0,arr) Where array is less than zero, make that value zero, otherwise leave as is
                              , 1.56891548, 0. , 1.61030401],
array([[ 0.45983701, 0.
       [ 0. , 0. , 0. , 0. , 0. ], [ 0.96909611, 0. , 0. , 0.69907836, 1.41859086],
       [ 0. , 0.
       [ 0. , 1.42554561, 1.30200218, 1.77784525, 0.8120543 ],
       [ 1.39031869, 0.14319058, 0.11438954, 0. , 0.
                                                                         11)
More statistical tools:
arr = np.array([[1,2,3],[4,5,6],[7,8,9]])
arr.sum() returns 45
arr.sum(0) returns array([12,15,18]) sums along vertical axes
arr.mean() returns 5.0
                                         Note there are no "median" or "mode" functions
arr.var() returns 6.6666666666666 variance
arr.std() returns 2.5819888974716112 standard deviation
Any and all for processing Boolean arrays:
bool arr = np.array([True, False, True])
bool arr.any() returns True
bool arr.all() returns False
Sort, Unique and In1d:
arr = randn(5,5)
arr.sort()
                                         sorts each row individually, in place
np.apply along axis (sorted, 0, arr) sorts each item horizontally
countries = np.array(['France', 'Germany', 'USA', 'Russia', 'USA', 'Mexico'])
np.unique(countries)
array(['France', 'Germany', 'Mexico', 'Russia', 'USA'],
      dtype='|S7')
np.in1d(['France','USA','Sweden'],countries)
array([ True, True, False], dtype=bool)
```

Array Input and Output

```
import numpy as np
```

```
Insert an element into an array (see http://docs.scipy.org/doc/numpy-1.10.0/reference/generated/numpy.insert.html)
a = np.array([[1, 1], [2, 2], [3, 3]])
а
array([[1, 1],
        [2, 2],
        [3, 311)
np.insert(a, 1, 5) inserts a 5 before index 1 and flattens the array (but not in-place!)
array([1, 5, 1, 2, 2, 3, 3])
np.insert(a, 1, 5, axis=1) inserts a 5 before index 1 along the vertical axis (but not in-place!)
array([[1, 5, 1],
        [2, 5, 2],
        [3, 5, 3]])
Saving an array to a binary (.npy) file
arr = np.arange(5)
                                 saves the array on disk in binary format (file extension .npy)
np.save('my array',arr)
arr = np.arange(10)
                                 here we create a different array with the same name
np.load('my array.npy')
                                 here we load the first array we created
array([0, 1, 2, 3, 4])
Saving multiple arrays into a zip (.npz) file
                                                     saves 2 copies of arr to one file
np.savez('two arrays.npz', x=arr, y=arr)
Loading multiple arrays:
archive array = np.load('two arrays.npz')
                                                     Note: .load works for binary and zip
archive array['x']
                                                     calls the first array from the file
array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
Saving and loading text files
arr = np.array([[1,2,3],[4,5,6]])
np.savetxt('my test text.txt',arr,delimiter=',')
arr = np.loadtxt('my test text.txt',delimiter = ',')
arr
array([[ 1., 2., 3.],
        [ 4., 5., 6.]])
```

```
PANDAS
```

```
import numpy as np
import pandas as pd
from pandas import Series, DataFrame this saves us from typing 'pd.Series' and 'pd.DataFrame' each time
```

WORKING WITH SERIES

Creating a Series (an array of data values and their index)

Creating a Series with a named index

```
coins = Series([.01,.05,.10,.25],index=['penny','nickel','dime','quarter'])
coins
        0.01
penny
nickel
        0.05
dime
        0.10
quarter
        0.25
dtype: float64
coins[coins>.07]
    0.10
dime
quarter 0.25
dtype: float64
'penny' in coins returns True (although 0.25 in coins returns False)
```

Converting a Series to a Python dictionary

Passing an index with the dictionary can reload a Series in order:

Use isnull and notnull to find missing data

```
pd.isnull(coins3['SBAnthony']) returns True
pd.notnull(coins3['penny']) returns True
```

Adding two Series together

```
series1 + series2 adds items by index, including null-value items
```

Labeling Series Indexes

```
coins3.index.name = 'Coins' puts a label above the index list (.values does not have a name method)
```

Checking for Unique Values and their Counts

```
ser1 = Series(list('abacab'))
ser1.unique() returns array(['a', 'b', 'c'], dtype=object)
ser1.value_counts() returns see the DataFrames section on value_counts for more info
a    3
b    2
c    1
dtype: int64
```

Rank and Sort

Sort by Index Name using .sort index:

```
ser1 = Series(range(3),index=['C','A','B'])
ser1.sort_index() returns ser1, but in index order (A:1,B:2,C:0)
Note: this does NOT sort ser1 in place.
For that use either ser1 = ser1.sort_index() or ser1.sort_index(inplace=True)
```

Sort by Value using .sort_values:

```
ser1.sort_values() returns ser1, but in value order (C:0,A:1,B:2)
Note:.order works, but throws a "FutureWarning: order is deprecated, use sort_values(...)" As above,
use either ser1 = ser1.sort index() or ser1.sort index(inplace=True) to sort in place
```

Rank

```
ser1.rank() returns an integer rank from 1 to len(ser1) for each index (low to high)
NOTE: in the case of ties, .rank returns floats (1, 2.5, 2.5, 4)
```

WORKING WITH DATAFRAMES

For more info: http://pandas.pydata.org/pandas-docs/stable/dsintro.html#dataframe

Creating a DataFrame

```
import numpy as np
import pandas as pd
from pandas import Series, DataFrame
dframe = DataFrame (np.arange(12).reshape(4,3))
dframe is constructed by casting a 4-row by 3-col numpy array as a pandas DataFrame, pre-filled with values 0-11.
Here the index defaults to [0,1,2,3], the columns to [0,1,2]
```

Constructing a DataFrame from a Dictionary:

```
data = {'City':['SF','LA','NYC'],'Population':[837000,3880000,8400000]}
city_frame = DataFrame (data)
Creates a DataFrame with columns labeled City and Population, indexes of [0,1,2]
```

Adding a Series to an existing DataFrame:

```
colors = Series(["Blue", "Red"], index=[4,1])
dframe['Color']=colors
dframe now has a Color column with Blue matched to index 4, Red to 1, and NaN after everything else.
```

Reading a DataFrame from a webpage (using edit/copy):

Grab NFL Win-Loss data from Wikipedia:

nfl frame

| | Rank | Team | Won | Lost | Tied | Pct. | First NFL Season | Total Games | Divison |
|---|------|---------------------|-----|------|------|----------|------------------|--------------------|-----------|
| 0 | 1 | Chicago Bears | 741 | 555 | 42 | 0.57 | 1920 | 1338 | NFC North |
| 1 | 2 | Dallas Cowboys | 480 | 364 | 6 | 0.568 | 1960 | 850 | NFC East |
| 2 | 3 | Green Bay Packers | 720 | 547 | 37 | 0.566 | 1921 | 1304 | NFC North |
| 3 | 4 | Miami Dolphins | 429 | 335 | 4 | 0.561 | 1966 | 768 | AFC East |
| 4 | 5 | San Francisco 49ers | 520 | 436 | 14 | .553 [a] | 1950 | 1019 | NFC West |

Note that pandas automatically adds an index in the left-most column. Data as of 1/19/16.

Grab column names:

```
Grab a specific column – 1 word name:
```

Grab a specific column – multiword names:

```
nfl frame.Team nfl frame['First Season']
```

```
Display specific data columns:
```

```
DataFrame (nfl_frame, columns=['Team', 'First Season', 'Total Games'])
    This returns a new DataFrame extracted from nfl_frame
```

NOTE: if you ask for a column that doesn't exist in the original, you get a column filled with null values (NaN)

Display a specific number of rows:

```
nfl_frame.head() retrieves the first 5 rows
nfl_frame.head(3) retrieves the first 3 rows
nfl_frame.tail() retrieves the last 5 rows
```

Grab a record by its index:

nfl frame.ix[3] returns an object with column names & values (the .ix method stands for "index")

Rename index and columns (dict method):

Rename a specific column:

```
nfl frame.rename(columns = {'First NFL Season':'First Season'}, inplace=True)
```

Index Objects

Set a Series index to be its own object:

```
coin_index = coins.index
coin_index
Index([u'penny', u'nickel', u'dime', u'quarter'], dtype='object')
coin_index[2] returns 'dime'
Note: Indexes are immutable (coin index[2]='fred' is not valid code)
```

Reindexing

```
ser1 = Series([1,2,3,4],index=['A','B','C','D'])
ser2 = ser1.reindex(['A','B','C','D','E','F'])
Creates a new Series, with null values for 'E' and 'F'
NOTE: this also converted the Series from dtype int64 to float64. ser2['C'] returns 3.0

ser2.reindex(['A','B','C','D','E','F','G'],fill_value=0)
Adds a new index 'G' with a value of 0. Indexes 'E' and 'F' are both still null values.
ser2.reindex(['B','A','C','D','E','F','G'])
Changes the order of index:value pairs (it doesn't reassign the index) B:2 is now ahead of A:1
ser2.reindex(['C','D','E','F'])
Removes A:1, B:2 and G:0 from Series ser2.
However: ser2.reindex(['A','B','C','D','E','F','G'])
brings back A:1 and B:2 (because ser2 is based on ser1) but not G:0. It assigns a null value to G.
```

Interpolating values between indices:

```
ser3 = Series(['USA', 'Mexico', 'Canada'], index=[0,5,10])
ser3.reindex(range(15), method='ffill') uses a "forward fill" method
ser3 now has 15 members. Index 0-4 = 'USA', 5-9 = 'Mexico' and 10-14='Canada'
```

```
Reindexing onto a DataFrame:
```

```
from numpy.random import randn
dframe = DataFrame(randn(25).reshape((5,5)),index=['A','B','D','E','F'],
    columns=['col1','col2','col3','col4','col5'])
dframe2 = dframe.reindex(['A','B','C','D','E','F'])
Inserts a new row 'C' between A and B filled with null values
```

Reindexing DataFrame columns:

```
dframe2.reindex(columns=['col1', 'col2', 'col3', 'col4', 'col5', 'col6']
Inserts a new column 'col6' at the end filled with null values (you have to call "columns" specifically)
```

Reindex quickly using .ix:

```
dframe.ix[[rows], [columns]]
you could say "newrows=['A',...'F']" and "newcols=['col1',...'col6']" and then dframe.ix[newrows,newcols]
```

Drop Entry – Rows:

```
ser1 = Series(np.arange(3),index=['a','b','c'])
ser1.drop('b') Displays the Series without row 'b' (although this row still belongs to the series)
Similarly, dframe1.drop('index1') drops a row from a DataFrame
```

Drop Entry – Columns:

```
dframe1.drop('col4', axis=1) axis=0 rows/ axis=1 columns, or axis=rows / axis=columns
```

Selecting Entries in a Series:

```
ser1 = 2*ser1 to avoid confusion in the future
ser1
A     0
B     2
C     4
dtype: int32
You can grab an entry by index name: ser1['B'] returns 2
    or by index value: ser1[1] returns 2
    or by a range of values: ser1[0:2] returns rows A:0 and B:2
```

ser1 = Series(np.arange(3),index=['A','B','C'])

You can grab entries by logic: ser1[ser1>3] returns row C:4

You can change values using logic: ser1[ser1>3] = 10 changes C

or by a list of index names: ser1[['A', 'B']] returns rows A:0 and B:2

Selecting Entries in a DataFrame:

```
dframe = DataFrame (np.arange (25).reshape ((5,5)),
    index=['NYC','LA','SF','DC','Chi'],columns=['A','B','C','D','E'])
You can grab entries by column name: dframe['B'] returns all rows with column B values
You can grab multiple columns with a list of names: dframe[['B','E']]
You can grab specific rows using Boolean: dframe[dframe['C']>8]
You can grab a specific cell by column and row: dframe['B']['LA']
To show a Boolean DataFrame: dframe>10
Returns the full DataFrame with True/False in each cell as appropriate
```

You can grab a row using .ix: dframe.ix['LA'] returns row LA as a Series with column names as its index NOTE: dframe.ix[1] also works to grab the 2nd row. dframe.ix['LA']['B'] grabs a single cell.

Data Alignment

```
ser1 = Series([0,1,2],index=['A','B','C'])
ser2 = Series([3,4,5,6],index=list('ABCD')) a nice little shortcut
                      ser2
ser1
    0
A
                      A
                           3
     1
В
                      В
                           4
C
    2
                      C
                           5
                           6
dtype: int64
                      D
                      dtype: int64
```

So what happens when we add these together?

```
ser1 + ser2
A     3
B     5
C     7
D     NaN
dtype: float64
```

Because ser1 didn't have a value for D, it replaced it with a null.

The same behavior occurs with DataFrames (null values are assigned for any unmatched field)

Use .add to assign fill values:

```
ser1.add(ser2, fill_value=0) this adds 0 to whatever hasn't matched
NOTE: ser2.add(ser1, fill value=0) returns the same thing!
```

When using .add/fill_value with dataframes, null values are assigned when there are no prior values in a cell (at the intersection where new rows from one DataFrame meet new columns from another)

Operations Between a Series and a DataFrame

```
dframe1 = DataFrame(np.arange(9).reshape(3,3),columns=list('ADC'),
   index=['NYC','SF','LA'])
ser1 = dframe1.ix[0] so ser1 takes the 'NYC' row and values
dframe1 - ser1 returns the dframe1 DataFrame, but now all the 'NYC' values = 0
```

A DataFrame column is itself a Series, so Series methods apply:

To count the unique values in a DataFrame column:

```
dframe['col1'].value_counts()

dframe['col1'].value_counts(ascending=True)

dframe['col1'].value_counts(sort=False)

dframe['col1'].value_counts(dropna=False)

returns the count from lowest to highest
returns the count in index order
includes a count of null values
```

For more info, incl normalize & bin parameters:

http://pandas.pydata.org/pandas-docs/version/0.17.1/generated/pandas.Series.value counts.html

To retrieve rows that contain a particular value:

```
dframe[dframe.col1=='value'] or
dframe[dframe['column 1']=='value']
```

Summary Statistics on DataFrames

```
arr = np.array([[1,2,np.nan],[np.nan,3,4]]) inserts null values
dframe1 = DataFrame(arr,index=['A','B'],columns = ['One','Two','Three'])
dframe1
```

| | One | Two | Three |
|---|-----|-----|-------|
| Α | 1 | 2 | NaN |
| В | NaN | 3 | 4 |

```
dframe1.sum()
                      dframe1.sum(axis=1)
```

One 1 5 7 Two В

Three 4 dtype: float64

dtype: float64

Two

dframe1.min() dframe1.idxmin() .idxmin returns the index of the lowest value

One 1 One A .max and .idxmax work as expected 2

Α

4 Three Three

dtype: float64 dtype: object

dframe1.cumsum()

redisplays the DataFrame with accumulation sums

| | One | Two | Three |
|---|-----|-----|-------|
| Α | 1 | 2 | NaN |
| В | NaN | 5 | 4 |

provides useful summary statistics dframe1.describe()

Two

| | One | Two | Three |
|-------|-----|----------|-------|
| count | 1 | 2.000000 | 1 |
| mean | 1 | 2.500000 | 4 |
| std | NaN | 0.707107 | NaN |
| min | 1 | 2.000000 | 4 |
| 25% | 1 | 2.250000 | 4 |
| 50% | 1 | 2.500000 | 4 |
| 75% | 1 | 2.750000 | 4 |
| max | 1 | 3.000000 | 4 |

For more information about Covariance and Correlation Check out these great videos! Video credit: Brandon Foltz.

from IPython.display import YouTubeVideo YouTubeVideo('xGbpuFNR1ME') #Covariance (26:22) YouTubeVideo('4EXNedimDMs') #Correlation (27:05)

Correlation and Covariance

import pandas_datareader.data as pdr pandas can get info off the web!
import datetime

OLD: import pandas.io.data as pdweb legacy code still works, but throws a FutureWarning

Get the closing prices:

Show preview:

prices.head()

returns the first 5 rows

| | BP | CVX | ХОМ |
|------------|-------|-------|-------|
| Date | | | |
| 2010-01-04 | 46.97 | 66.17 | 60.26 |
| 2010-01-05 | 47.30 | 66.63 | 60.50 |
| 2010-01-06 | 47.55 | 66.64 | 61.02 |
| 2010-01-07 | 47.53 | 66.39 | 60.83 |
| 2010-01-08 | 47.64 | 66.51 | 60.59 |

Get the volume trades:

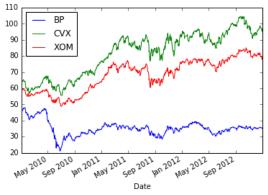
volume.head()

| | BP | CVX | XOM | |
|------------|---------|----------|----------|--|
| Date | | | | |
| 2010-01-04 | 3956100 | 10173800 | 27809100 | |
| 2010-01-05 | 4109600 | 10593700 | 30174700 | |
| 2010-01-06 | 6227900 | 11014600 | 35044700 | |
| 2010-01-07 | 4431300 | 9626900 | 27192100 | |
| 2010-01-08 | 3786100 | 5624300 | 24891800 | |

```
rets = prices.pct_change()
```

corr = rets.corr
%matplotlib inline

%matplotlib inline
prices.plot();



calculates the return using the .pct_change DataFrame method
gets the correlation of the stocks

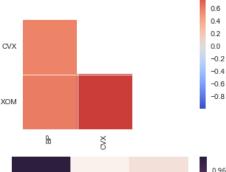
calls the plot method on the prices DataFrame

Plot the Correlation using Seaborn:

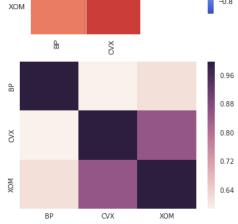
import the seaborn libraries import seaborn as sns import matplotlib.pyplot as plt import pyplot %matplotlib inline

triggers immediate matplotlib output

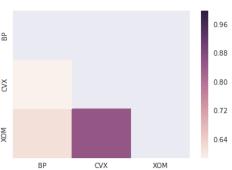
OLD: sns.corrplot(rets,annot=False,diag names=False) Returns a UserWarning: the 'corrplot' function has been deprecated in favor of 'heatmap'



NEW: sns.heatmap(rets.corr()) As expected, pretty strong correlations with each other!



Note: to mask half the plot and only show one diagonal: mask = np.zeros_like(rets.corr(), dtype=np.bool) mask[np.triu indices from(mask)] = True sns.heatmap(rets.corr(), mask=mask)



For further info visit

http://stanford.edu/~mwaskom/software/seaborn/generated/seaborn.heatmap.html

For more info:

http://dataskeptic.com/epnotes/ep79_covariance-and-correlation.php

MISSING DATA

Finding, Dropping missing data in a Series:

```
data = Series(['one','two', np.nan, 'four'])
                       data.isnull()
data
                                               data.dropna()
0
                       0 False
      one
                                               0
                                                     one
1
                       1 False
                                               1
      two
                                                     two
2
      NaN
                       2 True
                                               3
                                                    four
3
     four
                       3 False
                                               dtype: object
dtype: object
                       dtype: bool
```

Finding, Dropping missing data in a DataFrame (Be Careful!):

```
dframe = DataFrame([[1,2,3],[np.nan,5,6],[7,np.nan,9],[np.nan,np.nan,np.nan,np.nan]])
dframe.dropna(self, axis=0, how='any', thresh=None, subset=None, inplace=False)
dframe.dropna() will drop entire rows that contain at least one null value!
dframe.dropna(how='all') will drop only rows missing all data
dframe.dropna(axis=1) will drop entire columns that contain at least one null value
dframe.dropna(thresh=2) will drop rows that don't have at least 2 data points
Note that while inplace=False, none of these methods change dframe in place.
```

Filling missing data points:

```
dframe.fillna(1) fills any missing data point with a 1 dframe2.fillna({0:5,1:6,2:7,3:8}) will fill 5 in column 0, 6 in column 1, etc.
```

INDEX HIERARCHY

```
ser = Series(np.random.randn(6),index=[[1,1,1,2,2,2],['a','b','c','a','b','c']])
```

Display the series:

View the index:

```
ser.index
ser
                   MultiIndex(levels=[[1, 2], [u'a', u'b', u'c']],
  а
       -1.337299
   b
       -0.690616
                               labels=[[0, 0, 0, 1, 1, 1], [0, 1, 2, 0, 1, 2]])
   С
       1.792962
       0.457808
2
   а
       0.891199
   b
       -1.366387
   С
dtype: float64
```

Select specific subsets: Select an internal index level:

Now the cool part:

Create a DataFrame from a multilevel Series:

```
dframe = ser.unstack() returns a 2D frame, rows = (1,2), cols = (a,b,c)
```

Create a multilevel Series from a DataFrame:

```
dframe.unstack()
```

Multilevel Indexing on a DataFrame:

dframe2

| | | NY | | LA | SF |
|---|---|------|-----|-----|------|
| | | cold | hot | hot | cold |
| | 1 | 0 | 1 | 2 | 3 |
| а | 2 | 4 | 5 | 6 | 7 |
| b | 1 | 8 | 9 | 10 | 11 |
| D | 2 | 12 | 13 | 14 | 15 |

Adding names to row & column indices:

Operations on index levels:

dframe2.sum(level='Temp',axis=1)

| | Temp | cold | hot |
|--------|--------|------|-----|
| Index1 | Index2 | | |
| а | 1 | 3 | 3 |
| | 2 | 11 | 11 |
| b | 1 | 19 | 19 |
| | 2 | 27 | 27 |

Renaming columns and indices:

```
dframe.rename(index={0:'A',1:'B'}, inplace=True)
```

READING & WRITING FILES

Setting path names:

Set commonly used directories as raw data strings in the code:

```
path = r'C:\Users\Mike\Documents\Finance\'
file1 = pd.read_csv(path+'file.csv')
```

Comma Separated Value (csv) Files:

```
dframe = pd.read_csv('lect25.csv') previously saved in same directory as lecture notebook dframe = pd.read_table('lect25.csv', sep=',') can also read as table with comma delimiter pd.read_csv('lect25.csv', header=None) assigns an integer column index pd.read_csv('lect25.csv', header=None, nrows=2) takes only the first two rows dframe.to_csv('mydataout.csv') writes a DataFrame to a.csv file dframe.to_csv(sys.stdout, sep='_') lets you see the output directly without saving it dframe.to_csv(sys.stdout, columns=[0,1,2]) lets you send a specific set of columns
```

This is the pandas reader. For info on Python's csv reader/writer go to https://docs.python.org/2/library/csv.html

JSON (JavaScript Object Notation) Files:

Once you have the JSON, you can choose what info to load into a DataFrame

```
HTML Files: Note: requires beautiful soup, html5lib and lxml be installed
```

DataFrame columns: Bank Name, City, ST, CERT, Acquiring Institution, Closing Date, Updated Date, Loss Share Type, Agreement Terminated, Termination Date

```
NOTE: FOR NFL DATA, THE FOLLOWING CODE WAS NECESSARY:
```

```
website = 'http://en.wikipedia.org/wiki/NFL_win-loss_records'
nfl_frame_list = pd.read_html(website,match='Rank',header=0)
nfl_frame = nfl_frame_list[0]
```

Note: I was able to pass the argument 'Rank' without 'match=', but this may be library/parsing dependent.

For more info: http://pandas.pydata.org/pandas.pydata.

Excel Files: Note: requires xlrd and openpyxl be installed

Open an excel file as an object:

xlsfile = pd.ExcelFile('Lec_28_test.xlsx') file previously saved in notebook directory
Note: this wraps the original file into a special "ExcelFile" class object, which can then be passed to
.read_excel either sheet by sheet or all at once (performance benefit of reading original file only once)

Parse the first sheet of the excel file and set it as a DataFrame:

```
OLD: dframe = xlsfile.parse('Sheet1')
NEW: dframe = pd.read_excel(xlsfile, 'Sheet1') (xlsfile, 0) also works
dframe
```

Displays a 3x5 grid, Columns named "This is a test", "Unnamed: 1" and "Unnamed: 2". Rows indexed 0-4.

Note: Unnamed columns are assigned index positions! The tenth column would be "Unnamed: 9"

Passing sheets with the ExcelFile class as a context manager:

```
with pd.ExcelFile('path_to_file.xls') as xls:
    df1 = pd.read_excel(xls, 'Sheet1')
    df2 = pd.read_excel(xls, 'Sheet2')
```

For more info: http://pandas.pydata.org/pandas-docs/version/0.17.1/io.html#io-excel-reader

PANDAS CONCATENATE

numpy's concatenate lets you join arrays ("matrices" in the lecture): if arr1 is a 3x4 array,
np.concatenate([arr1,arr1],axis=1) creates a horizontal, 3x8 array
np.concatenate([arr1,arr1],axis=0) creates a vertical, 6x4 array (default)

in pandas, to concatenate two series:

pd.concat([ser1, ser2]) creates one long vertical series

If you concatenate two series along axis 1:

pd.concat([ser1,ser2], axis=1) the result is a DataFrame! ser1's values fall in column 0, ser2 in column 1

NOTE: if the two series being concatenated share a common index value, then

- the index value will be repeated in a vertical concatenation (axis = 0)
- the index value will appear once, and have values in both columns (axis=1)

You can add a hierarchical index using "keys":

```
concat1 = pd.concat([df1, df2, df3], keys= ['x', 'y', 'z'])
Concatenates DataFrames df1, df2 and df3 one above the other, adds an index hierarchy (x for df1, etc)
concat1['x'] will retrieve only those records belonging to 'x'
```

From: http://pandas.pydata.org/pandas-docs/stable/merging.html:

objs: list or dict of Series, DataFrame, or Panel objects. If a dict is passed, the sorted keys will be used as the keys argument, unless it is passed, in which case the values will be selected (see below)

axis: {0, 1, ...}, default 0. The axis to concatenate along

join: {'inner', 'outer'}, default 'outer'. How to handle indexes on other axis(es). Outer for union and inner for intersection

join_axes: list of Index objects. Specific indexes to use for the other n - 1 axes instead of performing inner/outer set logic

keys: sequence, default None. Construct hierarchical index using the passed keys as the outermost level If multiple levels passed, should contain tuples.

levels: list of sequences, default None. If keys passed, specific levels to use for the resulting MultiIndex. Otherwise they will be inferred from the keys

names: list, default None. Names for the levels in the resulting hierarchical index

verify_integrity: boolean, default False. Check whether the new concatenated axis contains duplicates. This can be very expensive relative to the actual data concatenation

ignore_index: boolean, default False. If True, do not use the index values on the concatenation axis. The resulting axis will be labeled 0, ..., n - 1. This is useful if you are concatenating objects where the concatenation axis does not have meaningful indexing information.

It is worth noting however, that concat (and therefore append) makes a full copy of the data, and that constantly reusing this function can create a significant performance hit. If you need to use the operation over several datasets, use a list comprehension.

MERGING DATA

See: http://pandas.pydata.org/pandas-docs/stable/merging.html#database-style-dataframe-joining-merging

Linking rows together by keys

```
dframe1 = DataFrame({'key':['X','Z','Y','Z','X','X'],'data_set_1': np.arange(6)})
dframe2 = DataFrame({'key':['Q','Z','Y','Z'],'data_set_2':[1,5,2,3]})
pd.merge(dframe1,dframe2)
```

| | data_set_1 | key | data_set_2 |
|---|------------|-----|------------|
| 0 | 1 | Z | 5 |
| 1 | 1 | Z | 3 |
| 2 | 3 | Z | 5 |
| 3 | 3 | Z | 3 |
| 4 | 2 | Υ | 2 |

Note that .merge automatically chooses overlapping columns to merge on (here it's 'key')
Where shared key values appear more than once, .merge provides every possible combination

Selecting columns and frames

pd.merge (dframe1, dframe2, on='key', how='left') Note: (...how='outer') grabs everything Returns a frame with all of data_set_1's keys, and null values (NaN) under data_set_2 where it lacked those keys From the docstring: how: {'left', 'right', 'outer', 'inner'}, default 'inner'

- * left: use only keys from left frame (SQL: left outer join)
- * right: use only keys from right frame (SQL: right outer join)
- * outer: use union of keys from both frames (SQL: full outer join)
- * inner: use intersection of keys from both frames (SQL: inner join)

Merging on multiple keys

| | key1 | key2 | left_data | right_data |
|---|------|------|-----------|------------|
| 0 | SF | one | 10 | 40 |
| 1 | SF | one | 10 | 50 |
| 2 | SF | two | 20 | NaN |
| 3 | LA | one | 30 | 60 |
| 4 | LA | two | NaN | 70 |

Handle duplicate key names with suffixes

If we had merged df_left and df_right on key1 only, there would be two columns named key2. By default, pandas sets them up as **key2 x** for left data, and **key2 y** for right data.

We can assign our own suffixes:

```
pd.merge(df_left,df_right,on='key1',suffixes=('_lefty','_righty'))
```

For more info: http://pandas.pydata.org/pandas-docs/dev/generated/pandas.DataFrame.merge.html

Merge on index (not column)

| | data | key | group_data |
|---|------|-----|------------|
| 0 | 0 | Χ | 10 |
| 3 | 3 | Χ | 10 |
| 1 | 1 | Υ | 20 |
| 4 | 4 | Υ | 20 |

This matched df_right's index values (X,Y) to df_left's "key" data, and retained df_left's index values (0-4).

This works because df_right's index contains unique values (df_left's data would never be duplicated)

From the docstring: "If joining columns on columns, the DataFrame indexes *will be ignored*. Otherwise if joining indexes on indexes or indexes on a column or columns, the index will be passed on."

Merge on multilevel index

From the docstring: left_index : boolean, default False

Use the index from the left DataFrame as the join key(s).

If it is a MultiIndex, the number of keys in the other DataFrame

(either the index or a number of columns) must match the number of levels

Merge key indicator (new in pandas 0.17.0):

```
merge(df1, df2, on='col1', how='outer', indicator=True)
Adds a categorical column merge with values "left_only", "right_only" or "both"
```

JOIN to join on indexes (row labels)

```
df_left.join(df_right) Column names must be unique. If not:

df_left.join(df_right, lsuffix='_L') use lsuffix and/or rsuffix, not suffixes

NOTE: IMO, don't pass an "on=" argument unless coping with hierarchical indices
```

COMBINING DATAFRAMES

Concatenate, Merge and Join bring two DataFrames together with tools for mapping values from each DataFrame. However, they lack the ability to *choose* between two corresponding values. That is, if dframe1 and dframe2 each have a value for row2, col2, which one wins? Can we choose a value over an empty cell?

The Long Way, using numpy's where method:

```
Series (np.where (pd.isnull (ser1), ser2, ser1), index=ser1.index) Where ser1 is null, take the value from ser2, otherwise use ser1. Apply the index from ser1.
```

The Shortcut, using pandas' combine first method:

```
ser1.combine first(ser2)
```

RESHAPING DATAFRAMES

Stack & Unstack methods

Note: stack filters out NaN by default. To avoid this use .stack (dropna=False)

PIVOTING DATAFRAMES

Consider a 12x3 dataframe. Column 1 'date' has 3 values repeating 4 times, column 2 'variable' has 4 values repeating 3 times, and column 3 'value' has random values.

```
dframe_piv = dframe.pivot('date', 'variable', 'value')
returns a 3x4 dataframe. Here, 'date' becomes a named index,
'variable' becomes the column headings, and 'value' fills the frame.
```

If we left the 'value' argument out, it would still fill frame but have 'value' shown as a label above the column headings.

Now consider a 12x7 DataFrame. Feed any three columns to the .pivot method (row, column, filler).

Alternatively, feed only two columns (row, column) and the remaining five will fill the table in turn.

For more info: https://en.wikipedia.org/wiki/Pivot_table

NOTE: the .pivot_table method on DataFrames behaves more like groupby. It aggregates values (default=mean).

DUPLICATES IN DATAFRAMES

MAPPING

Consider a DataFrame with a column called "city" and a Dictionary that matches cities up with states.

```
dframe['state'] = dframe['city'].map(state map)
```

Creates a new column called 'state' that uses keys from 'city' to grab values from the state_map dictionary.

If a city doesn't exist in the dictionary it assigns a null value.

REPLACE

```
ser1.replace('a','b') replaces 'a' with 'b' (entire entry?) ser1.replace([1,2,3],[5,6,7]) replaces 1s, 2s & 3s with 5s, 6s & 7s
```

RENAME INDEX using string operations

Change both index & column names while retaining the original:

```
dframe2 = dframe.rename(index=str.title, columns=str.lower)
```

<u>Use dictionaries to change specific names within the index and/or columns:</u> (note: keys are case sensitive!)

dframe.rename(index={dictionary}, columns={dictionary}) add inplace=True to change in place

BINNING

Using cut to design a category object

```
years = [1990,1991,1992,2008,2012,2015,1987,1969,2013,2008,1999]
decade bins = [1960, 1970, 1980, 1990, 2000, 2010, 2020] order matters!!
decade cat = pd.cut(years, decade bins)
                                            ...otherwise use sort(decade bins)
decate cat
[(1980, 1990], (1990, 2000], (1990, 2000], (2000, 2010], (2010, 2020], \ldots,
(1980, 1990], (1960, 1970], (2010, 2020], (2000, 2010], (1990, 2000)]
Length: 11
Categories (6, object): [(1960, 1970] < (1970, 1980] < (1980, 1990] < (1990,
2000] < (2000, 2010] < (2010, 2020]]
decade cat.categories
Index([u'(1960, 1970]', u'(1970, 1980]', u'(1980, 1990]', u'(1990, 2000]',
u'(2000, 2010]', u'(2010, 2020]'], dtype='object')
pd.value counts (decade cat)
                               ranks largest to smallest
(2010, 2020)
                 3
(1990, 2000]
                 3
(2000, 2010]
                 2
(1980, 1990]
                 2
(1960, 1970]
                 1
(1970, 1980]
                 0
dtype: int64
```

In the notation, () means "open" while [] means "closed/inclusive"

NOTE: As it stands, this example bins 1990 with 1985 and requires a sorted "decade bins" list. To avoid this:

```
decade_cat = pd.cut(years, sorted(decade_bins), right=False)
For some reason, if I change 1969 to 1959, the .value counts method creates a bin of (1958.9, 1987)
```

Passing data values to the cut:

Values that lie outside the bins are ignored (the cut array passes a null value) Floats are converted to integers (by chopping the decimal, not by rounding) You can't bin in alphabetical order.

OUTLIERS

Consider a 4-column data set with 1000 rows of random numbers:

np.random.seed(12345) seed the numpy generator (generates the same set of "random" numbers for each trial) dframe = DataFrame(np.random.randn(1000,4))

Grab a column from the dataset and see which values are greater than 3:

```
col = dframe[0]
col[np.abs(col)>3]
523    -3.428254
900    3.366626 in this column, rows 523 and 900 have abs values > 3
```

dframe [(np.abs (dframe) > 3) .any (1)] would grab rows where any column > 3

To cap the data at 3:

```
dframe[np.abs(dframe)>3] = np.sign(dframe) *3 multiply the sign by 3 (in place)
```

PERMUTATIONS

Create an array with a random permutation of 0,1,2,3,4:

```
array1 = np.random.permutation(5)
Note that this produces a permutation without replacement (each number appears only once in the array)
array1
array([2, 0, 4, 3, 1]) (for example)
```

Shuffle the rows of a DataFrame against this array:

```
dframe.take(array1)
```

IF ARRAY1 < DFRAME: dframe keeps only those rows represented by array1, and drops the rest.

IF ARRAY1 > DFRAME: throws an IndexError, indices are out of bounds (as opposed to filling a row with null values)

Note: if a row appears more than once in the array, it will appear more than once in the shuffled DataFrame.

Create a permutation with replacement:

```
array2 = np.random.randint(0, len(dframe), size=10)
This will make a 10-member array made up of randomly selected dframe rows.
Rows can appear more than once, and not at all.
```

Note that len(dframe) counts the number of rows, not the number of cells.

GROUPBY ON DATAFRAMES

| | dataset1 | dataset2 | k1 | k2 |
|---|----------|----------|----|-------|
| 0 | -0.45067 | -1.63403 | Χ | alpha |
| 1 | 0.268817 | 0.458236 | Χ | beta |
| 2 | 0.023818 | 0.212936 | Υ | alpha |
| 3 | -1.2282 | -1.36003 | Υ | beta |
| 4 | 0.032472 | -1.54512 | Z | alpha |

Create a SeriesGroupBy object:

```
group1 = dframe.groupby('k1') divides the DataFrame into groups around values in column 'k1'
group1
```

<pandas.core.groupby.SeriesGroupBy object at 0x000000000A6C9898>
Note that the GroupBy object is just stored data, not a DataFrame

Operations on a group return a DataFrame:

dframe.groupby('k1').mean()returns a DataFrame with index = k1, and mean values for dataset1 and dataset2

NOTE: Since 'k2' did not contain numerical values, it was dropped from the groupby.mean DataFrame Groupby.mean ignores null values. (the mean of x and null is x)

When we get to statistical analysis, is this a good way to obtain sample means to test for normal distribution?

Group data in one column 'dataset1' by another column 'k1'

```
group1 = dframe['dataset1'].groupby(dframe['k1'])
When specifying only one column from dframe, you also have to call "dframe['k1']"
```

Group by multiple column keys:

```
dframe.groupby(['k1','k2']).mean() returns a DataFrame with hierarchical index
```

Group one column by multiple column keys:

```
dataset2_group = dframe.groupby(['k1','k2'])[['dataset2']]
Note: you're calling the 'dataset2' column from the larger data set, which returns a DataFrame, because (dframe['dataset2'].groupby(dframe['k1','k2']) is not valid code.
```

Assign keys to 'dataset1' and group by them instead:

Name: dataset1, dtype: float64 Note that the output sorts by city then month alphabetically

Because we only grouped one column, groupby.mean returned a Series

Unlike the example above, this passed array indices to groupby, not column names

Other GroupBy methods:

Iterate over groups:

```
for name, group in dframe.groupby('k1'):
   print "This is the %s group" %name
   print group
   print '\n'
This is the X group
  dataset1 dataset2 k1
                            k2
0 -0.123544 1.924614 X alpha
1 -1.448666 0.477115 X beta
This is the Y group
  dataset1 dataset2 k1
                           k2
2 -1.139759 -1.378362 Y alpha
3 -0.617664 -0.105714 Y beta
This is the Z group
  dataset1 dataset2 k1
                            k2
4 -0.573748 0.409242 Z alpha
```

This operation supports multiple keys:

```
for (k1,k2),group in dframe.groupby(['k1','k2']):
    print "Key1 = %s Key2 = %s" %(k1,k2)
    print group
    print '\n' (sorts alphabetically by k1 then k2)
```

Create a dictionary from grouped data pieces:

```
group_dict = dict(list(dframe.groupby('k1')))
Here each unique member of k1 becomes a key, and its group DataFrame becomes a value!
```

For more info: http://pandas.pydata.org/pandas-docs/stable/groupby.html

Apply GroupBy using Dictionaries and Series

First, make a dataframe:

Add some null values:

```
animals.ix[1:2, ['W', 'Y']] = np.nan animals
```

| | W | X | Υ | Ζ |
|-------|-----|-----|-----|----|
| Dog | 0 | 1 2 | | 3 |
| Cat | NaN | 5 | NaN | 7 |
| Bird | 8 | 9 | 10 | 11 |
| Mouse | 12 | 13 | 14 | 15 |

Create a dictionary with "behavior" values:

```
behavior map = {'W': 'good', 'X': 'bad', 'Y': 'good', 'Z': 'bad'}
```

Group the DataFrame using the dictionary:

```
animal col = animals.groupby(behavior map, axis=1)
```

Now you can perform operations on animals based on the behavior_map dictionary values! The same thing can happen using Series.

Aggregation

The .agg(func) method lets you pass an aggregate function (like mean, max_minus_min, etc) to a GroupBy object. You can also pass string methods: grouped_frame.agg('mean')

Note: the .pivot_table method on DataFrames takes an "aggfunc=" argument (default is np.mean)

Refer to the Python Sample Code file for an example using UC Irvine's wine quality dataset on GroupBy aggregates.

Cross Tabulation

This is a special case of the .pivot table method on DataFrames

| | Sample | Animal | Intelligence |
|---|--------|--------|--------------|
| 0 | 1 | Dog | Smart |
| 1 | 2 | Dog | Smart |
| 2 | 3 | Cat | Dumb |
| 3 | 4 | Cat | Dumb |
| 4 | 5 | Dog | Dumb |
| 5 | 6 | Cat | Smart |

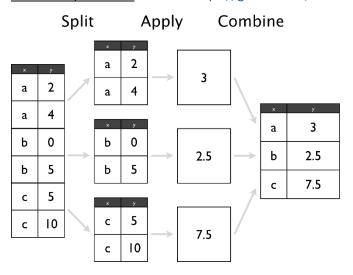
pd.crosstab (dframe.Animal, dframe.Intelligence, margins=True) margins=True adds the "All" info

| Intelligence | Dumb | Smart | All |
|--------------|------|-------|-----|
| Animal | | | |
| Cat | 2 | 1 | 3 |
| Dog | 1 | 2 | 3 |
| All | 3 | 3 | 6 |

Provides a frequency table by default, although crosstab does support aggfunc arguments as well

Split, Apply, Combine

A visual explanation: source = https://github.com/ramnathv/rblocks/issues/8



Split here is accomplished by the groupby command. If the function you're applying requires that members of the group be sorted, sort the dataframe first.

Apply can be a predefined function to be performed on each group in turn.

Combine is whatever gets returned once the apply finishes.

```
Using the same UC Irvine wine quality dataset as above (Aggregation – refer to the Python Sample Code file):
```

Build a DataFrame from the downloaded file

```
dframe wine = pd.read csv('winequality red.csv', sep=';')
```

<u>Create a function that assigns a rank to each wine based on alcohol content, with 1 being the highest alcohol content</u> def <u>ranker(df):</u>

```
df['alc_content_rank'] = np.arange(len(df)) + 1 index items 0-4 are ranked 1-5
return df
```

Sort the DataFrame by alcohol in descending order (highest at the top)

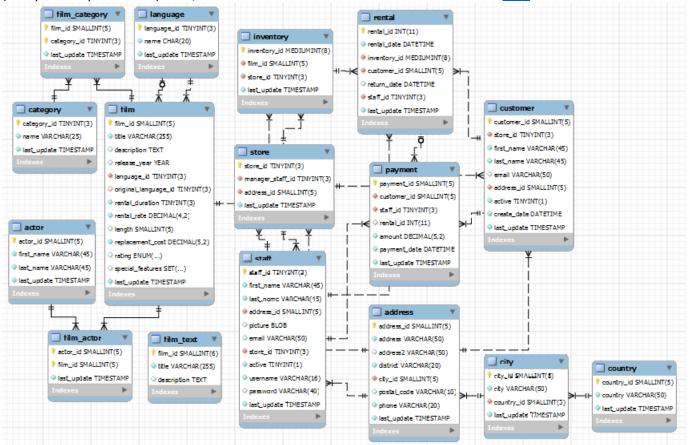
```
dframe_wine.sort_values(by='alcohol', ascending=False, inplace=True)
Group by quality and apply the ranking function
```

```
dframe_wine = dframe_wine.groupby('quality').apply(ranker)
```

dframe_wine[dframe_wine.alc_content_rank == 1].sort_values(by='quality')

SQL with Python

For this lecture we'll focus on using pandas, SQLAlchemy and the the SQLite sql browser for performing SQL queries. (Many other options exist). Also, I downloaded the sakila DVD rental database from here.



First, connect to the SQL database (using Python's built-in SQLite3 module):

```
import sqlite3
import pandas as pd
con = sqlite3.connect("sakila.db")
sql query = ''' SELECT * FROM customer '''
```

Use pandas to pass the sql query using connection from SQLite3

```
df = pd.read_sql(sql_query, con)
df.head()
```

| | custo mer_id | stor e_id | first_name | last_name | email | addre ss_id | acti ve | create_date | last_update |
|---|-----------------|--------------|------------|-----------|---|----------------|------------|--------------------------|-------------------------|
| 0 | 1 | 1 | MARY | SMITH | MARY.SMITH @sakilacustom er.org | 5 | 1 | 2/14/2006 10:04:36 PM | 9/14/2011 6:10:28 PM |
| 1 | 2 | 1 | PATRICIA | JOHNSON | PATRICIA.JO HNSON@sakil acustomer.org | 6 | 1 | 2/14/2006 10:04:36 PM | 9/14/2011 6:10:28 PM |
| 2 | 3 | 1 | LINDA | WILLIAMS | LINDA.WILLIA MS@sakilacus tomer.org | 7 | 1 | 2/14/2006 10:04:36 PM | 9/14/2011 6:10:28 PM |
| 3 | 4 | 2 | BARBARA | JONES | BARBARA.JO NES@sakilacu stomer.org | 8 | 1 | 2/14/2006 10:04:36 PM | 9/14/2011 6:10:28 PM |
| 4 | 5 | 1 | ELIZABETH | BROWN | ELIZABETH.B ROWN@sakila customer.org | 9 | 1 | 2/14/2006 10:04:36 PM | 9/14/2011 6:10:28 PM |

SQL Statements: Select, Distinct, Where, And & Or

In the statements above, we used SELECT and loaded the entire customer table

```
To save overhead, we can define a function for passing specific queries:
```

```
def sql to df(sql query):
     df = pd.read sql(sql_query, con)
    return df
query = ''' SELECT first name, last name
              FROM customer; '''
                                           rely on linebreaks & indents for improved readability
sql to df(query).head()
                                           also, SELECT, FROM etc. are not case-sensitive
Returns two specific columns
query = ''' SELECT DISTINCT country id
              FROM city'''
Returns distinct values from a specific column (not the entire record)
query = ''' SELECT *
              FROM customer
              WHERE store id = 1'''
Returns records that fit a specific criteria. Supports Boolean operators (=, <> or !=, <, >, etc.)
              WHERE first name = 'MARY'
Text values should be enclosed in single quotes (numerical values are not)
query = ''' SELECT *
```

WHERE release_year = 2006 AND rating = 'R' '''

FROM film

Supports conditional statements AND and OR

Aggregate functions include:

- AVG() Returns the average value.
- COUNT() Returns the number of rows.
- FIRST() Returns the first value.
- LAST() Returns the last value.
- MAX() Returns the largest value.
- MIN() Returns the smallest value.
- SUM() Returns the sum.

The usual syntax is:

Returns one item, with a count of the number of customers (index = 0)

Note that parentheses are required (they're optional after DISTINCT)

Wildcards are used with the LIKE operator

```
query = ''' SELECT *
    FROM customer
    WHERE first_name LIKE 'M%'; ''' use % to denote trailing text
    WHERE last_name LIKE ' ING'; ''' use_to denote leading text
```

Character Lists are enclosed in brackets

NOTE: Using [charlist] with SQLite is a little different than with other SQL formats, such as MySQL. In MySQL you would use:

```
WHERE value LIKE '[charlist]%'
In SQLite you use:
    WHERE value GLOB '[charlist]*'
query = ''' SELECT *
         FROM customer
         WHERE first_name GLOB '[AB]*'; '''
```

Returns records with any combination of A and/or B in the first name

Sorting with ORDER BY

```
query = ''' SELECT *
    FROM customer
    ORDER BY last_name ; ''' sort ascending
    ORDER BY last name DESC; ''' sort descending
```

Grouping with GROUP BY

The GROUP BY statement is used with the aggregate functions to group the results by one or more columns:

For more info: https://pymotw.com/2/sqlite3/

Web Scraping with Python

Practical considerations:

- 1.) You should check a site's terms and conditions before you scrape them.
- 2.) Space out your requests so you don't overload the site's server, doing this could get you blocked.
- 3.) Scrapers break after time web pages change their layout all the time, you'll more than likely have to rewrite your code.
- 4.) Web pages are usually inconsistent, more than likely you'll have to clean up the data after scraping it.
- 5.) Every web page and situation is different, you'll have to spend time configuring your scraper.

<u>Standard Imports</u> (BeautifulSoup4, lxml, requests)

```
from bs4 import BeautifulSoup
import requests
import pandas as pd
from pandas import Series, DataFrame
```

<u>Set the URL</u> (in this case, legislative reports from the University of California Web Page)

url = 'http://www.ucop.edu/operating-budget/budgets-and-reports/legislative-reports/2013-14-legislative-session.html'

Request content from webpage

```
result = requests.get(url)
c = result.content
```

Set as Beautiful Soup Object

soup = BeautifulSoup(c)

Use BeautifulSoup to search for the table we want to grab:

Use "inspect" in your web browser to find the particular keywords that correspond to the section you want

```
summary = soup.find("div", { 'class':'list-land', 'id':'content'})
tables = summary.find all('table')
```

Now we need to use Beautiful Soup to find the table entries.

A 'td' tag defines a standard cell in an HTML table. The 'tr' tag defines a row in an HTML table.

We'll parse through our tables object and try to find each cell using the findALL('td') method.

There are tons of options to use with findALL in beautiful soup. You can read about them here.

```
data = []
rows = tables[0].findAll('tr')
for tr in rows:
    cols = tr.findAll('td')
    for td in cols:
        text = td.find(text=True)
        print text,
        data.append(text)
```

Refer to the jupyter notebook to see output

Set up empty data list Set rows as first indexed object in tables with a row grab every HTML cell in every row

Check to see if text is in the row

Now we'll use a for loop to go through the list and grab only the cells with a pdf file in them. We also need to keep track of the index to set up the date of the report.

```
reports = [] Set up empty lists

date = []
index = 0

for item in data: Go find the pdf cells
    if 'pdf' in item:
        date.append(data[index-1]) Add the date and reports
        reports.append(item.replace(u'\xa0', u'')) Get rid of \xa0
    index += 1
```

You'll notice a line to take care of '\xa0' This is due to a unicode error that occurs if you don't do this. Web pages can be messy and inconsistent and it is very likely you'll have to do some research to take care of problems like these. Here's the link I used to solve this particular issue: StackOverflow Page

Now all that is left is to organize our data into a pandas DataFrame!

```
date = Series(date)
reports = Series(reports)
legislative_df = pd.concat([date,reports],axis=1)
legislative_df.columns = ['Date','Reports']
legislative_df.head()
Set up Dates and Reports as Series
Concatenate into a DataFrame
Set up the columns
Show the finished DataFrame (20 reports)
```

| | Date | Reports |
|---|------------|--|
| 0 | 8/1/2013 | 2013-14 (EDU 92495) Proposed Capital Outlay Pr |
| 1 | 9/1/2013 | 2014-15 (EDU 92495) Proposed Capital Outlay P |
| 2 | 11/1/2013 | Utilization of Classroom and Teaching Laborato |
| 3 | 11/1/2013 | Instruction and Research Space Summary & Analy |
| 4 | 11/15/2013 | Statewide Energy Partnership Program (pdf) |

For more info on HTML:

W3School Codecademy

Available webscraping tools:

https://import.io/
https://www.kimonolabs.com/