EE 379K: Data Science Lab

Lab 2 - 9/18/17

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1. Correlations

(a)

When given a data matrix, an easy way to tell if any two columns are correlated is to look at a scatter plot of each column against each other column. For a warm up, do this: Look at the data in DF1 in Lab2.zip. Which columns are (pairwise) correlated? Figure out how to do this with Pandas, and also how to do this with Seaborn.

In [1]: %matplotlib inline
 import numpy as np
 import matplotlib.pyplot as plt
 import pandas as pd
 from pandas.plotting import scatter_matrix
 import seaborn as sns

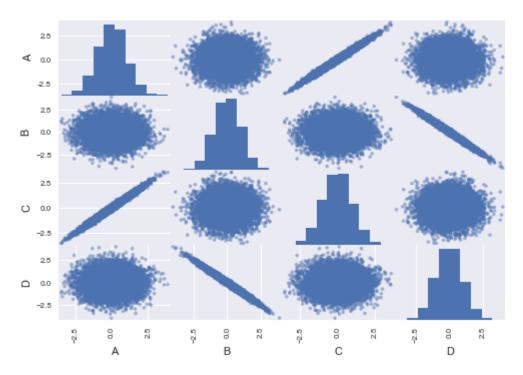
In [2]: #Read csv into pandas
 colNames = pd.Series(['A',' B','C', 'D'])
 df = pd.DataFrame.from_csv("Lab2_Data/DF1")
 df.columns = colNames;

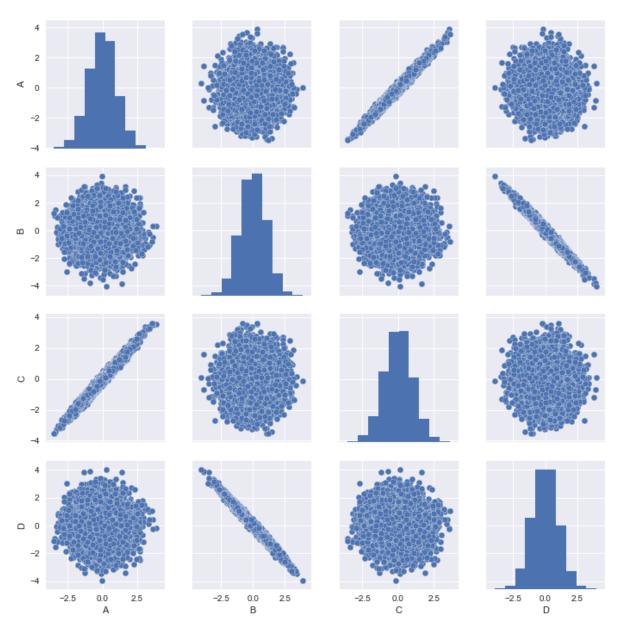
#verify data is correct
 df.head()

Out[2]:

	Α	В	С	D
0	1.038502	0.899865	0.835053	-0.971528
1	0.320455	-0.647459	0.149079	0.352593
2	0.055480	2.234771	0.271672	-2.108739
3	-0.007260	-0.524299	-0.126550	0.670827
4	-1.237390	-1.377017	-1.049932	1.342079

```
In [3]: scatter_matrix(df);
sns.pairplot(df);
sns.plt.show();
```





(b)

Compute the covariance matrix of the data. Write the explicit expression for what this is, and then use any command you like (e.g., np.cov) to compute the 4 \square 4 matrix. Explain why the numbers that you get t with the plots you got.

In [84]: df.cov()

Out[84]:

	Α	В	С	D
Α	1.001558	-0.004012	0.991624	0.004125
В	-0.004012	1.005378	-0.004099	-0.995457
С	0.991624	-0.004099	1.001589	0.004081
D	0.004125	-0.995457	0.004081	1.005168

$$\sum = egin{bmatrix} \phi_1^2 & \phi_{12} & \phi_{13} & \phi_{14} \ \phi_{21} & \phi_2^2 & \phi_{23} & \phi_{24} \ \phi_{31} & \phi_{32} & \phi_3^2 & \phi_{34} \ \phi_{41} & \phi_{42} & \phi_{43} & \phi_4^2 \end{bmatrix}$$

The covariance matrix is defined above, where the covariance between

$$Cov_{ij}$$

designates the correlation between the two columns.

In the dataset provided by DF1, we can see columns C and A have a positive correlation of 0.99 and columns B and D have a negative correlation of -0.99. This indicates that C and A and B and D are closely related

(c)

The above problem in reverse. Generate a zero-mean multivariate Gaussian random variable in 3 dimensions,

$$Z = (X1; X2; X3)$$

so that

(X1; X2)

and

(X1; X3)

are uncorre-lated, but

(X2; X3)

are correlated.

Specically: choose a covariance matrix that has the above correlations structure, and write this down. Then nd a way to generate samples from this Gaussian.

Choose one of the non-zero covariance terms (Cij, if C denotes your covariance matrix) and plot it vs the estimated covariance term, as the number of samples you use scales.

The goal is to get a visual representation of how the empirical covariance converges to the true (or family) covariance.

We can define the covariance matrix as follows:

follows:
$$\sum = \begin{bmatrix} \phi_1^2 & \phi_{12} & \phi_{13} \\ \phi_{21} & \phi_2^2 & \phi_{23} \\ \phi_{31} & \phi_{32} & \phi_3^2 \end{bmatrix}$$

In [85]: #Generate zero mean multivariate #X2 and X3 are correlated mean = [0, 0, 0]cov = [[5, 0, 0],[0, 5, 0.99], [0, 0.99, 5]] #Generate random samples from this gaussian rv = np.random.multivariate normal(mean,cov,100000) df = pd.DataFrame(rv, columns = ['X1', 'X2', 'X3']) samples = []covariances = [] #Plot vs estimated covariance term, as the number of samples you use scale for numsamples in range(0, 20000, 10): rv = np.random.multivariate_normal(mean,cov,numsamples) covar23 = pd.DataFrame(rv, columns = ['X1', 'X2', 'X3']).cov().loc['X2'] ['X3'] samples.append(numsamples) covariances.append(covar23) plt.title('Empirical Covariances') plt.xlabel('Sample Size') plt.ylabel('Covariance') plt.plot(samples,covariances)

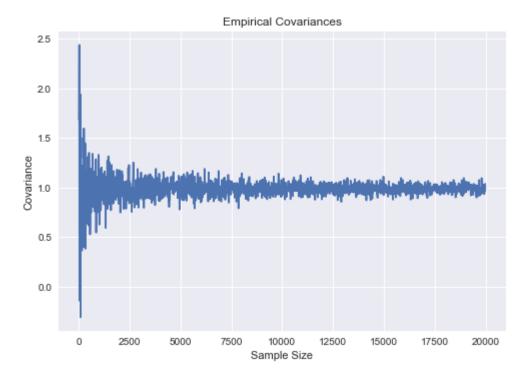
C:\Users\kevjy\Anaconda\lib\site-packages\numpy\lib\function_base.py:1110: Ru
ntimeWarning: Mean of empty slice.

avg = a.mean(axis)

C:\Users\kevjy\Anaconda\lib\site-packages\pandas\core\frame.py:5000: RuntimeW
arning: Degrees of freedom <= 0 for slice</pre>

baseCov = np.cov(mat.T)

Out[85]: [<matplotlib.lines.Line2D at 0x22b66a58>]



The above empirical covariance converges as n approaches infinity.

2. Outliers.

Consider the two-dimensional data in DF2 in Lab2.zip. Look at a scatter plot of the data. It contains two points that look like potential outliers. Which one is "more" outlying? Propose a transformation of the data that makes it clear that the point at (-1, 1) is more outlying than the point at (5.5, 5), even though the latter point is "farther away" from the nearest points. Plot the data again after performing this transformation. Provide discussion as appropriate to justify your choice of transformation. Hint: if y comes from a standard Gaussian in two dimensions (i.e., with covariance equal to the two by two identity matrix), and

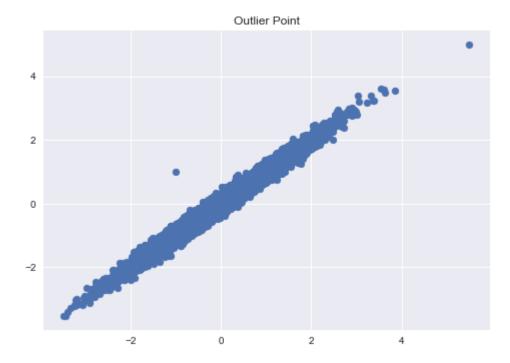
$$Q = \left[egin{array}{cc} 2 & rac{1}{2} \ rac{1}{2} & 2 \end{array}
ight]$$

what is the covariance matrix of the random variable z = Qy? If you are given z, how would you create a random Gaussian vector with covariance equal to the identity, using z?

```
In [86]: #Read from CSV file
df = pd.DataFrame.from_csv("Lab2_Data/DF2")

#plot of untransformed data
plt.scatter(df['0'],df['1'])
plt.title("Outlier Point")
```

Out[86]: <matplotlib.text.Text at 0x18b05b38>



```
In [87]: #covariance
    cov = df.cov()

#inverse covariance
    cov_inv = pd.DataFrame(np.linalg.pinv(cov.values))

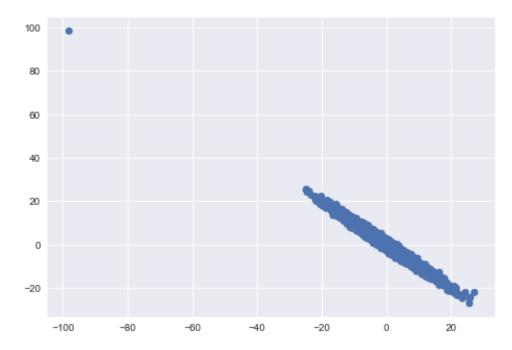
# Find Y = A^-1 * X where Y has the identity matrix
# Where Cov = A * A^T

# Transform the data and plot again
    data = pd.DataFrame(np.dot(df,cov_inv))

colnames = ['A','B']
    data.columns = colnames

plt.scatter(data['A'],data['B'])
```

Out[87]: <matplotlib.collections.PathCollection at 0x13287278>



From the graph above, we can see clearly that (-1, 1) is the clear outlier.

The Mahalanobis distance was used to scale distances so that distances along a direction where the dataset is very spread out are scaled down, and distances along directions where the dataset is tighly packed are scaled up.

As a result, a value of much higher variance will appear farther away with this transformation

Why the above statement works?

uataset.

The Mahalanobis distance uses covariance to scale distances so that distances along a direction where the dataset is very spread out are scaled down, and distances along directions where the dataset is tightly packed are scaled up. For example, in Figure $5.15(b)^{[221]}$ the Mahalanobis distance between B and A will be less than the Mahalanobis distance between C and A, whereas in Figure $5.15(c)^{[221]}$ the opposite will be true. The Mahalanobis distance is defined as

 $Mahalanobis(\mathbf{a}, \mathbf{b}) =$

(5.16)

Let's step through Equation $(5.16)^{[222]}$ bit by bit. First, this equation computes a distance between two instances **a** and **b**, each with m descriptive features. The first big term we come to in the equation is $[\mathbf{a}[1] - \mathbf{b}[1], ..., \mathbf{a}[m] - \mathbf{b}[m]]$. This is a row vector that is created by subtracting each descriptive feature value of instance **b** from the corresponding feature values of **a**. The next term in the equation, \sum^{-1} , represents the **inverse covariance matrix**²² computed across all instances in the dataset. Multiplying the difference in feature values by the **inverse covariance** matrix has two effects. First, the larger the **variance** of a feature, the less weight the difference between the values for that feature will contribute to the distance calculation. Second, the larger the correlation between two features, the less weight they contribute to the distance. The final

22 We explain covariance matrices in Section 3.5.2^[86]. The inverse covariance matrix is the matrix such that when the covariance matrix is multiplied by its inverse, the result is the identity matrix: $\sum \times \sum^{-1} = \mathbb{I}$. The identity matrix is a square matrix in which all the elements of the main diagonal are 1, and all other elements are 0. Multiplying any matrix by the identity matrix leaves the original matrix unchanged—this is the equivalent of multiplying by 1 for real numbers. So the effect of multiplying feature values be an inverse covariance matrix is to rescale the variances of all features to 1 and to set the covariance between all feature pairs to 0. Calculating the inverse of a matrix involves solving systems of linear equations and requires the use of techniques from linear algebra such as Gauss-Jordan elimination or LU decomposition. We do not cover these techniques here, but they are covered in most standard linear algebra textbooks such as Anton and Rorres (2010).

3. Even More Standard Error

 Even More Standard Error (This is to be completed only after you've completed the last written exercise below). In one of the written exercises below, you derive an expression

for what is called the *Standard Error*: where β denotes the "truth," $\hat{\beta}$ denotes the value we compute using least squares linear regression, and Z and e are as in the exercise below, you find:

$$\hat{\beta} - \beta = Ze$$
.

If we know the distribution of the noise (the distribution generating the noise vectors, e_i), then we know the distribution for the error, $(\hat{\beta} - \beta)$. This allows us to answer the question given in class: if we solve a regression and obtain value $\hat{\beta}$, how can we tell if it is statistically significant? The answer is: we compare the size of $\hat{\beta}$ to the spread introduced by the noise (i.e., the standard error), and we ask: what is the likelihood that the true $\beta = 0$, and what we observed was purely due to the noise.

If the noise is Gaussian (normal), i.e., $e_i \sim N(0, |\sigma^2|)$, and if the values of the x_i are normalized, then we expect error of the size σ/\sqrt{n} , as this is roughly the standard deviation of the expression for the error that you derive above. This means: if you have twice the data points, you should expect the error to be reduced by about 1.4 (the formula says that the standard deviation of the error would decrease by a factor of $1/\sqrt{2}$).

Compute this empirically, as follows: We will generate data for a regression problem, solve it, and see what the error is: Generate data as I did in the example from class: $x_i \sim N(0,1)$, $e_i \sim N(0,1)$. Generate y by $y_i = \beta_0 + x_i\beta + e_i$, where $\beta_0 = -3$ and $\beta = 0$. Note that since $\beta = 0$, this means that y and x are unrelated! The question we are exploring here is as follows: when we solve a regression problem, we are not going to find $\hat{\beta} = 0$ – we will find that $\hat{\beta}$ takes some other values, hopefully close to zero. How do we know if the value of $\hat{\beta}$ we get is statistically meaningful?

- By creating fresh data and each time computing β and recording β β, compute the empirical standard deviation of the error for n = 150 (the number we used in class). In class, in the exercise where I tried to find a linear regression of y vs. noise, we found β = -0.15. Given your empirical computation of the standard deviation of the error, how significant is the value -0.15?
- Now repeat the above experiment for different values of n. Plot these values, and on the same plot, plot $1/\sqrt{n}$. How is the fit?

Answer

Part 1

$$y_i = \beta_0 + x_i \beta + e_i$$

where $eta_0 = -3$ and eta = 0 $X_i = N(0,1)$

```
In [88]: def stdError(n):
             errors = []
             for i in range(1000):
                  #defining the constants
                  betanot = -3
                  beta = 0
                 Xnorm = np.random.normal(0, 1, n)
                  Enoise = np.random.normal(0,1, n)
                 #plugging into the formula
                 Y = betanot + beta*Xnorm + Enoise
                 #Formula derived in written section
                 betahat = np.dot(Xnorm, Y)/np.dot(Xnorm,Xnorm)
                 #calculate error
                  error = betahat - beta
                  errors.append(error)
             return np.std(errors)
```

```
In [89]: stdError(150)
```

Out[89]: 0.26638038764360272

Given that n = 150, the standard deviation is ~0.244 and the computed value $\hat{\beta}$ is ~0.-15.

We recognize that the calcualted value of $\hat{\beta}$ is within one standard deviation of 0.

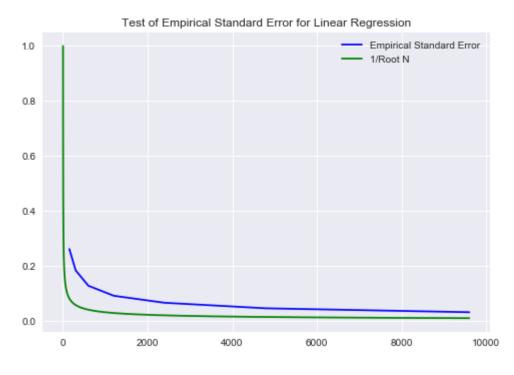
$$-0.266 < \hat{eta} < 0.266$$

the above expression indicates a good estimate. As a result, -0.15 is not significant given the standard deviation

Part 2

```
In [90]:
         #Gather data for Empirical Covariances
         yErrors = []
         numSamples = [150, 300, 600, 1200, 2400, 4800, 9600]
         for i in range(len(numSamples)):
             yErrors.append(stdError(numSamples[i]))
         #Gather data for 1/root(i)
         xSqrt = []
         ySqrt = []
         for i in range(1,9600):
             xSqrt.append(i)
             ySqrt.append(1/np.sqrt(i))
         #plot
         plt.plot(numSamples, yErrors, color = 'b', label = 'Empirical Standard Error')
         plt.plot(xSqrt, ySqrt, color = 'g', label = '1/Root N')
         plt.legend()
         plt.title('Test of Empirical Standard Error for Linear Regression')
```

Out[90]: <matplotlib.text.Text at 0x10cea278>



4. Names and Frequencies

The goal of this exercise is for you to get more experience with Pandas, and to get a chance to explore a cool data set. Download the file Names.zip from Canvas. This contains the frequency of all names that appeared more than 5 times on a social security application from 1880 through 2015.

- Write a program that on input k and XXXX, returns the top k names from year XXXX.
- Write a program that on input Name returns the frequency for men and women of the name Name.
- It could be that names are more diverse now than they were in 1880, so that a name may be relatively
 the most popular, though its frequency may have been decreasing over the years. Modify the above to
 return the relative frequency.
- Find all the names that used to be more popular for one gender, but then became more popular for another gender.
- (Optional) Find something cool about this data set.

```
In [8]: k = int(input("Enter the number of top names: "))
    year = int(input("Enter the year would like to search: "))
    while(year < 1880 or year > 2015):
        year = int(input("Please enter another year between 1880 and 2015"))
    df = pd.read_csv("Names/yob"+str(year)+".txt", sep=',', names=['name', 'gende r', 'freq'])

#print
print(df.sort_values('freq', ascending=False).head(k))
```

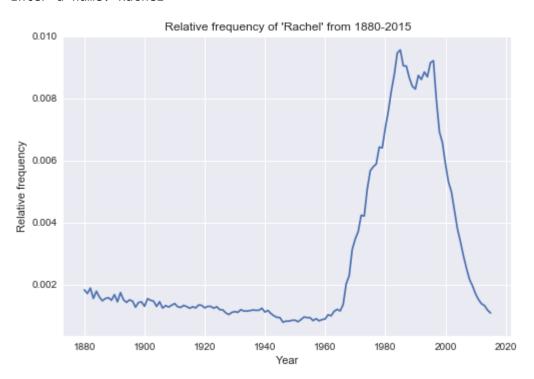
```
Enter the number of top names: 5
Enter the year would like to search: 1999
         name gender
                       freq
                   M 35346
16938
        Jacob
16939 Michael
                   M 33906
                   M 30417
16940 Matthew
16941
       Joshua
                   M 27254
        Emily
                   F 26537
```

```
name = raw_input("Enter the name: ")
filenames = ['Names/yob%s.txt' % year for year in range(1880, 2016)]
dfs = []
for filename in filenames:
    dfs.append(pd.read_csv(filename, sep=',', names=['name', 'gender',
'freq']))
name_count_female = 0
name_count_male = 0
for df in dfs:
    df name = df.loc[df['name'] == name]
    females = df_name.loc[df_name['gender'] == 'F'].freq
    males = df_name.loc[df_name['gender'] == 'M'].freq
    if len(females) > 0:
        name_count_female += females.values[0]
    if len(males) > 0:
        name_count_male += males.values[0]
print 'Female: %d, Male: %d' % (name_count_female, name_count_male)
```

Enter the name: James Female: 23215, Male: 5120990

```
In [8]: name = raw input("Enter a name: ")
        filenames = ['Names/yob%s.txt' % year for year in range(1880, 2016)]
        dfs = []
        for filename in filenames:
            dfs.append(pd.read csv(filename, sep=',', names=['name', 'gender',
         'freq']))
        df freq = pd.DataFrame()
        vear = 1880
        name count female = 0
        name count male = 0
        for df in dfs:
            df_name = df.loc[df['name'] == name]
            females = df name.loc[df name['gender'] == 'F'].freq
            males = df name.loc[df name['gender'] == 'M'].freq
            if len(females) > 0:
                 name_count_female = females.values[0]
            if len(males) > 0:
                name_count_male = males.values[0]
            #get specific name's male and female count
            df female = df.loc[df['gender'] == 'F'].sum()
            df_male = df.loc[df['gender'] == 'M'].sum()
            total count female = df female[2]
            total count male = df male[2]
            #get total male and female count
            rel freq female = name count female / float(total count female)
            rel_freq_male = name_count_male / float(total_count_male)
            #calculate relative freq of name
            df_name_freq = pd.DataFrame({'Year' : year, 'F_freq' : [rel_freq_female],
         'M freq' : [rel freq male]})
            df freq = df freq.append(df name freq)
            year += 1
            #reset
            name count female = 0
            name count male =0
        plt.title("Relative frequency of '"+ name + "' from 1880-2015")
        plt.xlabel("Year")
        plt.vlabel("Relative frequency")
        plt.plot(df freq.Year, (df freq.M freq + df freq.F freq))
        plt.show()
        print(df_freq)
```

Enter a name: Rachel



F freq M freq Year 0 0.001824 0.000000 1880 0 0.001707 0.000000 1881 0 0.001882 0.000000 1882 0 0.001549 0.000000 1883 0 0.001783 0.000000 1884 0 0.001593 0.000000 1885 0 0.001474 0.000000 1886 0 0.001548 0.000000 1887 0 0.001573 0.000000 1888 0 0.001491 0.000000 1889 0 0.001670 0.000000 1890 0 0.001439 0.000000 1891 0 0.001738 0.000000 1892 0 0.001508 0.000000 1893 0 0.001422 0.000000 1894 0 0.001502 0.000000 1895 0 0.000000 0.001463 1896 0 0.001268 0.000000 1897 0 0.001418 0.000000 1898 0 0.001390 0.000047 1899 0 0.001264 0.000033 1900 0 0.001542 0.000000 1901 0.000000 0 0.001488 1902 0 0.001462 0.000000 1903 0 0.001293 0.000000 1904 0 0.001444 0.000000 1905 0 0.001239 0.000000 1906 0 0.001318 0.000000 1907 0 0.001271 0.000000 1908 0 0.001331 0.000000 1909 0 0.009010 0.000031 1986 0 0.009000 0.000034 1987 0 0.008620 0.000032 1988 0 0.008331 0.000053 1989 1990 0 0.008275 0.000019 0 0.008718 0.000015 1991 0 0.008589 0.000013 1992 0 0.008829 0.000013 1993 0 0.008674 0.000013 1994 0 0.009125 0.000015 1995 0 0.009193 0.000013 1996 0 0.007921 0.000012 1997 0 0.006905 0.000009 1998 0.006554 0.000009 1999 0 0 0.005878 0.000006 2000 0 0.005297 0.000004 2001 0 0.004969 0.000008 2002 0 0.004369 0.000007 2003 0 0.000017 2004 0.003767 0 0.003364 0.000007 2005 0 0.002909 0.000003 2006 0 0.002511 0.000003 2007 0 0.002160 0.000002 2008 0 0.001952 0.000003 2009 0.001698 0.000000 2010

```
0.001512 0.000000 2011
        0
            0.001372 0.000004 2012
            0.001318 0.000000 2013
            0.001168 0.000003
                                2014
            0.001080 0.000000 2015
        [136 rows x 3 columns]
In [2]: filenames = ['Names/yob%s.txt' % year for year in range(1880, 2016)]
        dfs = []
        initialyear = 1880
        for filename in filenames:
            currentDF = pd.read csv(filename, sep=',', names=['name', 'gender',
            currentDF['Year'] = pd.Series(initialyear, index=currentDF.index)
            dfs.append(currentDF)
            initialyear += 1
        bigDF = pd.concat(dfs)
        #find all values that appear more than the 136 (expected number names if it wa
        s just one gender)
        dfpiv = pd.pivot_table(bigDF,index=['name'],aggfunc='count')
        multiNamePiv = dfpiv[dfpiv['freq'] > (2016-1880)]
        print multiNamePiv.index
        Index([u'Aaron', u'Abbie', u'Abby', u'Abel', u'Abigail', u'Abraham', u'Ada',
               u'Adair', u'Adam', u'Addie',
               u'Yancy', u'Yolanda', u'Young', u'Yvette', u'Yvonne', u'Zachary',
               u'Zane', u'Zelma', u'Zoe', u'Zola'],
              dtype='object', name=u'name', length=1672)
```

In [76]: #WARNING: Long Running Code femaleToMale = []; maleToFemale = []; for name in multiNamePiv.index: male1948 = bigDF[(bigDF['name'] == name) & (bigDF['Year'] < 1948) &</pre> (bigDF['gender'] == 'M')].freq.sum() female1948 = bigDF[(bigDF['name'] == name) & (bigDF['Year'] < 1948) & (big</pre> DF['gender'] == 'F')].freq.sum() male2015 = bigDF[(bigDF['name'] == name) & (bigDF['Year'] > 1948) & (bigDF['gender'] == 'M')].freq.sum() female2015 = bigDF[(bigDF['name'] == name) & (bigDF['Year'] > 1948) & (big DF['gender'] == 'F')].freq.sum() if(male1948 > female1948 and male2015 < female2015):</pre> femaleToMale.append(name) print "Male name before 1948 but Female name after " + name if(female1948 > male1948 and female2015 < male2015):</pre> maleToFemale.append(name) print "Female name before 1948 but Male name afterwards " + name

Male name before 1948 but Female name after Allison Female name before 1948 but Male name afterwards Alpha Male name before 1948 but Female name after Alva Male name before 1948 but Female name after Arden Female name before 1948 but Male name afterwards Artie Male name before 1948 but Female name after Ashley Male name before 1948 but Female name after Aubrey Female name before 1948 but Male name afterwards Audie Male name before 1948 but Female name after Avery Male name before 1948 but Female name after Blair Male name before 1948 but Female name after Carlie Male name before 1948 but Female name after Charley Male name before 1948 but Female name after Clair Male name before 1948 but Female name after Courtney Male name before 1948 but Female name after Dee Male name before 1948 but Female name after Elisha Female name before 1948 but Male name afterwards Frankie Male name before 1948 but Female name after Gale Male name before 1948 but Female name after Hollie Male name before 1948 but Female name after Ivev Male name before 1948 but Female name after Ivory Male name before 1948 but Female name after Jackie Female name before 1948 but Male name afterwards Jessie Male name before 1948 but Female name after Jodie Male name before 1948 but Female name after Kelly Male name before 1948 but Female name after Lacy Male name before 1948 but Female name after Lavern Female name before 1948 but Male name afterwards Lavon Male name before 1948 but Female name after Leigh Female name before 1948 but Male name afterwards Lennie Male name before 1948 but Female name after Lesley Male name before 1948 but Female name after Leslie Male name before 1948 but Female name after Lindsay Male name before 1948 but Female name after Lindsey Female name before 1948 but Male name afterwards Maxie Male name before 1948 but Female name after Morgan Female name before 1948 but Male name afterwards Ocie Female name before 1948 but Male name afterwards Ollie Male name before 1948 but Female name after Paris Female name before 1948 but Male name afterwards Pat Female name before 1948 but Male name afterwards Robbie Male name before 1948 but Female name after Rosario Male name before 1948 but Female name after Stacy Male name before 1948 but Female name after Sydney Female name before 1948 but Male name afterwards Toby Male name before 1948 but Female name after Tracy Female name before 1948 but Male name afterwards Trinidad

Male name before 1948 but Female name after Allison Female name before 1948 but Male name afterwards Alpha Male name before 1948 but Female name after Alva Male name before 1948 but Female name after Arden Female name before 1948 but Male name afterwards Artie Male name before 1948 but Female name after Ashley Male name before 1948 but Female name after Aubrey Female name before 1948 but Male name afterwards Audie Male name before 1948 but Female name after Avery Male name before 1948 but Female name after Blair Male name before 1948 but Female name after Carlie Male name before 1948 but Female name after Charley Male name before 1948 but Female name after Clair Male name before 1948 but Female name after Courtney Male name before 1948 but Female name after Dee Male name before 1948 but Female name after Elisha Female name before 1948 but Male name afterwards Frankie Male name before 1948 but Female name after Gale Male name before 1948 but Female name after Hollie Male name before 1948 but Female name after Ivey Male name before 1948 but Female name after Ivory Male name before 1948 but Female name after Jackie Female name before 1948 but Male name afterwards Jessie Male name before 1948 but Female name after Jodie Male name before 1948 but Female name after Kelly Male name before 1948 but Female name after Lacy Male name before 1948 but Female name after Lavern Female name before 1948 but Male name afterwards Lavon Male name before 1948 but Female name after Leigh Female name before 1948 but Male name afterwards Lennie Male name before 1948 but Female name after Lesley Male name before 1948 but Female name after Leslie Male name before 1948 but Female name after Lindsay Male name before 1948 but Female name after Lindsey Female name before 1948 but Male name afterwards Maxie Male name before 1948 but Female name after Morgan Female name before 1948 but Male name afterwards Ocie Female name before 1948 but Male name afterwards Ollie Male name before 1948 but Female name after Paris Female name before 1948 but Male name afterwards Pat Female name before 1948 but Male name afterwards Robbie Male name before 1948 but Female name after Rosario Male name before 1948 but Female name after Stacy Male name before 1948 but Female name after Sydney Female name before 1948 but Male name afterwards Toby Male name before 1948 but Female name after Tracy Female name before 1948 but Male name afterwards Trinidad

```
In [78]:
         #this code works, but take's forever to analyze. commented out
         runningCountMales = 0
         runningCountFemales = 0
         moreFemaleThanMale =False;
         swapHappened = False;
         femaleToMaleList = []
         maleToFemaleList = []
         for name in multiNamePiv.index:
             for year in range(1880,2016):
                 df name = biqDF.loc[(biqDF['name'] == name) & (biqDF['Year'] == year)]
                 females = df_name.loc[df_name['gender'] == 'F'].freq
                 males = df name.loc[df name['gender'] == 'M'].freq
             if (len(females) > 0 and len(males) > 0):
                  if(females.values[0] > males.values[0]):
                      moreFemaleThanMale = True
                 else:
                      moreFemaleThanMale = False
                  if(moreFemaleThanMale and not swapHappened and males.values[0] > femal
         es.values[0]):
                      swapHappened = True;
                      femaleToMaleList.append(name);
                  if(not moreFemaleThanMale and not swapHappened and females.values[0] >
          males.values[0]):
                      swapHapened = True;
                      maleToFemaleList.append(name);
         print femaleToMaleList
```

"\nrunningCountMales = 0\nrunningCountFemales = 0\n\nmoreFemaleThanMale =Fals Out[78]: e;\nswapHappened = False;\n\nfemaleToMaleList = []\nmaleToFemaleList = []\nfo r name in multiNamePiv.index:\n for year in range(1880,2016):\n df_ name = bigDF.loc[(bigDF['name'] == name) & (bigDF['Year'] == year)]\n females = df name.loc[df name['gender'] == 'F'].freq\n males = df name.loc[df_name['gender'] == 'M'].freq\n if (len(females) > 0 and len(males) if(females.values[0] > males.values[0]):\n > 0):\n moreFema leThanMale = True\n else:\n moreFemaleThanMale = False\n if(moreFemaleThanMale and not swapHappened and males.values[0] > female s.values[0]):\n swapHappened = True;\n femaleToMaleLis t.append(name);\n if(not moreFemaleThanMale and not swapHappened and f emales.values[0] > males.values[0]):\n swapHapened = True;\n maleToFemaleList.append(name);\nprint femaleToMaleList

5. Visualization Tools and Missing/Hidden Values.

Visualization is important both for exploring the data, as well as for explaining what you have done. There are a huge number of such tools now available. This exercise walks through various functionalities of matplotlib and pandas.

- The first part of this exercise was created by Dataquest. Run through the commands given in this tutorial: https://www.dataquest.io/blog/matplotlib-tutorial/ (https://www.dataquest.io/blog/matplotlib-tutorial/) and under- stand the code.
- Suppose that you would now like to plot some of the results by state. As you will see, the state information is sometimes missing, and other times it comes in varying forms. Figure out how to aggregate the results by state. The challenge here: how many of the tweets can you (correctly) assign to a state? Note: depending on how well you want to do (i.e., how many tweets you want to correctly assign to their state), this is not an easy problem!

```
In [10]: tweets = pd.read_csv("Lab2_data/tweets.csv")
```

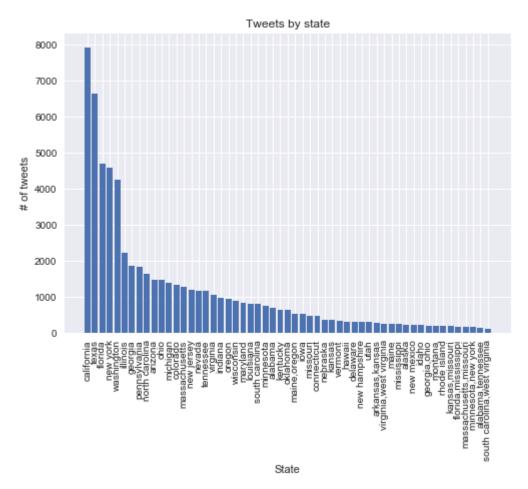
```
In [11]: | def get_state(row):
             state = []
             text = row["user location"].lower()
             if "alabama" in text or r"\bal\b" in text or "birmingham" in text or "mobi
         le" in text or "huntsville" in text:
                 state.append("alabama")
             if "alaska" in text or r"\bak\b" in text or "anchorage" in text or "fairba
         nks" in text or "juneau" in text:
                 state.append("alaska")
             if "arizona" in text or r"\baz\b" in text or "phoenix" in text or "tuscon"
          in text or "mesa" in text:
                 state.append("arizona")
             if "arkansas" in text or r"\bar\b" in text or "little rock" in text or "fo
         rt smith" in text or "fayetteville" in text:
                 state.append("arkansas")
             if "california" in text or r"\bca\b" in text or "los angeles" in text or
         "san diego" in text or "san jose" in text or "san francisco" in text:
                 state.append("california")
             if "colorado" in text or r"\bco\b" in text or "enver" in text or "colorado
          springs" in text or "aurora" in text:
                 state.append("colorado")
             if "connecticut" in text or r"\bct\b" in text or "bridgeport" in text or
         "new haven" in text or "hartford" in text:
                 state.append("connecticut")
             if "delaware" in text or r"\bde\b" in text or "wilmington" in text or "dov
         er" in text or "newark" in text:
                 state.append("delaware")
             if "florida" in text or r"\bfl\b" in text or "jacksonville" in text or "mi
         ami" in text or "tampa" in text:
                 state.append("florida")
             if "georgia" in text or r"\bga\b" in text or "atlanta" in text or "august
         a" in text or "columbus" in text:
                 state.append("georgia")
             if "hawaii" in text or r"\bhi\b" in text or "honolulu" in text or "hilo" i
```

Lab2

Lab2

n text or "kailua" in text: state.append("hawaii") if "idaho" in text or r"\bid\b" in text or "boise" in text or "nampa" in t ext or "idaho falls" in text: state.append("idaho") if "illinois" in text or r"\bil\b" in text or "chicago" in text or "auror a" in text or "rockford" in text: state.append("illinois") $\textbf{if} \ \texttt{"indiana"} \ \textbf{in} \ \texttt{text} \ \textbf{or} \ \texttt{r"} \\ \texttt{bin} \\ \texttt{b"} \ \textbf{in} \ \texttt{text} \ \textbf{or} \ \texttt{"indianapolis"} \ \textbf{in} \ \texttt{text} \ \textbf{or} \ \texttt{"fo} \\$ rt wayne" in text or "evansville" in text: state.append("indiana") if "iowa" in text or r"\bia\b" in text or "des moines" in text or "cedar r apids" in text or "davenport" in text: state.append("iowa") if "kansas" in text or r"\bks\b" in text or "wichita" in text or "overland park" in text or "kansas city" in text: state.append("kansas") if "kentucky" in text or r"\ky\b" in text or "louisville" in text or "lexi ngton" in text or "owensboro" in text: state.append("kentucky") if "louisiana" in text or r"\bla\b" in text or "new orleans" in text or "s hreveport" in text or "baton rouge" in text: state.append("louisiana") if "maine" in text or r"\bme\b" in text or "portland" in text or "lewisto n" in text or "bangor" in text: state.append("maine") if "maryland" in text or r"\bmd\b" in text or "baltimore" in text or "fred erick" in text or "gaithersburg" in text: state.append("maryland") if "massachusetts" in text or r"\bma\b" in text or "boston" in text or "wo rcester" in text or "springfield" in text: state.append("massachusetts") if "michigan" in text or r"\bmi\b" in text or "detroit" in text or "grand rapids" in text or "warren" in text: state.append("michigan") if "minnesota" in text or r"\bmn\b" in text or "minneapolis" in text or "s aint paul" in text or "rochester" in text: state.append("minnesota") if "mississippi" in text or r"\bms\b" in text or "jackson" in text or "gul fport" in text or "biloxi" in text: state.append("mississippi") if "missouri" in text or r"\bmo\b" in text or "kansas city" in text or "sa int louis" in text or "springfield" in text: state.append("missouri") if "montana" in text or r"\bmt\b" in text or "bilings" in text or "missoul a" in text or "great falls" in text: state.append("montana") if "nebraska" in text or r"\bne\b" in text or "omaha" in text or "lincoln" in text or "bellevue" in text: state.append("nebraska") if "nevada" in text or r"\bnv\b" in text or "las vegas" in text or "reno" in text or "henderson" in text: state.append("nevada") if "new hampshire" in text or r"\bnh\b" in text or "machester" in text or "nashua" in text or "concord" in text: state.append("new hampshire") if "new jersey" in text or r"\bnj\b" in text or "newark" in text or "jerse

```
y city" in text or "paterson" in text:
        state.append("new jersey")
   if "new mexico" in text or r"\bnm\b" in text or "albuquerque" in text or
"las cruces" in text or "rio rancho" in text:
        state.append("new mexico")
   if "new york" in text or r"\bny\b" in text or "new york city" in text or
"buffalo" in text or "rochester" in text:
       state.append("new york")
   if "north carolina" in text or r"\bnc\b" in text or "charlotte" in text or
 "raleigh" in text or "greensboro" in text:
        state.append("north carolina")
   if "north dakota" in text or r"\bnd\b" in text or "fargo" in text or "bism
arck" in text or "grand forks" in text:
        state.append("north dakota")
   if "ohio" in text or r"\boh\b" in text or "columbus" in text or "clevelan
d" in text or "cincinnati" in text:
        state.append("ohio")
   if "oklahoma" in text or r"\bok\b" in text or "oklahoma city" in text or
"tulsa" in text or "norman" in text:
        state.append("oklahoma")
   if "oregon" in text or r"\bor\b" in text or "portland" in text or "salem"
in text or "eugene" in text:
        state.append("oregon")
   if "pennsylvania" in text or r"\bpa\b" in text or "philadelphia" in text o
r "pittsburgh" in text or "allentown" in text:
        state.append("pennsylvania")
   if "rhode island" in text or r"\bri\b" in text or "providence" in text or
"warwick" in text or "cranston" in text:
        state.append("rhode island")
   if "south carolina" in text or r"\bsc\b" in text or "charleston" in text o
r "columbia" in text or "north charleston" in text:
        state.append("south carolina")
   if "south dakota" in text or r"\bsc\b" in text or "sioux falls" in text or
 "rapid city" in text or "aberdeen" in text:
        state.append("south dakota")
   if "tennessee" in text or r"\btn\b" in text or "memphis" in text or "mobil
e" in text or "huntsville" in text:
        state.append("tennessee")
   if "texas" in text or r"\btx\b" in text or "austin" in text or "houston" i
n text or "dallas" in text:
        state.append("texas")
   if "utah" in text or r"\but\b" in text or "salt lake city" in text or "wes
t valley city" in text or "provo" in text:
        state.append("utah")
   if "vermont" in text or r"\bvt\b" in text or "burlington" in text or "sout
h burlington" in text or "rutland" in text:
        state.append("vermont")
   if "virginia" in text or r"\bva\b" in text or "virginia beach" in text or
"norfolk" in text or "chesapeake" in text:
        state.append("virginia")
   if "washington" in text or r"\bwa\b" in text or "seattle" in text or "spok
ane" in text or "tacoma" in text:
        state.append("washington")
   if "west virginia" in text or r"\bwv\b" in text or "charleston" in text or
 "huntington" in text or "parkersburg" in text:
        state.append("west virginia")
   if "wisconsin" in text or r"\bwi\b" in text or "milwaukee" in text or "mad
```



california texas florida new york	7926 6630 4705 4581
washington	4261
illinois	2222
georgia	1856
pennsylvania	1815
north carolina	1638
arizona	1469
ohio	1459
michigan	1390
colorado	1320
massachusetts	1270
new jersey	1200
nevada	1164
tennessee virginia	1161 1046
indiana	958
oregon	937
wisconsin	875
maryland	821
louisiana	799
south carolina	798
minnesota	743
alabama	697
kentucky	641
oklahoma	627
maine,oregon	516
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arkansas,kansas,oklahoma,texas1illinois,tennessee1illinois,new york1illinois,massachusetts1

Name: state, Length: 312, dtype: int64

1. Standard Error:

It is important to develop an intuition for how much error we should expect" when we solve a particular statistical problem. As the number of sample increase, we should expect the error to decrease. But by how much? In the rst lab, you generated samples from a univariate (Problem 3) and multivariate (Problem 4) Gaussian with given parameters, and then you were asked to estimate those parameters from the data you generated. In this exercise, we derive explicitly the relationship that you (should have) observed doing those exercises.

• Suppose $Z=N(\mu;\theta^2)$, i.e., Z is a univariate Gaussian (a.k.a. normal) random variable with mean μ and variance θ^2 . Suppose that you see n samples from Z, i.e., you see data z_1,\ldots,z_n . Let

$$z_{avg} = \sum_{i=1}^n rac{z_i}{n}$$

denote the sample mean.

We want to answer: how close is z_{ava} to μ ?

Note that z_{avq} is a random variable so we need to quantify in a probabilistic way how close z_{avq} is to μ .

- Suppose Z ~ N(0,1). This is also called a standard normal random variable. For n = 10,000, compute the probability that z_{avg} > 0.1, z_{avg} > 0.01, and z_{avg} > 0.01
- Now for the general case: suppose $Z=N(\mu;\theta^2)$, and for general n, compute the probability that $z_{avg}>n^{\frac{-1}{3}}$, $z_{avg}>n^{\frac{-1}{2}}$, and $z_{avg}>n^{\frac{-2}{3}}$. For your calculations, you can let n scale if that makes things easier.

Answer

$$P(z_{avg}>c)$$

Re-arrange variables

$$P(S_n > c * n)$$

Using Central Limit Theorem

$$P(rac{S_n}{\sqrt{n} * heta} > rac{0.1 * n}{\sqrt{n} * heta}) \ P(Z_n > 0.1 * \sqrt{n})$$

Standard Normal

$$1 - P(Z_n < c * \sqrt{10000})$$

Replace:

c = 0.1

$$P(z_{avg} > 0.1) \ 1 - P(Z_n < 10) = 0$$

c = 0.01

$$P(z_{avg} > 0.01) \ 1 - P(Z_n < 1) = 0.16$$

c = 0.001

$$P(z_{avg}>0.001) \ 1-P(Z_n<0.1)=0.46$$

General Case

$$Z=N(\mu; heta^2) \ P(Z_{avg}>c)$$

Using Central Limit Theorem

$$P(rac{S_n - (n*\mu)}{\sqrt{n}* heta} > rac{c*n - (n*\mu)}{\sqrt{n}* heta})$$

For $c = n^{-1/3}$

$$egin{aligned} P(Z_{avg} - \mu > n^{-1/3}) \ \hat{x} &= n^{-1/3} + \mu \ Z &= rac{n^{-1/3} + \mu - \mu}{\delta / \sqrt(n)} \ Z &= rac{n^{-1/3}}{\delta} n^{1/2} = rac{n^{1/6}}{\delta} \ &= P(Z < rac{-n^{1/6}}{\delta}) \end{aligned}$$

For $c=n^{-1/2}$

$$egin{aligned} P(Z_{avg} - \mu > n^{-1/2}) \ \hat{x} &= n^{-1/2} + \mu \ Z &= rac{n^{-1/2} + \mu - \mu}{\delta / \sqrt(n)} \ Z &= rac{n^{-1/2}}{\delta} n^{1/2} = rac{n^{-1/2}}{\delta} \ &= P(Z < rac{-1}{\delta}) \end{aligned}$$

For $c=n^{-2/3}$

$$egin{aligned} P(Z_{avg} - \mu > n^{-2/3}) \ \hat{x} &= n^{-2/3} + \mu \ Z &= rac{n^{-2/3} + \mu - \mu}{\delta / \sqrt(n)} \ Z &= rac{n^{-2/3}}{\delta} n^{1/2} = rac{n^{-1/6}}{\delta} \ &= P(Z < rac{-n^{-1/6}}{\delta}) \end{aligned}$$

2. More Standard Error Consider a one dimensional regression problem, where the offset is zero. Thus, we are trying to fit a function of the form $h(x) = x \cdot \beta$. Suppose that the truth is a noisy version of this – that is, the true model according to which data are generated is:

$$y_i = x_i \cdot \beta + e_i.$$

Everything in the above equation is a scalar, i.e., $y_i, x_i, \beta, e_i \in \mathbb{R}$. Here, e_i represents independent noise that is not modeled by the linear relationship.

When we have n data points, the least squares objective reads:

$$\min_{\beta} : \frac{1}{n} \sum_{i=1}^{n} (x_i \beta - y_i)^2.$$

Show that this is a quadratic function in β , that is, if we expand it, it has the form

$$A\beta^2 + B\beta + C.$$

- Compute A, B, and C explicity, i.e., as explicit functions of the data, $\{x_i, y_i\}$. Note that these should not be functions of β . Show that $A \geq 0$ regardless of the values of the data.
- Since A ≥ 0, this is a quadratic function whose graph opens up. This means that it is convex, and therefore the solution is characterized as the solution obtained by setting the first derivative (w.r.t. β) equal to zero. Do this, and therefore explicitly solve for the solution β̂. This is the one-dimensional form of what is known as the normal equations. Hint: we did this problem in class.
- Now using the one dimensional expression from the second part, and plugging in the relationship $y_i = x_i \cdot \beta + e_i$, write

$$\hat{\beta} = \beta + Z\mathbf{e},$$

where e denotes the vector of all the errors, e_i , added in each stage, and where Z is a matrix of appropriate dimension. What is Z, explicitly?

(Bonus) Repeat the last two questions in the general case. That is, derive the normal equations and the standard error for the general (vector) case, where our model is

$$y_i = x_i^{\top} \beta + e_i,$$

where now $x_i, \beta \in \mathbb{R}^p$, and $x_i^{\top} \beta$ denotes the dot product.

Answer:

Part 1

When we have n data points, the least square objective reads:

$$mineta = rac{1}{n} \sum_{i=1}^n (x_ieta - y_i)^2$$

We can show that eta is a quadratic formula of the form $Aeta^2+Beta+C$

Multiplying out

$$mineta = rac{1}{n}[\sum_{i=1}^n (x_i^2eta^2 - 2X_ieta y_i + y_i^2]$$

Expand out

$$mineta = eta^2 rac{1}{n} \sum_{i=1}^n x_i^2 - eta rac{2}{n} \sum_{i=1}^n 2x_i y_i + rac{1}{n} \sum_{i=1}^n y_i^2$$

where

$$A = rac{1}{n} \sum_{i=1}^n x_i^2 \ B = rac{2}{n} \sum_{i=1}^n 2x_i y_i \ C = rac{1}{n} \sum_{i=1}^n y_i^2$$

Notice A is a quadratic and $A\geq 0$

Part 2

Solve for β

$$egin{aligned} rac{d}{deta}(rac{1}{n}\sum_{i=1}^n(X_ieta-y_i)^2) &= rac{2}{n}\sum_{i=1}^n(X_ieta-y)X_i) = 0 \ &\sum_{i=1}^n(X_i^2eta-X_iY_i) = 0 \ &\therefore eta &= rac{\sum_{i=1}^nX_iY_i}{\sum_{i=1}^nX_i^2} \end{aligned}$$