Estimating Distributions

Credit: Ryan Henning

- Review common moments: μ , σ^2
- Motivation: Why estimate distributions?
- Four standard methods:
 - Method of Moments (MOM)
 - Maximum Likelihood Estimation (MLE)
 - Maximum a Posteriori (MAP)
 - Kernel Density Estimation (KDE)
- Parametric vs nonparametric methods

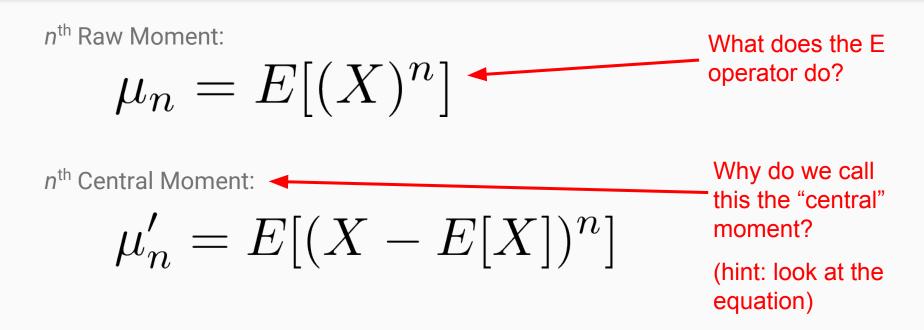


Review common moments

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Review: *Moments* of a random variable *X*





Fill in the blank:

Fill in the blank:

$$\mu = E[X]$$

For discrete random variables with *k* possible outcomes:

$$E[X] = \sum_{i=1}^{k} x_i P(X = x_i)$$
 P is the "Probability ____ Function"

For continuous random variables:

$$E[X] = \int_{-\infty}^{\infty} x f(x) dx$$
 f is the "Probability ____ Function"



$$\sigma^2 = E[(X - E[X])^2]$$

For discrete random variables with *k* possible outcomes:

$$\sigma^2 = \sum_{i=1}^{\kappa} (x_i - \mu)^2 P(X = x_i)$$

For continuous random variables:

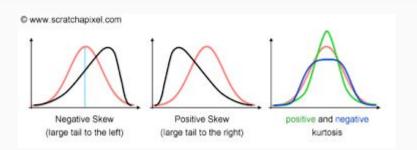
$$\sigma^2 = \int_{-\infty}^{\infty} (x - \mu)^2 f(x) \, dx$$

Variance measures the "spread" of data. Can you see that in the equations?

Normalized moments



$$\frac{\mu_n}{\sigma^n} = \frac{\mathrm{E}[(X-\mu)^n]}{\sigma^n}.$$



The n'th normalized moment is the n'th central moment divided by σ^n

The normalised **third** central moment is called the **skewness**, often γ . The normal distribution has a skewness of 0.

The **kurtosis** κ is defined to be the normalised **fourth** central moment.

The excess kurtosis is the kurtosis minus 3.

The normal distribution has kurtosis of 3 and excess kurtosis of 0.

Why Estimate Distributions?

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Why estimate distributions?

Example 1: You have data on how many people order cakes every day at your bakery, and you want to estimate the probability of selling out.

Example 2: You have data on how often your car breaks down, and you want to know your chances of safely crossing the country in it.

Example 3: You have data on how many people visit your website each day, and you want to know the probability of your servers being overloaded.

Method of Moments (MOM)

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Method of Moments (MOM)

Overview:

1. Assume an underlying distribution for your domain.

E.g. Poisson, Bernoulli, Binomial, Gaussian

2. Compute the relevant sample moments.

E.g. Mean, Variance

3. Plug the sample moments into the PMF/PDF of the assumed distribution.

Example 1: Method of Moments



Your website visitor log shows the following number of visits for each of the last seven days: [6, 4, 7, 4, 9, 3, 5]. What's the probability of zero visitors tomorrow?

Which underlying distribution should we assume?

Poisson! Let's look at Wikipedia to remind ourselves what it is. :)

What moment should we estimate?

The mean. Our mean estimate will become the estimate for the only parameter used in the Poisson distribution: λ

Your website visitor log shows the following number of visits for each of the last seven days: [6, 4, 7, 4, 9, 3, 5]. What's the probability of zero visitors tomorrow?

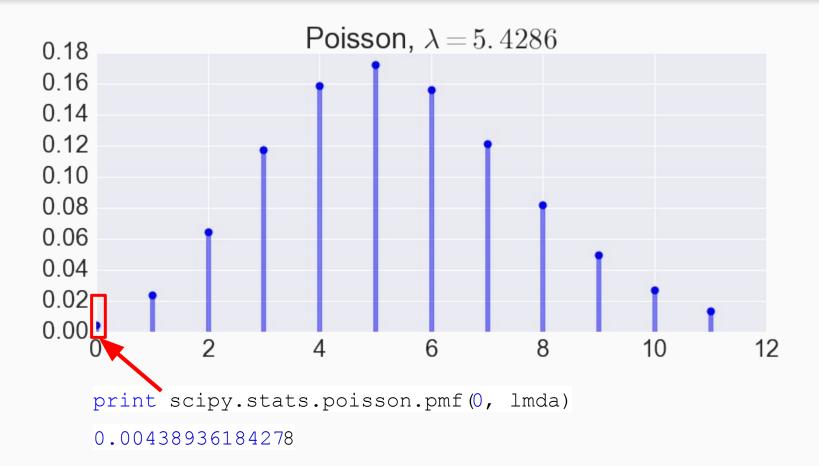
From our samples, estimate the mean:

$$\bar{x} = (6+4+7+4+9+3+5)/7 = 5.43$$

Our mean estimate is used to estimate λ :

$$\lambda = 5.43$$







You flip a coin 100 times. It comes up heads 52 times. What's the MOM estimate that in the next 100 flips the coin will be heads <= 45 times?

Which underlying distribution should we assume?

Binomial... note: We really only have one binomial sample here.

What moment should we estimate?

The mean. We actually only have one sample here where the result is 52. So the mean is 52.



You flip a coin 100 times. It comes up heads 52 times. What's the MOM estimate that in the next 100 flips the coin will be heads <= 45 times?

From our one binomial sample, we know:

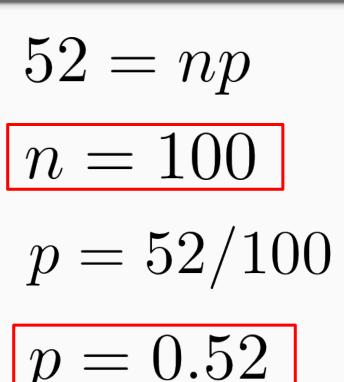
$$\bar{x} = 52$$

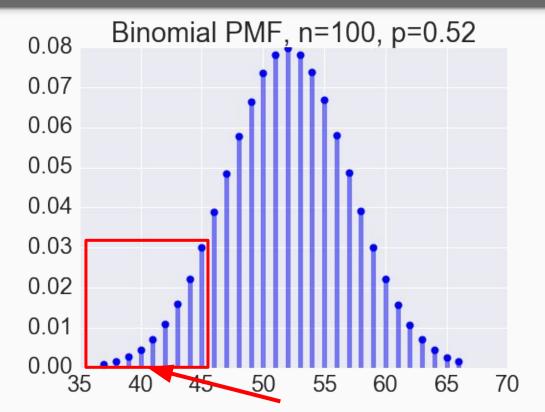
The binomial distribution has mean:

$$\mu = np$$

What does MOM say to do next?

Use the sample moments to estimate the distribution parameters. In this case, the parameter we need to estimate is p.





Probability in the next 100 flips the coin will be heads <= 45 times:

print scipy.stats.binom.cdf(45, n, p) \Rightarrow 0.096653350327

Maximum Likelihood Estimation (MLE)

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Maximum Likelihood Estimation (MLE)

Overview:

Law of Likelihood:

If P(X|H1) > P(X|H2), then the evidence supports H1 over H2.

Question:

Which hypothesis does the evidence most strongly support?

Answer:

The hypothesis H that maximizes P(X|H), which is found via MLE.

Maximum Likelihood Estimation (MLE)

Overview:

- 1. **Assume an underlying distribution for your domain.** (just like with MOM) E.g. Poisson, Bernoulli, Binomial, Gaussian
- Define the likelihood function.
 We want to know the likelihood of the data we observe under different distribution parameterizations.
- 3. Choose the parameter set that maximizes the likelihood function.



$$f(x_1, x_2, \dots, x_n | \theta) = f(x_1 | \theta) f(x_2 | \theta) \dots f(x_n | \theta)$$

 θ parameters of the assumed pdf f(.)

True because we assume X is i.i.d. Recall, what does i.i.d. Mean? What's the i. part? What's the i.d. part?

$$\mathcal{L}(\theta|x_1, x_2, \dots, x_n) = f(x_1, x_2, \dots, x_n|\theta) = \prod_{i=1}^n f(x_i|\theta)$$

 $\mathcal{L}(.)$ is the same as f(.) just with θ as variables and x as fixed parameters

We will find θ to maximize the log-likelihood function:

$$\hat{\theta}_{\text{mle}} = \arg\max_{\theta \in \Theta} \log \left(\mathcal{L}(\theta | x_1, \dots, x_n) \right)$$

Whoa whoa... what? Let's talk about it.



You flip a coin 100 times. It comes up heads 52 times. What's the MLE estimate that in the next 100 flips the coin will be heads <= 45 times?

Yep, same example...

Which underlying distribution should we assume?

Binomial... (like last time)

... now we need to define our likelihood function...



$$X_{i} \stackrel{iid}{\sim} Bin(n, p) \qquad i = 1, 2, \dots, n \qquad f(x_{i}|p) = \binom{n}{x_{i}} p^{x_{i}} (1 - p)^{n - x_{i}}$$

$$log \mathcal{L}(p) = \sum_{i=1}^{n} \left[log \binom{n}{x_{i}} + x_{i} log p + (n - x_{i}) log (1 - p) \right]$$

$$\frac{\partial log \mathcal{L}(p)}{\partial p} = \sum_{i=1}^{n} \left[\frac{x_{i}}{p} - \frac{n - x_{i}}{1 - p} \right] = 0$$

$$\hat{p}_{MLE} = \boxed{\bar{x} \atop n}$$

Maximum a Posteriori (MAP)

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Maximum a Posteriori (MAP)

MAP is MLE if we have prior knowledge about the parameter distribution

MLE → MAP, just use Bayes' Theorem

$$f(\theta|x) = \frac{f(x|\theta)g(\theta)}{\int_{\theta' \in \Theta} f(x|\theta')g(\theta') \, d\theta'} \propto f(x|\theta)g(\theta)$$
 MLE is just this part.

MLE vs MAP

MLE solves:

$$\hat{\theta}_{\text{mle}} = \arg\max_{\theta \in \Theta} f(x_1, x_2, \dots, x_n | \theta)$$

MAP solves:

$$\hat{\theta}_{\text{map}} = \arg\max_{\theta \in \Theta} f(x_1, x_2, \dots, x_n | \theta) g(\theta)$$

What if all θ are equally likely?

MAP includes the prior belief.



MOM vs MLE

- MOM is inferior to MLE, because maximum likelihood estimators have higher probability of being close to the quantities to be estimated (unbiased estimator for large samples).
- MLE may be intractable in some situations, whereas the MOM estimates can be quickly and easily calculated by hand or using a computer.
- MOM estimates may be used as the first approximations to the solutions of the MLE method, and successive improved approximations may then be found by the Newton-Raphson method (numerical).
- Sometimes, MOM estimates may be outside of the parameter space; i.e., they are unreliable, which is never a problem with the MLE method.
- MOM estimates are not necessarily sufficient statistics, i.e., they sometimes fail to take into account all relevant information in the sample (like prior distribution, see MAP).
- MOM may be preferred to MLE for estimating some structural parameters (e.g., parameters of a utility function, instead of parameters of a known probability distribution), when appropriate probability distributions are unknown.

Kernel Density Estimation (KDE)

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Nonparametric Techniques

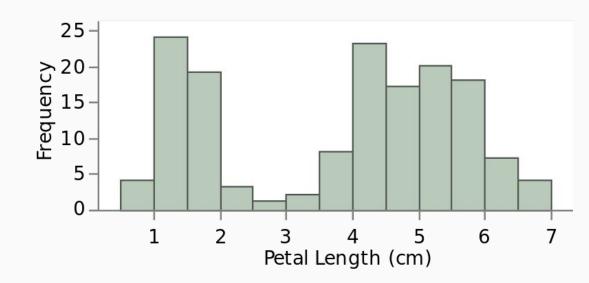
Question: How can we model data that does not follow a known distribution?

Answer: Use a nonparametric technique.



Histograms

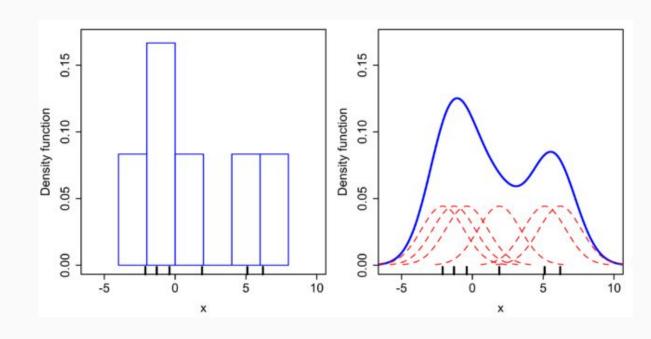
A histogram groups continuous data into discrete intervals and displays relative frequencies. But it's not a smooth distribution. :(



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Kernel Density Estimation (KDE)

KDE is a nonparametric way to estimate the PDF of a random variable. KDE smooths the histogram by summing "kernel functions" (usually Gaussians) instead of binning into rectangles.

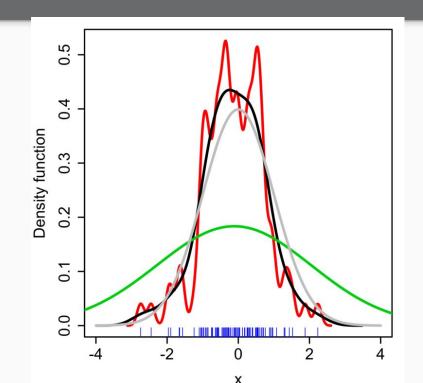


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Kernel Density Estimation (KDE)

Kernel functions have a *bandwidth* parameter to control under- and over-fitting.

Each curve on the right shows an estimated PDF with different bandwidths.



Parametric vs Nonparametric Methods

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Estimating Distributions

Parametric vs Nonparametric Methods

Parametric: We assume an underlying distribution, then we use our data to estimate the parameters of that underlying distribution. E.g. Using:

- Method of Moments (MOM)
- Maximum Likelihood Estimation (MLE)
- Maximum a Posteriori (MAP)

Nonparametric: We don't assume any *single* underlying distribution, but instead we fit a combination of distributions to the observed data. E.g. Using:

Kernel Density Estimation (KDE)



Parametric methods:

- Based on assumptions about the distribution of the underlying population and the parameters from which the sample was taken.
- 2. If the data deviates strongly from the assumptions, could lead to incorrect conclusions.

Nonparametric methods:

- NOT based on assumptions about the distribution of the underlying population.
- 2. Generally not as powerful -- less inference can be drawn.
- 3. Interpretation can be difficult... what does the wiggly curve mean?

Sampling

Credit: Ryan Henning

- Population Inference & Sampling
- Central Limit Theorem
- Confidence Intervals
- Bootstrapping



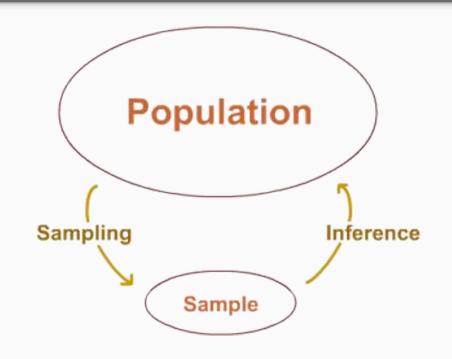
Population Inference & Sampling

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Population Inference

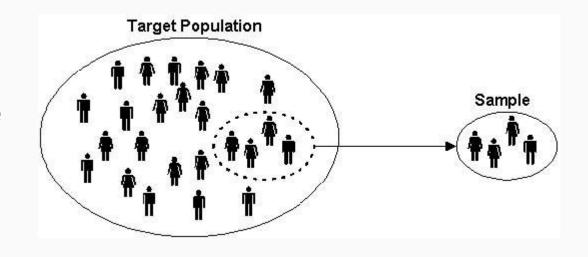
- Start with a question/hypothesis
- Design an experiment
- Collect data
- Analyze
- Check the results
- Repeat? Redesign?



Collecting data: Taking a sample

A sample should be representative of the population.

Random sampling is often the best way to achieve this. Ideally: each subject has an equal chance of being in the sample.





Random sampling is surprisingly hard to do...

Scenario: You want to estimate the percentage of dog owners in Austin.

Method 1: Go to the nearby dog park and ask **random** people if they own dogs until you have *n* responses.

Method 2: Stand on 6th and Congress and ask **random** people if they own dogs until you have *n* responses.

Method 3: Repeat *n* times: Pick a **random** neighborhood in Austin (weighted by census data per neighborhood), go to that neighborhood, ask **random** people you see if they own dogs until you get 1 response.

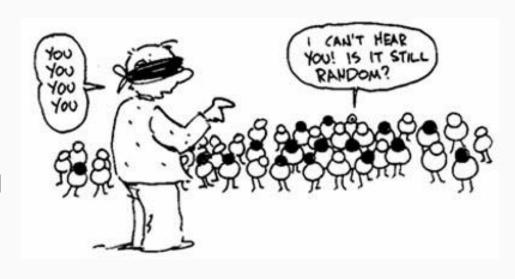
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Random sampling... just do the best you can.

Often it's impossible to do *perfect* random sampling.

So...

- 1. do the best you can,
- 2. call out possible objections, and
- 3. make a case for why you think your results are valid.





Random sampling in the digital age...

You might think that random sampling in a digital context is easier, and you're right! But there are still gotchas.

<u>Scenario:</u> Slack is testing a new features ("channel polling", a way to survey people in a channel). They'd like to test the feature on only a subset of their users (n), then draw inference about their entire userbase.

Method 1: SELECT user_id FROM users LIMIT n;

Method 2: SELECT user_id FROM users ORDER BY RANDOM() LIMIT n;

Central Limit Theorem

$$\bar{X} \sim N(\mu, \frac{\sigma^{-}}{n})$$

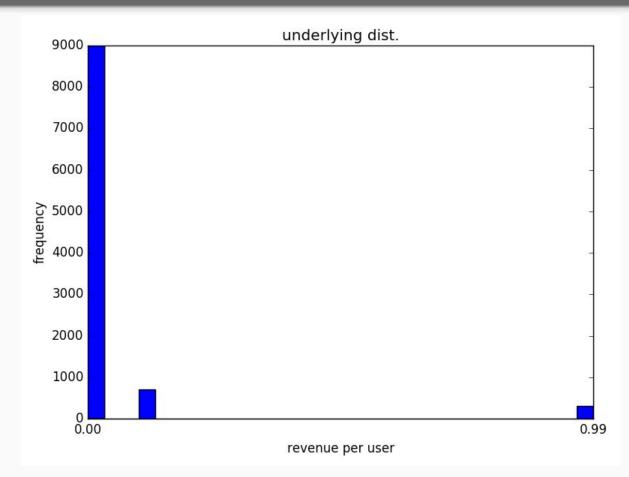
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Distribution of website revenue per visitor



Underlying Distribution:

Random variable: X = revenue per visitor	P(X):
X = \$0.00 (no revenue)	90%
X = \$0.10 (ad-click)	7%
X = \$0.99 (app purchase)	3%



Distribution of sample means

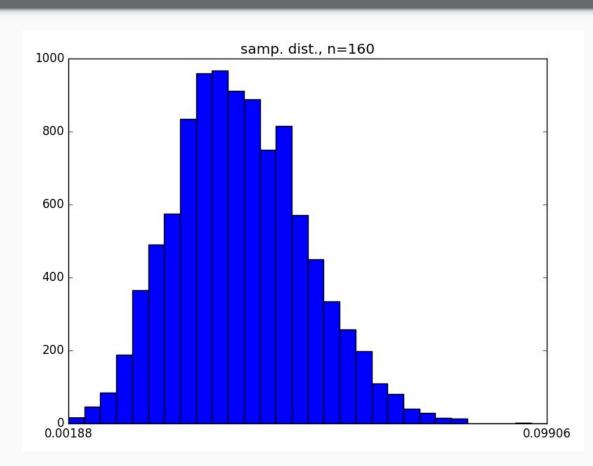


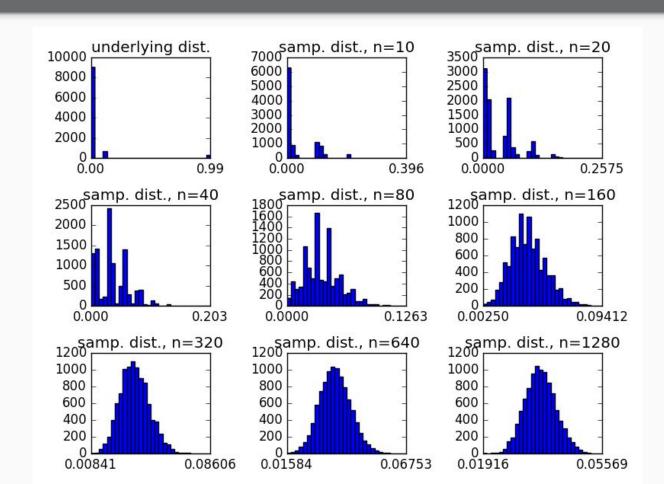
Collect n samples from the website revenue distribution, calculate the sample mean $\overline{\mathcal{X}}$

Repeat 10,000 times, we get:

$$\bar{x}_0, \bar{x}_1, \ldots, \bar{x}_{9999}$$

Plot all 10,000 sample means.





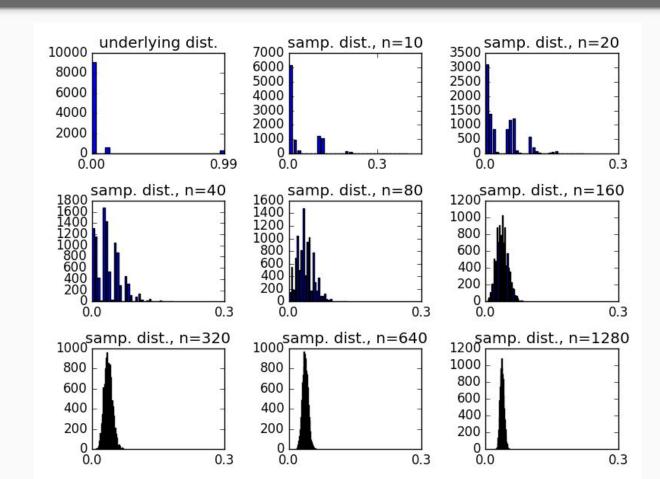
$$\bar{X} \sim N(\mu, \frac{\sigma^2}{n})$$

The distribution of sample means (aka, the "sampling distribution") is normally distributed. *

* Under certain conditions; e.g. sufficiently large sample sizes, and i.i.d. r.v.

Central Limit Theorem: What happens when the sample size increases?





Same charts as the previous slide, but now the scale of each x-axis is the same!

Now we can see: What happens when the sample size increase?

Central Limit Theorem: Std. Dev precise relationship to sample mean



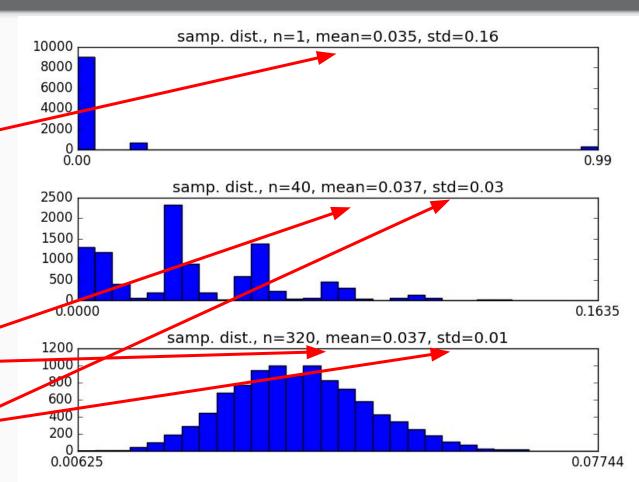
Let the underlying distribution have mean and std. dev.

$$\mu$$
 and σ

The sampling distribution's mean and std. dev. will equal:

$$\mu' = \mu$$

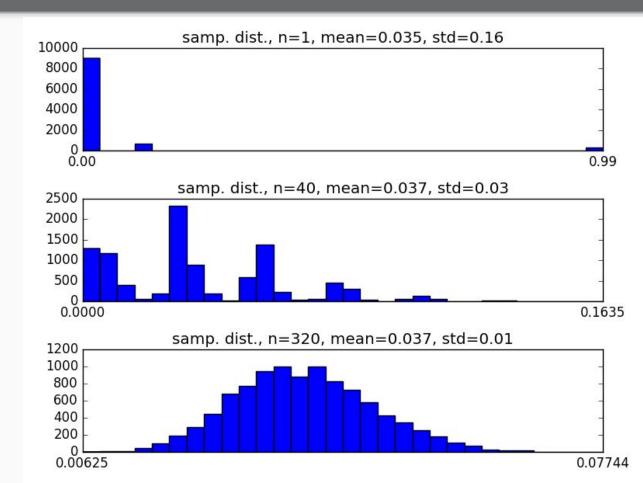
$$\sigma' = \sigma/\sqrt{n}$$





Intuitively, why does the mean stay the same in each histogram?

Intuitively, why does the std. dev. decrease as the sample size increases?



Confidence Intervals

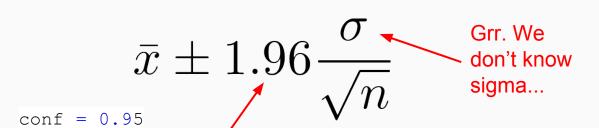
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Confidence Interval

A confidence interval (CI) is an interval estimate of a population parameter.

The typical level of confidence is 95%, but they can be calculated for any level.

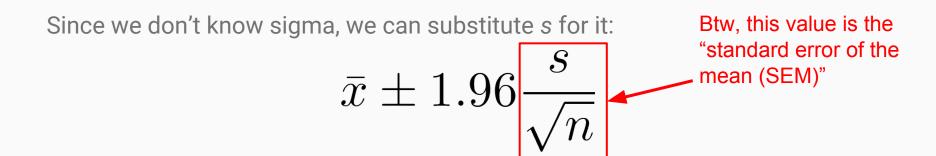
For example, a 95% CI for the population mean is given by:



Why assuming the normal distribution here?

Where does the sqrt(n) come from?

Confidence Interval (con't)



When n is small (<30), we should use the t-distribution instead of the normal:

$$\bar{x} \pm t_{(\alpha/2, n-1)} \frac{s}{\sqrt{n}}$$

Bootstrapping

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Bootstrap Sampling

Estimates the *sampling distribution* of an estimator by sampling with replacement from the original sample.

Recall how is the

Advantages:

Recall, how is the "sampling distribution" distributed when the estimator is the <u>sample average</u>?

- Completely automatic
- Available regardless of how complicated the estimator may be

Often used to estimate the standard errors and confidence intervals of an unknown population parameter.

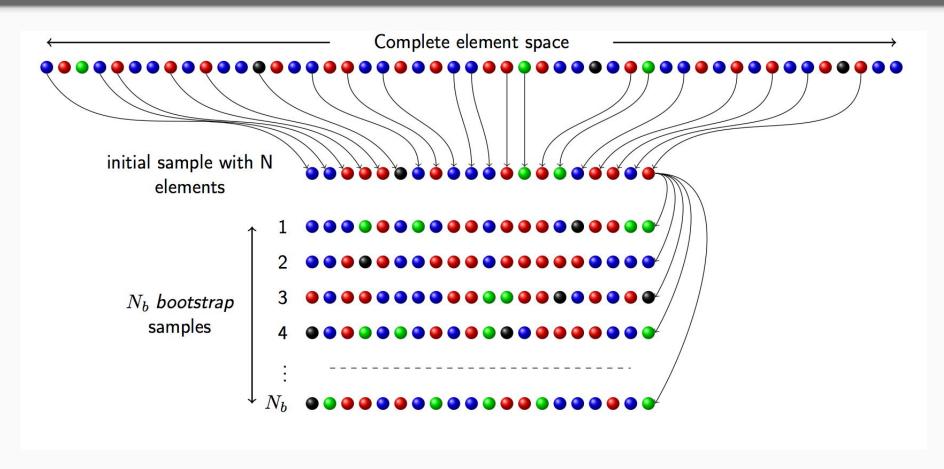


Bootstrap Sampling

Method:

- Start with your dataset of size n
- 2. Sample from your dataset **with replacement** to create 1 bootstrap sample of size *n*
- 3. Repeat *B* times
- 4. Each bootstrap sample can then be used as a separate dataset for estimation and model fitting





Bootstrap Variance



1. Draw a bootstrap sample:

$$X_1^*, X_2^*, \dots, X_n^*$$

2. Calculate bootstrap estimate of your parameter (the parameter you're interested in): $\hat{\theta}^* = t(X_1^*, X_2^*, \dots, X_n^*)$

3. Repeat steps 1 and 2, B times to get:

$$\hat{\theta}_1^*, \hat{\theta}_2^*, \dots, \hat{\theta}_B^*$$

4. Calculate the bootstrapped variance:

$$s_{\rm boot}^2 = \frac{1}{B} \sum_{b=1}^B (\hat{\theta}_b^* - \bar{\theta}^*)^2 \qquad \text{where } \bar{\theta}^* = \frac{1}{B} \sum_{b=1}^B \hat{\theta}_b^*$$

Bootstrap Confidence Intervals

Percentile method using percentile values resulting from bootstrap set

$$(\hat{\theta}_{\alpha/2}^*, \hat{\theta}_{1-\alpha/2}^*)$$

Interval assuming approximately normal bootstrap sampling distribution:

$$\bar{\theta}^* \pm 1.96 \, s_{\text{boot}}$$

Not great because we have to assume a normal distribution. We used bootstrap to not have to assume a distribution!



When to Bootstrap

When the theoretical distribution of the statistic (parameter) is complicated or unknown. (E.g. Median or Correlation, if we estimate the sample mean and sample size large enough, we know we can use normal or t-distribution, no bootstrapping needed)

When the sample size is too small for traditional methods (normality assumption of sampling distribution not accurate)

Favor accuracy over computational cost.