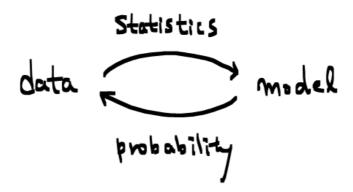
Statistics

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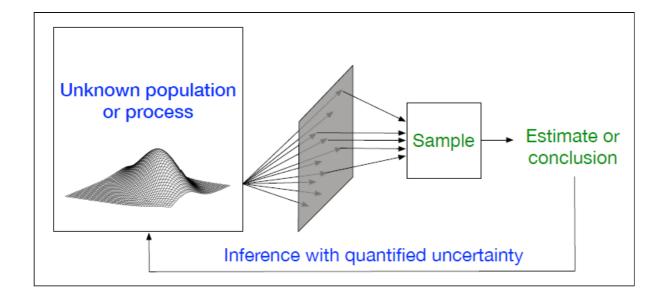


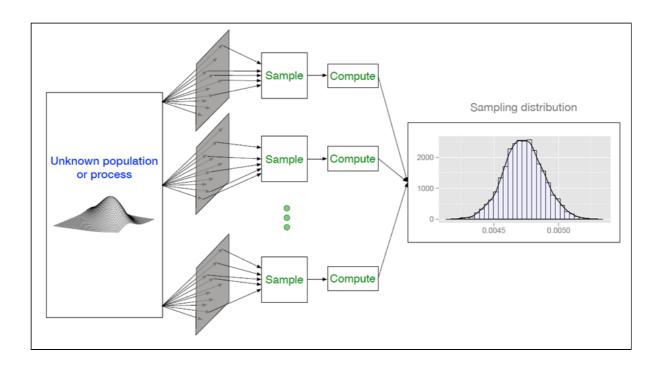
1. Populations and Samples

- A population includes all the elements from a set of data
- A parameter is a quantity computed from a population
 - mean, *μ*
 - variance, σ^2
- A sample is a subset of the population.
 - one or more observations
- A statistic is a quantity computed from a sample
 - sample mean, \bar{x}
 - sample variance, s^2
 - sample correlation, S_{xy}

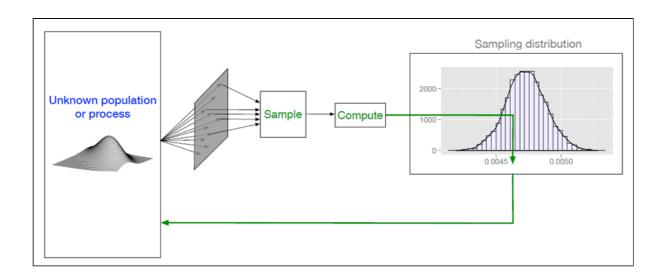
2. Inference

- True population or process is modeled probabilistically.
- Sampling supplies us with realizations from probability model.
- Compute something, but recognize that we could have just as easily gotten a different set of realizations.





• We want to infer the characteristics of the true probability model from our **one** sample.

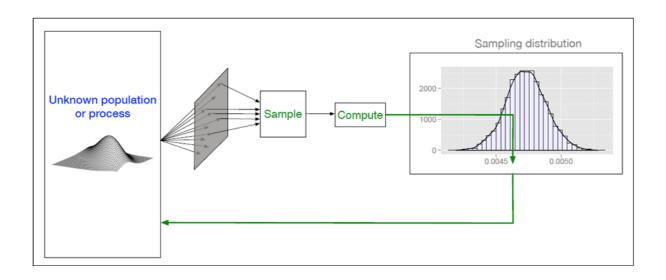


3. Law of Large Numbers

• Sample mean converges to the population mean as sample size gets large

$$ar x o \mu_x \qquad ext{as} \qquad m o \infty$$

· True for any probability density functions



wikipedia (http://en.wikipedia.org/wiki/Law of large numbers)

3.1. Sample Mean and Sample Size

· sample mean and sample variance

$$ar{x} = rac{x_1 + x_2 + \ldots + x_m}{m} \ s^2 = rac{\sum_{i=1}^m (x_i - ar{x})^2}{m-1}$$

- suppose $x \sim U[0,1]$

In [1]:

import numpy as np
import matplotlib.pyplot as plt

%matplotlib inline

In [2]:

```
# statistics
# numerically understand statisticcs

m = 100
x = np.random.rand(m,1)

#xbar = 1/m*np.sum(x, axis=0)
#np.mean(x, axis=0)
xbar = 1/m*np.sum(x)
np.mean(x)

varbar = (1/(m - 1))*np.sum((x - xbar)**2)
np.var(x)

print(xbar)
print(np.mean(x))
print(varbar)
print(np.var(x))
```

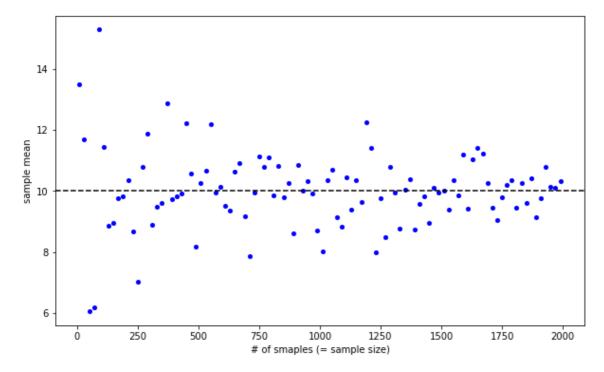
- 0.458759386078
- 0.458759386078
- 0.0749424145354
- 0.07419299039

In [3]:

```
# various sample size m
m = np.arange(10,2000,20)
means = []

for i in m:
    x = np.random.normal(10, 30, i)
    means.append(np.mean(x))

plt.figure(figsize=(10,6))
plt.plot(m, means, 'bo', markersize = 4)
plt.axhline(10, c='k', linestyle='dashed')
plt.xlabel('# of smaples (= sample size)', fontsize = 10)
plt.ylabel('sample mean', fontsize = 10)
plt.show()
```



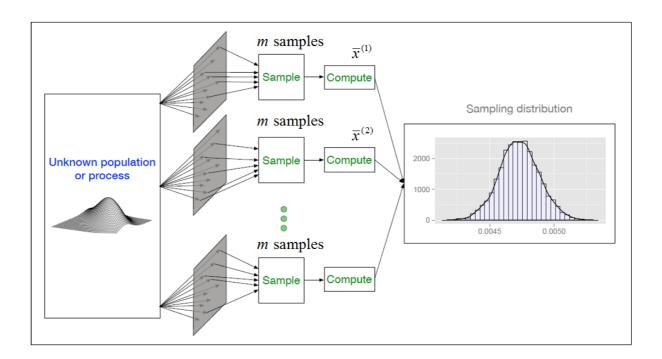
4. Central limit theorem

- Sample mean (not samples) will be approixmatedly normally distributed as a sample size $m o \infty$ $\bar x = \frac{x_1+x_2+\ldots+x_m}{m}$

$$ar{x}=rac{x_1+x_2+\ldots+x_m}{m}$$

- More samples provide more confidence (or less uncertainty)
- Note: true regardless of any distribution of population

$$ar{x}
ightarrow N\left(\mu_x, \left(rac{\sigma}{\sqrt{m}}
ight)^2
ight)$$

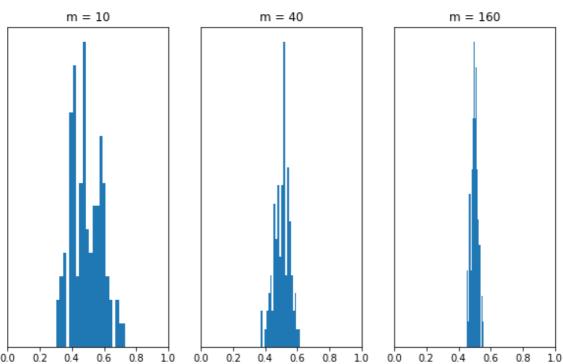


• wikipedia (http://en.wikipedia.org/wiki/Central limit theorem)

4.1. Variance Gets Smaller as m is Larger

- · Seems approximately Gaussian distributed
- numerically demostrate that sample mean follows the Gaussin distribution

```
N = 100
m = np.array([10, 40, 160]) # sample of size m
S1 = []
                                          # sample mean (or sample average)
S2 = []
S3 = []
for i in range(N):
                 S1.append(np.mean(np.random.rand(m[0],1)))
                 S2.append(np.mean(np.random.rand(m[1],1)))
                 S3.append(np.mean(np.random.rand(m[2],1)))
plt.figure(figsize=(10, 6))
plt.subplot(1,3,1), \ plt.hist(S1,\ 21), \ plt.xlim([0,\ 1]), \ plt.title('m = '+ \ str(m[0])), \ plt.xlim([0,\ 1]), \ plt.title('m = '+ \ str(m[0])), \ plt.xlim([0,\ 1]), \ p
lt.yticks([])
plt.subplot(1,3,2), plt.hist(S2, 21), plt.xlim([0, 1]), plt.title('m = '+ str(m[1])), p
lt.yticks([])
plt.subplot(1,3,3), plt.hist(S3, 21), plt.xlim([0, 1]), plt.title('m = '+ str(m[2])), p
lt.yticks([])
plt.show()
```



5. How to Generate Random Numbers (Samples or data)

• Data sampled from population/process/generative model

In [5]:

```
## random number generation (1D)
m = 1000;

# uniform distribution U(0,1)
x1 = np.random.rand(m,1);

# uniform distribution U(a,b)
a = 1;
b = 5;
x2 = a + (b-a)*np.random.rand(m,1);

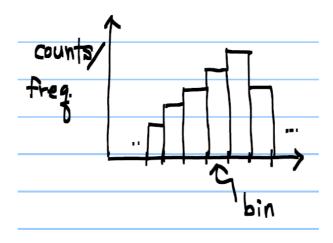
# standard normal (Gaussian) distribution N(0,1^2)
# x3 = np.random.normal(0, 1, m)
x3 = np.random.randn(m,1);

# normal distribution N(5,2^2)
x4 = 5 + 2*np.random.randn(m,1);

# random integers
x5 = np.random.randint(1, 6, size = (1,m));
```

Histogram: graphical representation of data distribution

 \Rightarrow rough sense of density of data



6. Multivariate Statistics

$$x^{(i)} = egin{bmatrix} x_1^{(i)} \ x_2^{(i)} \ dots \end{bmatrix}, \quad X = egin{bmatrix} - & (x^{(i)})^T & - \ - & (x^{(i)})^T & - \ dots \ dots & dots \ - & (x^{(m)})^T & - \ \end{bmatrix}$$

•
$$m$$
 observations $\left(x^{(i)},x^{(2)},\cdots,x^{(m)}\right)$ sample mean $\bar{x}=\dfrac{x^{(1)}+x^{(2)}+\cdots+x^{(m)}}{m}=\dfrac{1}{m}\sum_{i=1}^mx^{(i)}$

sample variance
$$S^2 = rac{1}{m-1} \sum_{i=1}^m (x^{(i)} - ar{x})^2$$

(Note: population variance
$$\sigma^2 = \frac{1}{N} \sum_{i=1}^{N} (x^{(i)} - \mu)^2$$

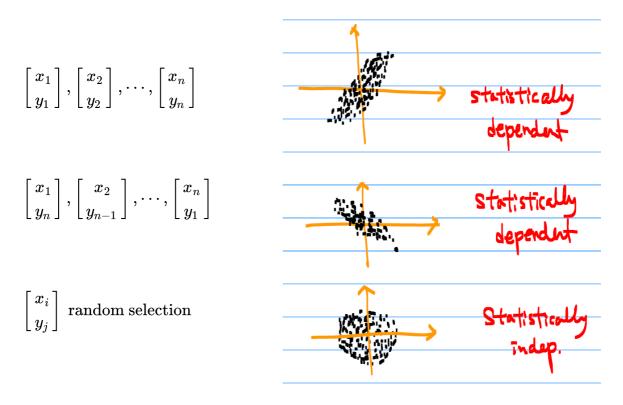
6.1. Two random variables

$$\begin{aligned} \text{Sample Variance} : S_x &= \frac{1}{m-1} \sum_{i=1}^m \left(x^{(i)} - \bar{x} \right)^2 \\ \text{Sample Covariance} : S_{xy} &= \frac{1}{m-1} \sum_{i=1}^m \left(x^{(i)} - \bar{x} \right) \left(y^{(i)} - \bar{y} \right) \\ \text{Sample Covariance matrix} : S &= \begin{bmatrix} S_x & S_{xy} \\ S_{yx} & S_y \end{bmatrix} \\ \text{sample correlation coefficient} : r &= \frac{S_{xy}}{\sqrt{S_{xx} \cdot S_{yy}}} \end{aligned}$$

- strength of **linear** relationship between two variables, x and y
- assume

$$x_1 \le x_2 \le \dots \le x_n$$

 $y_1 \le y_2 \le \dots \le y_n$



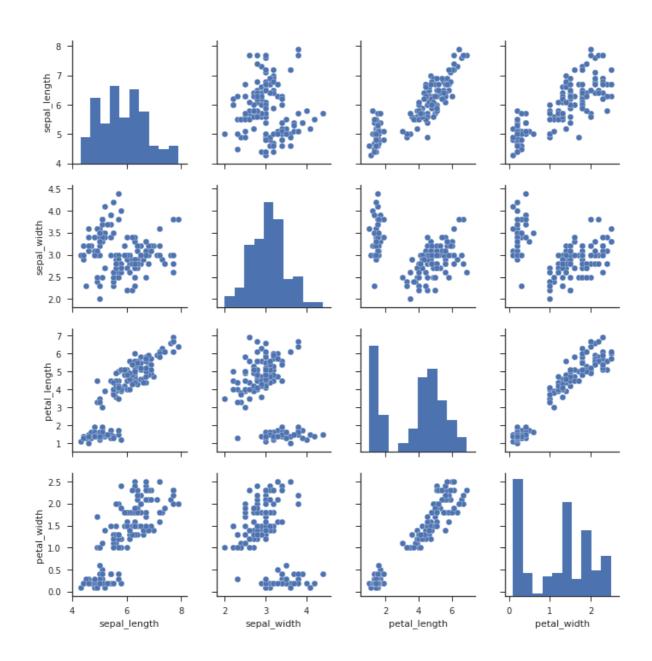
6.2. Correlation coefficient

- ullet +1
 ightarrow close to a straight line
- ullet -1
 ightarrow close to a straight line
- · Indicate how close to a linear line, but
- · No information on slope

· does not tell anything about causality

6.3. Correlation Coefficient Plot

- · Plots correlation coefficients among pairs of variables
- http://rpsychologist.com/d3/correlation/ (http://rpsychologist.com/d3/correlation/)

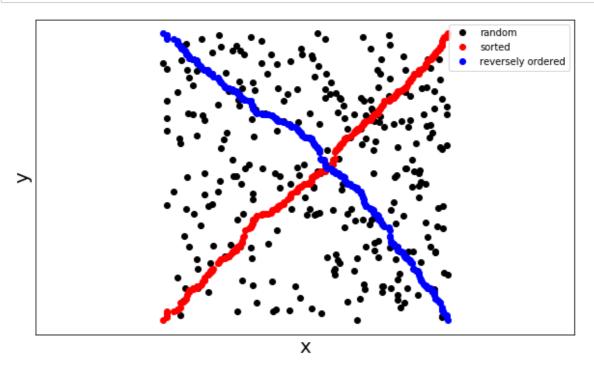


6.4. Covariance Matrix

$$\sum = \begin{bmatrix} E[(X_1 - \mu_1)(X_1 - \mu_1)] & E[(X_1 - \mu_1)(X_2 - \mu_2)] & \cdots & E[(X_1 - \mu_1)(X_n - \mu_n)] \\ E[(X_2 - \mu_2)(X_1 - \mu_1)] & E[(X_2 - \mu_2)(X_2 - \mu_2)] & \cdots & E[(X_2 - \mu_2)(X_n - \mu_n)] \\ \vdots & \vdots & \ddots & \vdots \\ E[(X_n - \mu_n)(X_1 - \mu_1)] & E[(X_n - \mu_n)(X_2 - \mu_2)] & \cdots & E[(X_n - \mu_n)(X_n - \mu_n)] \end{bmatrix}$$

In [6]:

```
# correlation coefficient
m = 300
x = np.random.rand(m)
y = np.random.rand(m)
xo = np.sort(x)
yo = np.sort(y)
yor = -np.sort(-y)
plt.figure(figsize=(10, 6))
plt.plot(x, y, 'ko', label='random')
plt.plot(xo, yo, 'ro', label='sorted')
plt.plot(xo, yor, 'bo', label='reversely ordered')
plt.axis('equal')
plt.xticks([])
plt.yticks([])
plt.xlabel('x', fontsize=20)
plt.ylabel('y', fontsize=20)
plt.legend()
plt.show()
print(np.corrcoef(x,y))
print(np.corrcoef(xo,yo))
print(np.corrcoef(xo,yor))
```



In [7]:

```
# correlation coefficient
m = 300
x = 2*np.random.randn(m)
y = np.random.randn(m)
xo = np.sort(x)
yo = np.sort(y)
yor = -np.sort(-y)
plt.figure(figsize=(10, 6))
plt.plot(x, y, 'ko', label='random')
plt.plot(xo, yo, 'ro', label='sorted')
plt.plot(xo, yor, 'bo', label='reversely ordered')
plt.xticks([])
plt.yticks([])
plt.xlabel('x', fontsize=20)
plt.ylabel('y', fontsize=20)
plt.axis('equal')
plt.legend()
plt.show()
print(np.corrcoef(x,y))
print(np.corrcoef(xo,yo))
print(np.corrcoef(xo,yor))
```

