Question: With limited resources, which of the following Options would give you a more accurate reflection of the Parameters?:

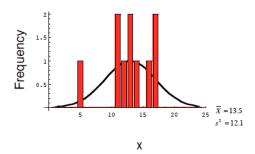
1. 1000 samples, each of 10 individuals

or

2. 10 samples, each of 1000 individuals?

Frequency distributions NOT sampling distributions

Sample size 10 from Normal distribution with μ =13 and σ^2 =16



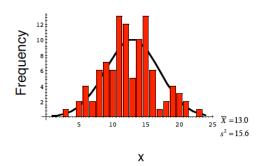
Another sample of 10 from same distribution

Frequency

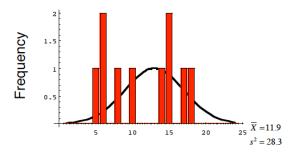
X

 $s^2 = 13.0$

A sample of 100 from the same population distribution

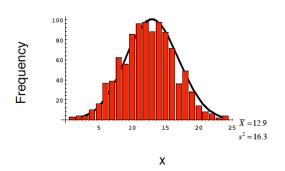


A third sample of 10 from the same distribution



Х

A sample of 1000 from the same population distribution



n	$ar{X}$	s ²
10	13.5	12.1
10	13.3	13.0
10	11.9	28.3
100	13.0	15.6
1000	12.9	16.3

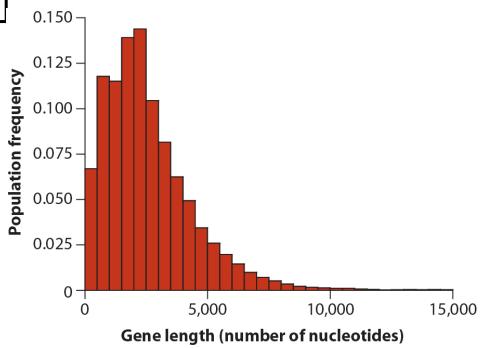
Now we understand the concept of the sampling distribution of an estimate, so let's move on to an example....

Example: The length of human genes (page 96). From build 35, there are 20,290 genes. Since we **know** the size of **all** predicted Genes (**the entire population of gene lengths**), we are in the unique situation of calculating parameters directly from the data instead of inferring their values from samples. We can compare the values we get from samples to the <u>real</u>, true parameter values.

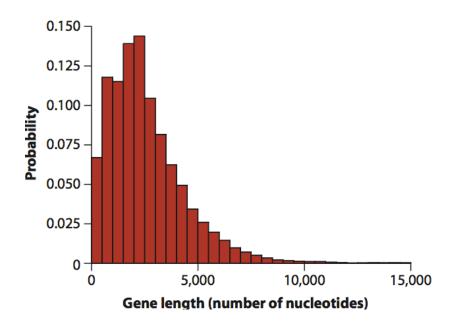
http://phylo.bio.ku.edu/biostats/geneLenDemo.html

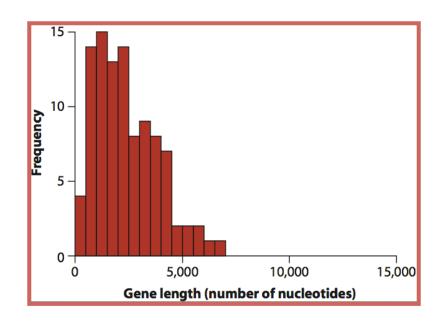
<u>Example:</u> The length of human genes. From build 35, there are 20,290 genes. Since we *know* the size of *all* predicted genes, we are in the unique situation of calculating parameters directly from the data instead of inferring their values from samples.

<u>Name</u>	<u>Parameter</u>	<u>Value</u>
<u>Mean</u>	μ	2622.0
<u>S.D.</u>	σ	2036.9



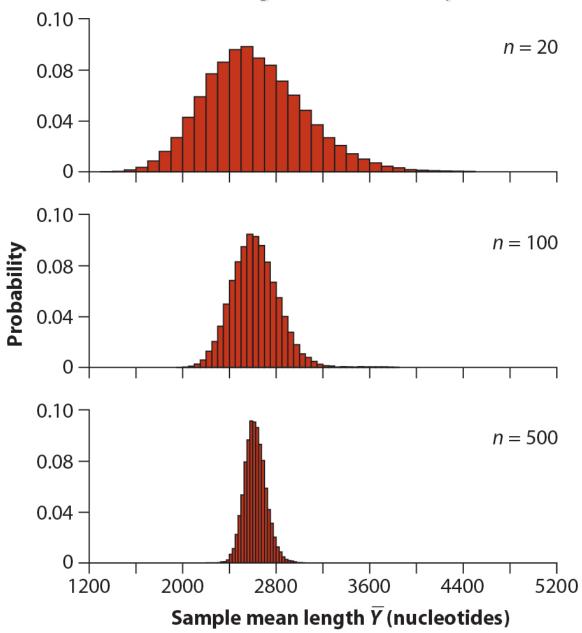
<u>Example:</u> The length of human genes. From build 35, there are 20,290 genes. Since we *know* the size of *all* predicted genes, we are in the unique situation of calculating parameters directly from the data instead of inferring their values from samples.





<u>Name</u>	<u>Parameter</u>	<u>Value</u>
<u>Mean</u>	μ	2622.0
<u>S.D.</u>	σ	2036.9

<u>Name</u>	<u>Parameter</u>	<u>Value</u>
<u>Mean</u>	\overline{X}	2411.8
<u>S.D.</u>	S	1463.5



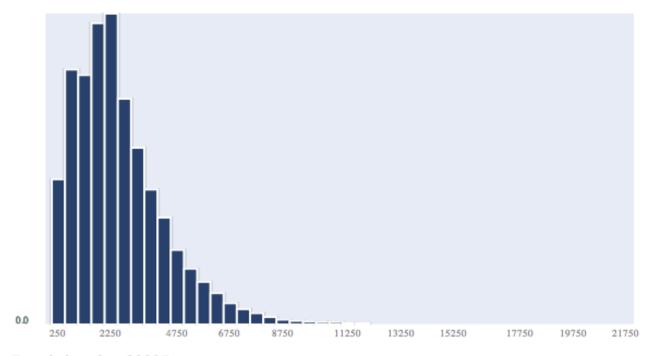
Question: With limited resources, would you rather have:

1. 1000 samples, each of 10 individuals

or

2. 10 samples, each of 1000 individuals?

Gene lengths in human genes (some long genes were excluded from consideration to make it easier to make these graphs):



Population size: 20287

Population mean length: $\mu = 2614.36$

Population standard deviation of length: $\sigma = 1897.30$

http://phylo.bio.ku.edu/biostats/geneLenDemo.html

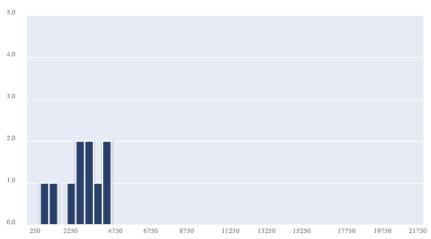
Question: With limited resources, would you rather have:

1. 1000 samples, each of 10 individuals or 2. 10 samples, each of 1000 individuals?

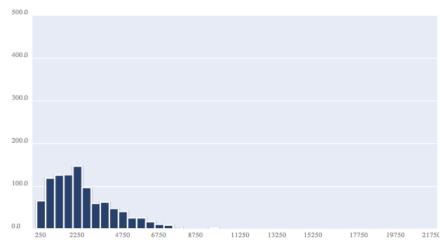
1000 samples of 10 individuals

10 samples of 1000 individuals

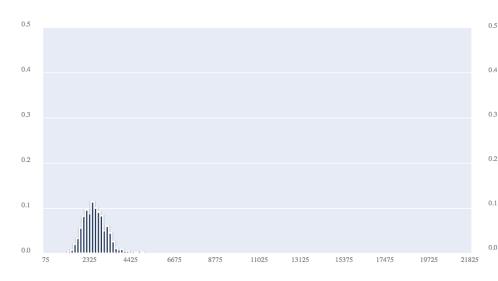
Gene lengths in the last random sample:



Mean length in sample: $\bar{Y} = 2812.90$ Sample standard deviation: s = 1193.99 Gene lengths in the last random sample:

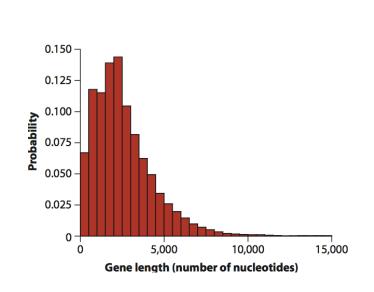


Mean length in sample: $\bar{Y} = 2633.77$ Sample standard deviation: s = 1988.42

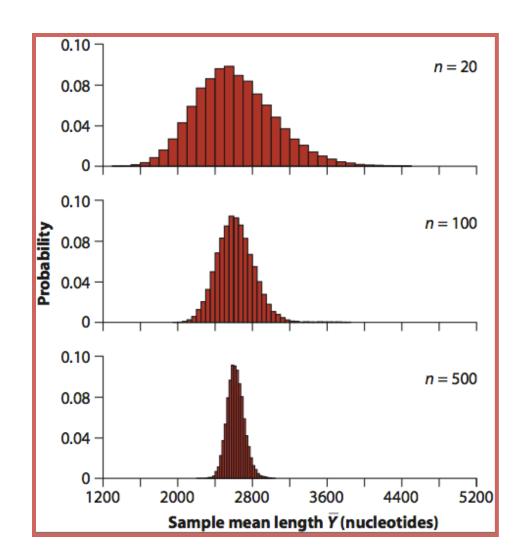




<u>Example:</u> The length of human genes (page 84). From build 35, there are 20,290 genes. Since we *know* the size of *all* predicted genes, we are in the unique situation of calculating parameters directly from the data instead of inferring their values from samples.



<u>Name</u>	<u>Parameter</u>	<u>Value</u>
<u>Mean</u>	μ	2622.0
<u>S.D.</u>	σ	2036.9

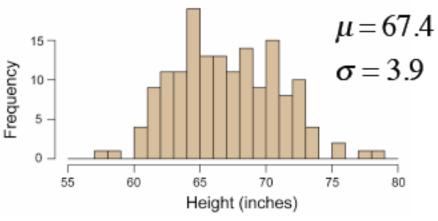


Standard Error (of the mean): the standard deviation of the sampling distribution of some statistic

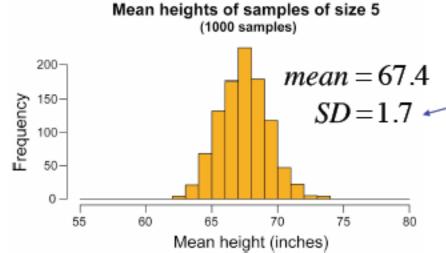
$$\sigma_{\bar{Y}} = \frac{\sigma}{\sqrt{n}}$$

*Some statistic could be: mean, difference between two means, correlation coefficient etc

Measuring Uncertainty



O



$$\mu_{\overline{\gamma}} = \mu = 67.4$$

$$\sigma_{\overline{Y}} = \frac{\sigma}{\sqrt{n}} = \frac{3.9}{\sqrt{5}} = 1.7$$

The math works!

The problem is, we rarely know σ .

Measuring Uncertainty

Estimate of the standard error (of the mean):

$$SE_{\bar{Y}} = \frac{S}{\sqrt{n}}$$

Measuring Uncertainty

