# Lab 2

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# **Programming Questions**

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

### 1. Correlations

#### **Pandas**

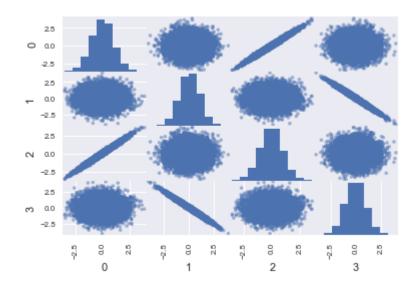
```
In [2]: path = 'Data\DF1'
df=pd.read_csv(path,index_col=0,na_values='?')
df.head()
```

Out[2]:

		0	1	2	3
	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]: from pandas.plotting import scatter_matrix
# plot each column against each other using pandas
scatter_matrix(df)
```

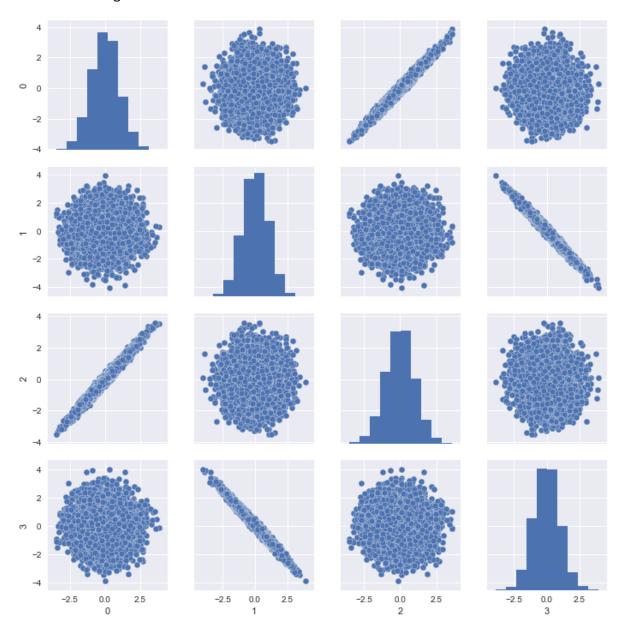
```
Out[3]: array([[<matplotlib.axes._subplots.AxesSubplot object at 0x000000000BFE87B8>,
                <matplotlib.axes. subplots.AxesSubplot object at 0x00000000C1E9AC8>,
                <matplotlib.axes. subplots.AxesSubplot object at 0x000000000C15E8D0>,
                <matplotlib.axes._subplots.AxesSubplot object at 0x000000000C3AB6D8</pre>
        >],
               [<matplotlib.axes._subplots.AxesSubplot object at 0x000000000C479080>,
                <matplotlib.axes._subplots.AxesSubplot object at 0x000000000C51DC88>,
                <matplotlib.axes._subplots.AxesSubplot object at 0x000000000C5F07B8>,
                <matplotlib.axes. subplots.AxesSubplot object at 0x000000000C6DCE80</pre>
        >],
               [<matplotlib.axes._subplots.AxesSubplot object at 0x000000000C7F29B0>,
                <matplotlib.axes. subplots.AxesSubplot object at 0x000000000CCE3198>,
                <matplotlib.axes._subplots.AxesSubplot object at 0x000000000CDE7DD8>,
                <matplotlib.axes._subplots.AxesSubplot object at 0x000000000CE90940</pre>
        >],
               [<matplotlib.axes._subplots.AxesSubplot object at 0x000000000CF5C5C0>,
                <matplotlib.axes._subplots.AxesSubplot object at 0x00000000D117E80>,
                <matplotlib.axes. subplots.AxesSubplot object at 0x000000000D224828</pre>
        >]], dtype=object)
```



#### Seaborn

In [4]: # plot each column against each other using seaborn
sns.pairplot(df)

Out[4]: <seaborn.axisgrid.PairGrid at 0xd24ef28>



### **Covariance Matrix**

```
In [5]: df.cov()
```

Out[5]:

	0	1	2	3
0	1.001558	-0.004012	0.991624	0.004125
1	-0.004012	1.005378	-0.004099	-0.995457
2	0.991624	-0.004099	1.001589	0.004081
3	0.004125	-0.995457	0.004081	1.005168

The covariance matrix is an n x n matrix where each element is  $cov(x_i,x_i)$ 

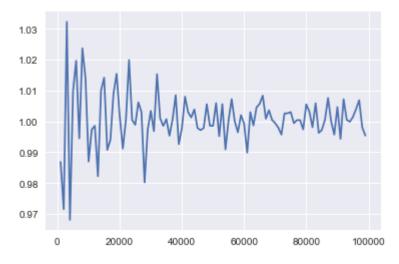
For example:  $\frac{12}^2 = \frac{1^2}^2 \cdot \frac{12}^2 \cdot$ 

Columns 0 and 1 have a low covariance magnitude, so their plot seems to have no relation. This is the case for 0 and 3, 1 and 2, and 2 and 3 as well.

Columns 0 and 2 have a high covariance, so their plot looks like a line with a positive slope.

Columns 1 and 3 have a high negative covariance, so their plot looks like a line with a negative slope.

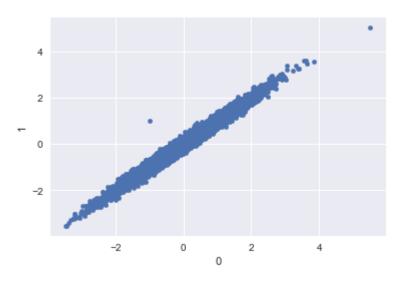
#### **Covariance vs Sample Size**



As the number of samples increases, the calculated covariance approaces the true variance.

## 2. Outliers

```
In [7]: plt.clf()
  path = 'Data\DF2'
  df=pd.read_csv(path,index_col=0,na_values='?')
  df.plot(x='0',y='1',kind='scatter')
```

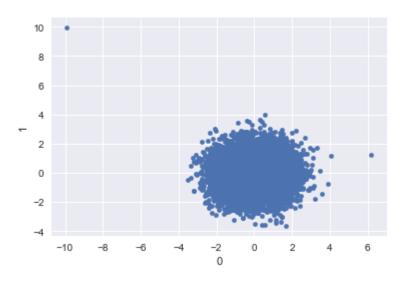


```
In [8]: cov = df.cov()

    from scipy.linalg import sqrtm
    cov_inv = np.linalg.inv(sqrtm(cov))
    data = pd.DataFrame(np.dot(df,cov_inv))

data.plot(x = 0, y = 1,kind='scatter')
```

Out[8]: <matplotlib.axes.\_subplots.AxesSubplot at 0xf79bd30>



$$\mathbf{C}_{\mathbf{z}} \triangleq E\{(\mathbf{z} - \bar{\mathbf{z}})(\mathbf{z} - \bar{\mathbf{z}})^{T}\}\$$

$$= E\left\{\left[(\mathbf{Q}\mathbf{y}) - (\mathbf{Q}\bar{\mathbf{y}})\right]\left[(\mathbf{Q}\mathbf{y}) - (\mathbf{Q}\bar{\mathbf{y}})\right]^{T}\right\}\$$

$$= E\left\{\left[\mathbf{Q}(\mathbf{y} - \bar{\mathbf{y}})\right]\left[\mathbf{Q}(\mathbf{y} - \bar{\mathbf{y}})\right]^{T}\right\}\$$

$$= E\left\{\mathbf{Q}(\mathbf{y} - \bar{\mathbf{y}})(\mathbf{y} - \bar{\mathbf{y}})^{T}\mathbf{Q}^{T}\right\}\$$

$$= \mathbf{Q}E\left\{(\mathbf{y} - \bar{\mathbf{y}})(\mathbf{y} - \bar{\mathbf{y}})^{T}\right\}\mathbf{Q}^{T}\$$

$$= \mathbf{Q}C_{\mathbf{y}}\mathbf{Q}^{T}$$

If Cy is the identity matrix and Cz is know, then Q can be determined by taking the matrix square root of Cz. We then multiply points in Z by  $Q^{-1}$ , and we will obtain Y. Now the point (-1,1) is more outlying than (5.5,5).

#### 3. Even More Standard Error

#### Constant n

```
In [9]:
    for i in range(10000):
        # take samples of e and x
        eSamples = np.random.normal(0,1,150)
        xSamples = np.random.normal(0,1,150)
        # compute y
        ySamples = -3 + eSamples
        # calculate bHat using the equation from written problem 2
        bHat = np.dot(xSamples, ySamples)/np.dot(xSamples,xSamples)
        error = bHat - 0
        a.append(error)
    #find the standard deviation of the errors
        np.std(a)
```

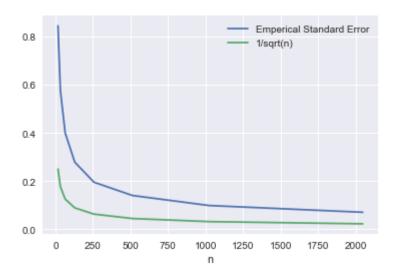
Out[9]: 0.25957849857841037

The value -0.15 is not significant since the standard deviation is greater than |-0.15|

#### Varying n

```
In [11]:
         xAxis = [2**i for i in range(4,12)]
         stdError = []
         invSqrt = [1/np.sqrt(i) for i in xAxis]
         for n in xAxis:
             errors = []
             # perform calculation from last step with varying n
             for i in range(10000):
                 eSamples = np.random.normal(0,1,n)
                 xSamples = np.random.normal(0,1,n)
                 ySamples = -3 + eSamples
                 bHat = np.dot(xSamples, ySamples)/np.dot(xSamples,xSamples)
                 error = bHat - 0
                 errors.append(error)
             stdError.append(np.std(errors))
         # plot emperical error and 1/sqrt(n) against n
         plt.plot(xAxis,stdError, label = 'Emperical Standard Error')
         plt.plot(xAxis,invSqrt, label = '1/sqrt(n)')
         plt.legend()
         plt.xlabel('n')
```

Out[11]: <matplotlib.text.Text at 0x1048f4a8>



The shape of the Emperical Standard Error matches the inverse square root of the number of samples

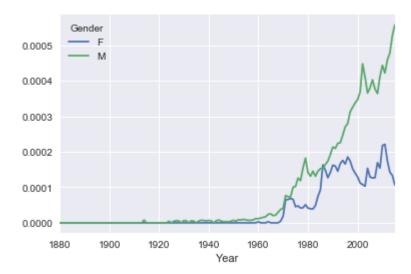
## 4. Names and Frequencies

```
In [13]: # top k names for a given year
         def topNames(k, year):
             if(not year in range(1880,2016)):
                  return None
             df=pd.read_csv('Names\yob{}.txt'.format(year),header=None,na_values='?')
             df.sort_values(2,ascending=False)
             return(df[0][:k])
         topNames(10, 2000)
Out[13]: 0
                  Emily
         1
                 Hannah
         2
                Madison
         3
                 Ashley
         4
                  Sarah
         5
                 Alexis
         6
               Samantha
         7
                Jessica
              Elizabeth
         8
                 Taylor
         Name: 0, dtype: object
In [14]:
         # name frequency from 1880 to 2015 based on gender
         def nameFreq(name):
             names = {'M':0,'F':0}
             for year in range(1880,2016):
         df=pd.read_csv('Names\yob{}.txt'.format(year),header=None,na_values='?')
                  # find all the males named name, and count them
                 males = df.loc[df[0] == name].loc[df[1] == 'M'][2].get values()
                 females = df.loc[df[0] == name].loc[df[1] == 'F'][2].get_values()
                  if(males.size != 0):
                      names['M'] += males[0]
                  if(females.size != 0):
                      names['F'] += females[0]
             return names
         nameFreq('Ali')
```

Out[14]: {'F': 9501, 'M': 23614}

```
In [201]:
          # relative name frequency from 1880 to 2015 based on gender
          def relativeNameFreq(name):
              # combine all files into one dataframe
              dataFrames = []
              for year in range(1880,2016):
                  df=pd.read_csv('Names\yob{}.txt'.format(year),names=['Name', 'Gender',
            'Freq'])
                  df['Year']=pd.Series(year,index=df.index)
                  dataFrames.append(df)
              df = pd.concat(dataFrames)
              # get the total names in a year
              totalNames = df.pivot_table(values='Freq',index='Year',columns='Gender',ag
          gfunc=np.sum)
              # get the number of people named name in a year
              totalGivenName =
          df.loc[df['Name']==name].pivot_table(values='Freq',index='Year',columns='Gende
          r',aggfunc=np.sum)
              # get the fraction of people named name in a year
              relativeFreq = totalGivenName.div(totalNames).fillna(0)
              return relativeFreq
          relativeNameFreq('Ali').plot()
```

Out[201]: <matplotlib.axes.\_subplots.AxesSubplot at 0x27b716a0>



```
In [202]: # find all names that changed in gender popularity from 1880 to 2015
          def nameGenderSwitch():
              # combine all files into one dataframe
              dataFrames = []
              for year in range(1880,2016):
                  df=pd.read_csv('Names\yob{}.txt'.format(year),names=['Name', 'Gender',
            'Freq'])
                  df['Year']=pd.Series(year,index=df.index)
                  dataFrames.append(df)
              df = pd.concat(dataFrames)
              # get the number of people named a name in a year
              totalNames = df.pivot_table(values='Freq',index=['Name','Year'],columns='G
          ender',aggfunc=np.sum).fillna(0)
              totalNames['M-F'] = np.sign(totalNames['M']-totalNames['F'])
              totalNames = totalNames['M-F']
              index = totalNames.axes[0]
              ret = {'MToF': 0, 'FToM':0}
              for a in set(index.get_level_values(0)):
                   if totalNames[a].is_monotonic_increasing and
          len(totalNames[a].unique()) > 1:
                       ret['FToM'] += 1
                       # if m < f then m > f
                        print "{} was a female name until {}, and now is a male name".fo
          rmat(a,totalNames[a][totalNames[a] != -1].index[0])
                  elif totalNames[a].is_monotonic_decreasing and len(totalNames[a].uniqu
          e()) > 1:
                       ret['MToF'] += 1
                       # if f < m then m < f
                        print "{} was a male name until {}, and now is a female name".fo
          rmat(a,totalNames[a][totalNames[a] != 1].index[0])
              return ret
          nameGenderSwitch()
```

### Out[202]: {'FToM': 756, 'MToF': 609}

#### Examples:

Niger was a female name until 1994, and now is a male name

Chesleigh was a male name until 2010, and now is a female name

Chade was a male name until 1985, and now is a female name

### 5. Visualization Tools and Missing/Hidden Values

```
In [209]: tweets = pd.read_csv('tweets.csv')
          tweets['user_location'].head()
Out[209]: 0
                   Wheeling WV
          1
                           NaN
          2
                           NaN
                        global
          3
          4
               California, USA
          Name: user_location, dtype: object
In [213]: def get_state(row):
              state = []
              text = row['user location'].lower()
              if "west virginia" in text or r"\bwv\b" in text or "Wheeling" in text:
                  state.append("WV")
              if "texas" in text or r"\btx\b" in text or "austin" in text or "houston" i
          n text:
                  state.append("TX")
              if "new york" in text or r"\bny\b" in text or "new york city" in text:
                   state.append("NY")
              return ",".join(state)
          tweets['user_location'] = tweets['user_location'].astype(str)
          tweets['state'] = tweets.apply(get state,axis=1)
          tweets['state'].value_counts()
                   226352
Out[213]:
          TX
                     6285
          NY
                     4592
                      249
          WV
          TX,NY
          Name: state, dtype: int64
```

We can find the number of states by searching for certain terms in the user\_location field. We can assign more states with more strict criteria and checking for more states/cities.

a)  $P(Z_{avg} > c) = P(\frac{Z_{avg} - u}{6/\sqrt{5}}) \approx (-\Phi(\frac{c - u}{6/\sqrt{5}})$ 

Zn = Zavg - M

secopopopopopo

かかか

10

つか

o C

M.

P(Zn > C-MIN)

P(Zn > .1. J10000) 21 - \$\phi(10) \pi 0.16
P(Zn > .01.100) 21 - \$\phi(1) \pi 0.16
P(Zn > .001.100) 21 - \$\phi(1) \pi 0.46

b)  $P(2n > \frac{n^{1/3} - \mu}{6/\sqrt{n}}) \approx 1 - \phi(\frac{n^{-1/3} - \mu}{6/\sqrt{n}})$   $P(2n > \frac{n^{1/3} - \mu}{6/\sqrt{n}}) \approx 1 - \phi(\frac{n^{-1/3} - \mu}{6/\sqrt{n}})$  $P(2n > \frac{n^{-2/3} - \mu}{6/\sqrt{n}}) \approx 1 - \phi(\frac{n^{-2/3} - \mu}{6/\sqrt{n}})$ 

2 a) † Σ(xiβ-yi)2= † Σxi2β2= = Σxiyiβ+ = yi2 A= $\frac{1}{1}$  $\Rightarrow \lim_{x \to \infty} \sum_{x \to \infty} \sum_$ > mint | XB - 3112 (c)  $\hat{\beta} = \frac{\sum_{x_i \mid x_i \mid \beta + e_i}}{\sum_{x_i \mid x_i}}$ c) \( \hat{\beta} = (x^{\tau} x)^{-1} x^{\tau} (\beta x \beta + \hat{\epsilon}) \\
= (x^{\tau} x)^{-1} (x^{\tau} x) \beta + (x^{\tau} x)^{\tau} x^{\tau} \hat{\epsilon} \) = ExiZB + Exei ExiZ = B + (xTX) XTE  $= \beta + \frac{\sum_{i} e_{i}}{\sum_{k_{i}}^{2}}$ == (xTx) xTe = B + X8. ex Z = X.XT