A/B Testing and Beyond

Designed Experiments for Data Scientists





Week 2

Tuesday September 11th, 2018





Outline

- Recap
- Statistical Prerequisites
 - Random Variables and Distributions
 - Point and Interval Estimation
 - Hypothesis Testing





RECAP





Recap

- Response variables
- Factors (3 types) Levels
- Experimental Conditions
- Experimental Units
- Experiments vs. Observational Studies
- QPDAC
- Randomization
- Replication
- Blocking





STATISTICAL PREQUISITES





A random variable is a function that assigns real numbers to the outcomes of a random process.

$$Y:\Omega\to\mathbb{R}$$

We typically denote random variables with upper case letters, such as Y, and the values they can take on with lower case letters, such as y.

We refer to the possible values a random variable can take on as its support set.





- A discrete random variable Y is a random variable that can only take on a finite or countably infinite number of values.
 - Example: y = 0,1,2,...,n or y = 0,1,2,...
 - We typically associate discrete random variables with counting things
- A continuous random variable Y is a random variable that can take on any value in any subinterval of the real numbers.
 - Example: $y \ge 0$ or $y \in [0,1]$ or $-\infty < y < \infty$
 - We typically associate continuous random variables with measuring things





Example 1: Suppose we send an email survey to n=30 individuals and we're interested in the the number of these individuals that respond to the survey. Let Y represent the number of survey responses. In this case the support set is $y=0,1,2,\ldots,30$, and so Y is a discrete random variable.





Example 2: Interest often lies in measuring lifetimes of people, products, and processes. Suppose that, in particular, we are interested in the lifetime of an iPhone's battery. Let Y represent the lifetime (in hours) of an iPhone battery on a single charge. In this case the support set is theoretically $y \ge 0$, which is a continuous subinterval of the real numbers, and so Y is a continuous random variable.





Because Y takes on values randomly, interest lies in quantifying the probability that Y assumes a particular value or lies in some interval

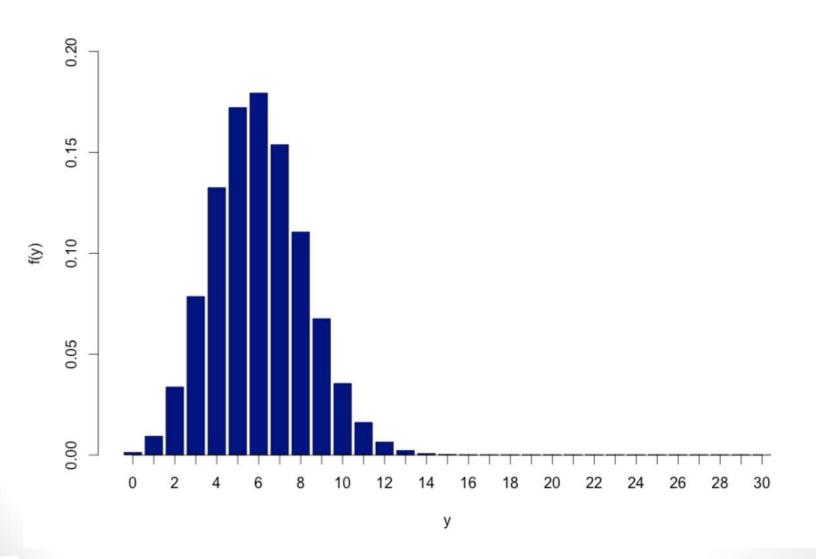
$$P(Y = a) = ?$$
 or $P(a < Y < b) = ?$

In general, questions like these are answered with probability functions f(y). The manner in which they are used depends on whether Y is discrete or continuous.

But in both cases a plot of f(y) vs. y is useful for visualizing the probability distribution

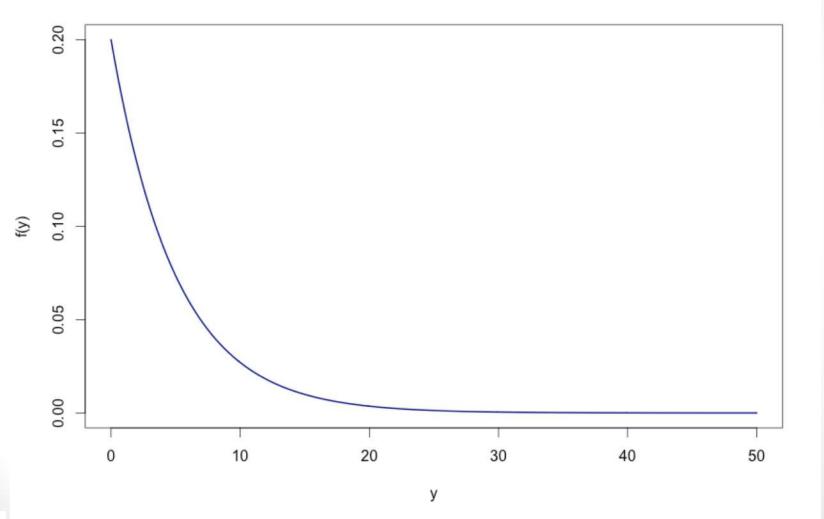
















A probability mass function (PMF) is the name given to the probability function for a discrete random variable Y, and it is defined as

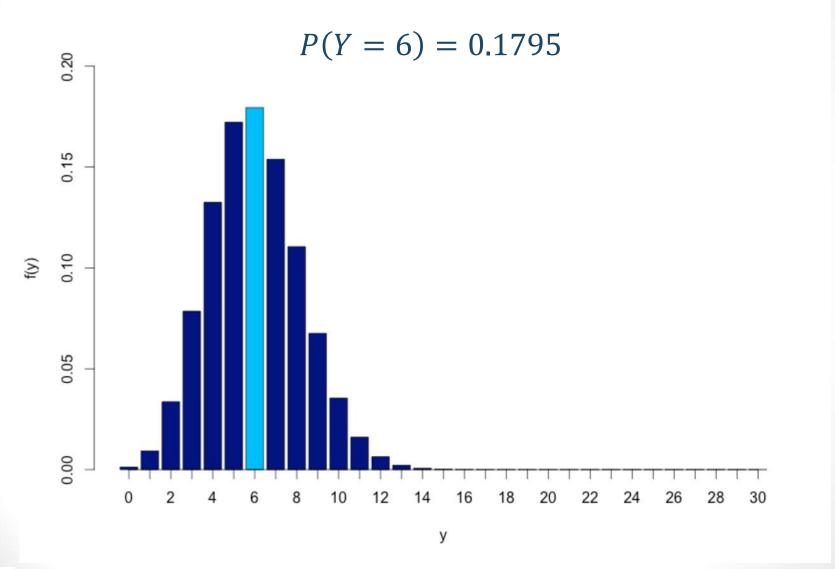
$$f(y) = P(Y = y)$$

for $y \in A$, where the support set A depends on the context of the problem.

- The PMF describes the probabilistic behavior of *Y*;
- It allocates probability to every element of the support set.
- The PMF provides a way to characterize the probability distribution of *Y* .

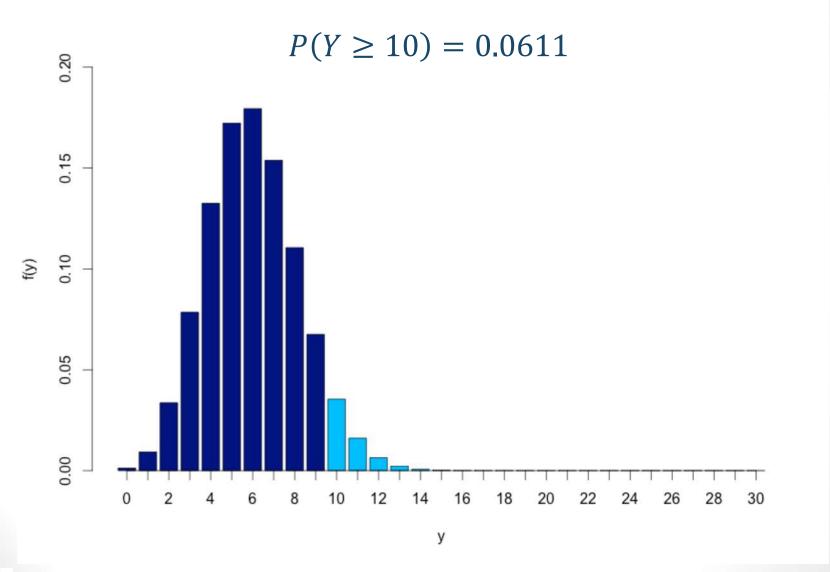
















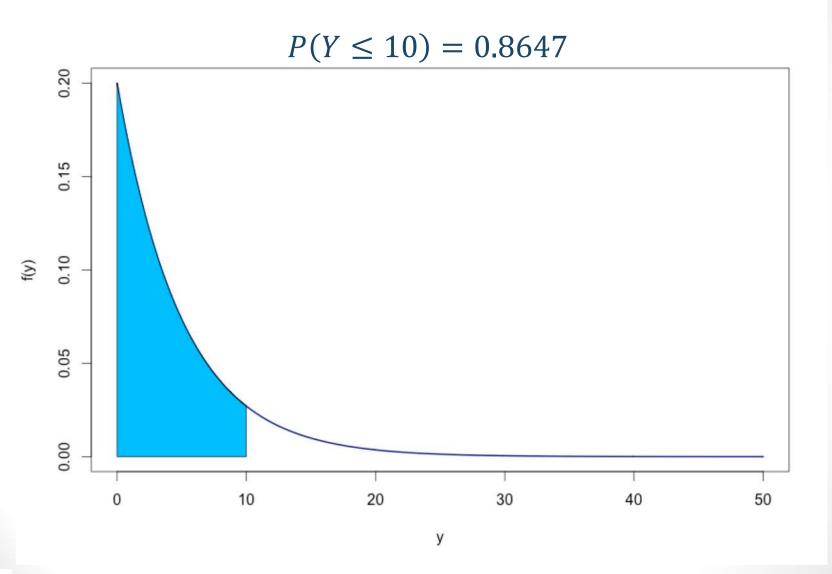
A probability density function (PDF) is the name given to the probability function for a continuous random variable Y, and it is also denoted by

and defined for $y \in A$, where the support set A depends on the context of the problem.

- The difference between a PDF and a PMF lies in how it is used to calculate probabilities.
- If *Y* is a continuous random variable, we calculate probabilities associated with it by calculating areas beneath the associated PDF.

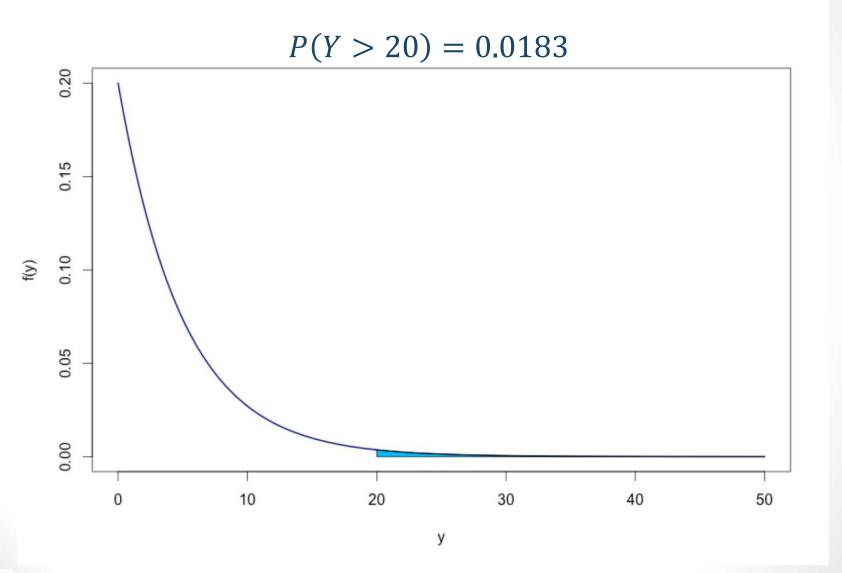
















- Probability distributions are efficiently summarized by plots of the probability function, but it is also useful to characterize a probability function with a closed-form expression.
- Such expressions exist for several well-known probability distributions which are useful for describing a host of real-life random phenomenon.
- We discuss some such distributions, focusing on ones that are used routinely in the context of experimentation.





The Binomial Distribution

 $Y \sim BIN(n, \pi)$ counts the number of "success" in a sequence of n independent Bernoulli trials

$$f(y) = \binom{n}{y} \pi^y (1 - \pi)^{n - y}$$

for y = 0,1,2,...,n and $\pi \in [0,1]$.

In R:

$$P(Y = x) = dbinom(x, size = n, prob = pi)$$

In Python (scipy.stats):

$$P(Y = x) = \text{binom.pmf}(k = x, n = n, p = pi)$$





The Bernoulli Distribution

A $Y \sim BIN(1,\pi)$ random variable is said to have a Bernoulli Distribution. Thus, this is a special case of the binomial distribution when n=1

$$f(y) = \pi^{y} (1 - \pi)^{1 - y}$$

for y = 0.1 and $\pi \in [0.1]$.

In R:

$$P(Y = x) = \text{dbinom}(x, \text{size} = 1, \text{prob} = \text{pi})$$

In Python (scipy.stats):

$$P(Y = x) = \text{binom.pmf}(x, n = 1, p = pi)$$





The Normal Distribution

A $Y \sim N(\mu, \sigma^2)$ distribution has PDF given by

$$f(y) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{\frac{-(y-\mu)^2}{2\sigma^2}}$$

for $-\infty < y < \infty$, $-\infty < \mu < \infty$ and $\sigma > 0$.

In R:

 $P(Y \le q) = pnorm(q, mean = mu, sd = sigma)$ In Python (scipy.stats):

 $P(Y \le q) = \text{norm.cdf}(q, \text{loc} = \text{mu, scale} = \text{sigma})$



The Standard Normal Distribution

This distribution arises as a special case of the normal distribution when $\mu=0$ and $\sigma=1$ and is typically denoted by $Z{\sim}N(0,1)$ with PDF

$$f(z) = \frac{1}{\sqrt{2\pi}} e^{\frac{-z^2}{2}}$$

for $-\infty < z < \infty$.

In R:

$$P(Z \le q) = pnorm(q)$$

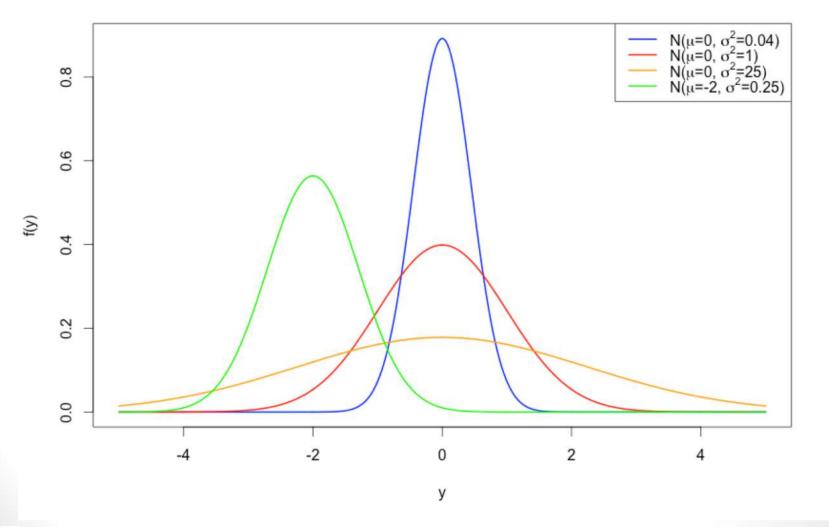
In Python (scipy.stats):

$$P(Y \le q) = \text{norm.cdf}(q)$$





The Normal distribution







The Student's *t*-Distribution

A $Y \sim t_{(v)}$ random variable has PDF given by

$$f(y) = \frac{\Gamma\left(\frac{\nu+1}{2}\right)}{\sqrt{\nu\pi}\Gamma\left(\frac{\nu}{2}\right)} \left(1 + \frac{y^2}{\nu}\right)^{\frac{-(\nu+1)}{2}}$$

for $-\infty < y < \infty$ and $\nu > 0$.

In R:

$$P(Y \le q) = pt(q, df = v)$$

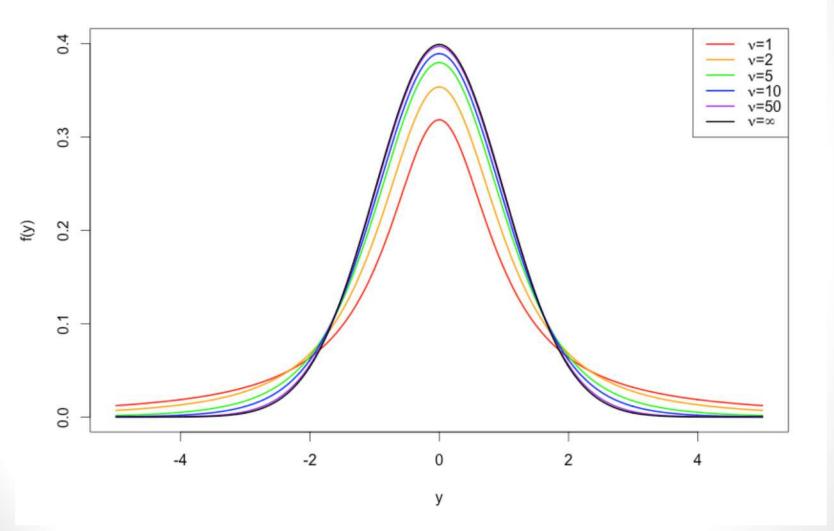
In Python (in scipy.stats):

$$P(Y \le q) = \text{t.cdf}(q, df = v)$$





The *t*-distribution







The Chi-Squared Distribution

A $Y \sim \chi^2_{(\nu)}$ random variable has PDF given by

$$f(y) = \frac{y^{\frac{k}{2}-1}e^{-y/2}}{2^{k/2}\Gamma\left(\frac{\nu}{2}\right)}$$

for $y \ge 0$ and a positive integer ν .

In R:

$$P(Y \le q) = \text{pchisq}(q, \text{df} = v)$$

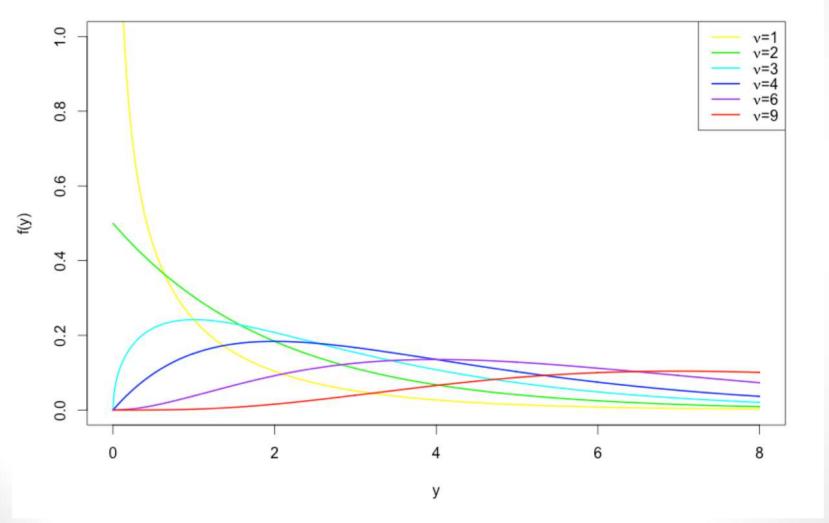
In Python (in scipy.stats):

$$P(Y \le q) = \text{chi2.cdf}(q, \text{df} = v)$$





The Chi-squared distribution







The *F*-distribution

A $Y \sim F(v_1, v_2)$ random variable has PDF given by

$$f(y) = \frac{\Gamma(\frac{\nu_1 + \nu_2}{2})}{\Gamma(\frac{\nu_1}{2})\Gamma(\frac{\nu_2}{2})} \left(\frac{\nu_1}{\nu_2}\right)^{\frac{\nu_1}{2}} y^{\frac{\nu_1}{2} - 1} \left(1 + \frac{\nu_1}{\nu_2} y\right)^{\frac{-\nu_1 + \nu_2}{2}}$$

for $y \ge 0$ and where v_1 and v_2 are positive integers.

In R:

$$P(Y \le q) = pf(q, df1 = v1, df2 = v2)$$

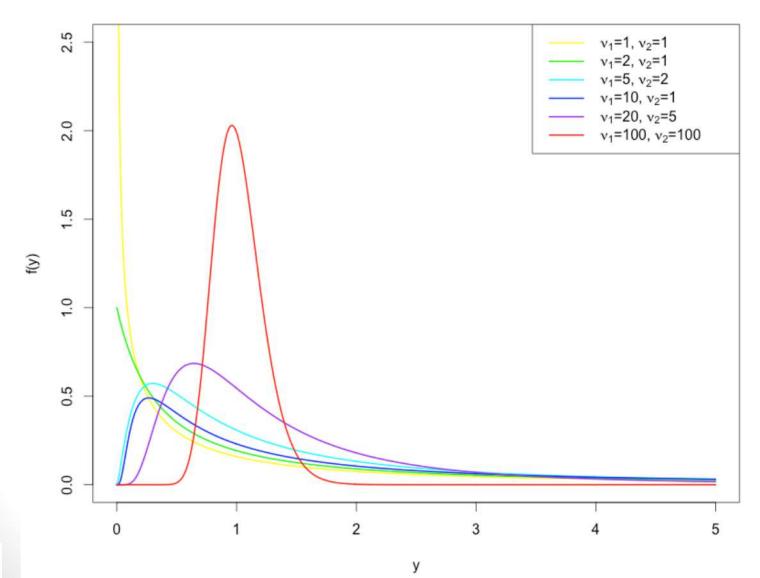
In Python (in scipy.stats):

$$P(Y \le q) = \text{f.cdf}(q, \text{dfn} = \text{v1, dfd} = \text{v2})$$





The *F*-distribution







- The general shapes of these distributions are dictated by their PDFs, and by the parameters that govern them.
- However, the shape of ANY probability distribution, discrete or continuous, can be described by its moments.
- But to understand moments, we must first understand expected values.





- The expected value of a random variable Y, denoted $\mathrm{E}[Y]$, is thought of as:
 - the "average" value of the random variable
 - the "most typical" outcome of the random process that governs Y
 - a measure of a probability distribution's "center"
- Mathematically, the expected value of Y is defined as:
 - $E[Y] = \sum_{all \ y} yf(y)$ if Y is discrete, and
 - $E[Y] = \int_{all \ v} y f(y) dy$ if Y is continuous, and





- We then define the k^{th} moment of Y, or equivalently, the k^{th} moment of Y's distribution, as $\mathrm{E}[Y^k]$.
 - 1st moment: E[Y] quantifies center
 - 2^{nd} moment: $E[Y^2]$ quantifies spread
 - 3^{rd} moment: $E[Y^3]$ quantifies skewness
 - 4^{th} moment: $E[Y^4]$ quantifies kurtosis
- However, the first two moments are used most frequently in practice to describe a distribution's shape





• While the 2^{nd} moment $E[Y^2]$ itself provides information about the dispersion of a distribution, it is most commonly used in the calculation of the variance of Y:

$$Var[Y] = E[(Y - E[Y])^{2}]$$

= $E[Y^{2}] - E[Y]^{2}$

- The variance is interpreted as the expected squared deviation from the mean
- Note that the dispersion of a distribution is also commonly communicated in terms of the standard deviation of *Y*:

$$SD[Y] = \sqrt{Var[Y]}$$





Distribution	E[Y]	Var[Y]
$Y \sim BIN(n, \pi)$	$n\pi$	$n\pi(1-\pi)$
$Y \sim N(\mu, \sigma^2)$	μ	σ^2
$Y \sim t_{(v)}$	0	$\nu/(\nu-2)$
$Y \sim \chi^2_{(\nu)}$	ν	2ν





Statistical Inference

In real life we do not observe probability models, we observe data that have been sampled from some population.

Statistical Inference occurs when one uses sample data to draw conclusions about the population from which the sample was drawn.

If a population can reasonably be modeled by some well-studied probability distribution, then drawing conclusions about it is straightforward.





Statistical Inference

- As we saw previously, much of distribution's information is contained in its shape, and the shape of a given distribution relies entirely on one or more parameters.
- More generally, we think of all statistical models as being governed by one or more unknown parameters.
- Because parameter values are unknown in practice interest lies in
 - Estimating these parameters in light of the observed data
 - ii. Testing hypotheses about the parameters





When a data scientist says that they are fitting a model to some data, what they really mean is:

- They've assumed a certain model or probability distribution is appropriate for describing some characteristic or relationship in a population.
- They have collected data (i.e., a sample from the population) with which they intend to study this characteristic or relationship.
- They intend to use the observed data to estimate the unknown parameters associated with the model or distribution.





The goal of point estimation is to use observed data to obtain reasonable values of a model's unknown parameters that are consistent with the data that were actually observed.

We typically use Greek letters, like θ , to denote parameters and we use $\hat{\theta}$ to denote an estimate of the parameter calculated using sample data.

The notation $\widehat{\theta}$ is read as "theta-hat"





In general, a variety of estimation methods may be used to obtain parameter estimates:

- The Method of Moments
- Maximum Likelihood Estimation
- Least Squares Estimation.
- etc. ...

All estimation procedures have advantages and disadvantages, and so it is important to choose the one that is appropriate for your data and your problem.





However, there is no guarantee that $\hat{\theta}$ is anywhere close to the true value of θ .

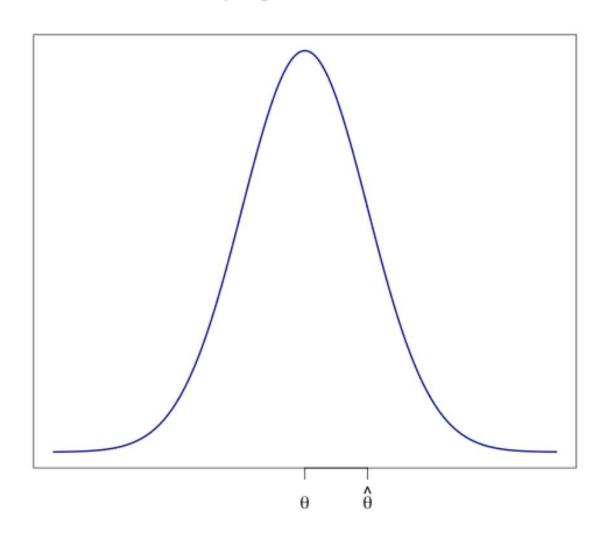
A point estimate of θ is just a single value, based on one sample of data.

If we were to draw a second sample and repeat the exact same estimation procedure we would very likely obtain a slightly different value of $\hat{\theta}$ than before, simply due to sampling variation.





Sampling Distribution of $\boldsymbol{\theta}$







The point estimate $\hat{\theta}$ communicates no information about the sampling distribution on its own.

For this reason we typically accompany point estimates with interval estimates, also know as confidence intervals.

A confidence interval accounts for sampling variation and provides an interval within which we can be confident the true value of θ lies.

Thus a point estimate provides the **best estimate** of θ based on the observed data and a confidence provides a **range of plausible values** for θ accounting for sampling variation.





The most common confidence interval for a parameter θ is a 95% confidence interval.

We interpret such intervals by saying that we are 95% confident that the true value of θ is contained in this interval.

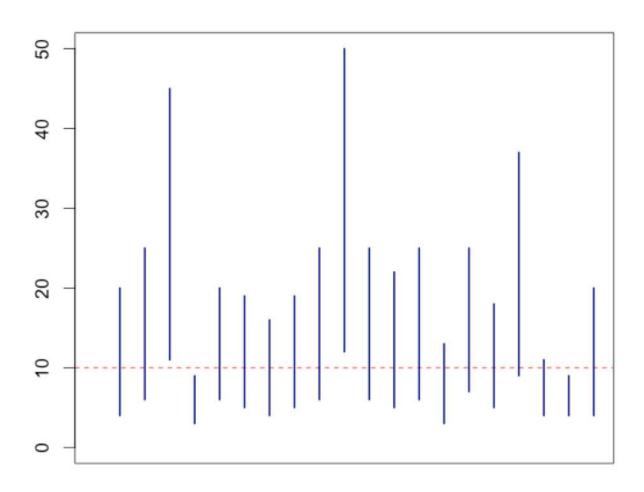
But what does it mean to be "95% confident"?

What this means is that if we were to make infinitely many such intervals, 95% of them would contain θ .





Confidence Interval Illustration







With estimation we knew nothing about the parameter of interest, so we estimated it from observed data.

However, we may have a belief as to what θ is, and if we'd like to formally evaluate this belief, we should perform a hypothesis test.

Formally, a hypothesis is a statement about a parameter that we'd either like to prove or disprove, using the collected data.

A hypothesis is tested by comparing ones own data with a hypothesized statistical distribution.





We typically state such a hypothesis as

$$H_0: \theta = \theta_0 \text{ versus } H_A: \theta \neq \theta_0$$

where

- H_0 is the null hypothesis, and it is the statement we believe to be true, and that we want to test using observed data.
- H_A is the alternative hypothesis, and is the complement of H_0 .

Exactly one of these hypotheses is true, and we use observed data to try and empirically uncover the truth





The previous hypothesis statement corresponded to a two-sided hypothesis.

The following statements correspond to one-sided hypotheses

$$H_0: \theta \leq \theta_0 \text{ versus } H_A: \theta > \theta_0$$

or

$$H_0: \theta \ge \theta_0 \text{ versus } H_A: \theta < \theta_0$$

Which hypothesis statement is appropriate depends on the context of the problem and the question that the hypothesis test is designed to answer





Formally, we use observed data to decide whether to reject or not reject H_0 .

In order to draw such a conclusion, we define a test statistic T which is a random variable that satisfies three properties

- it must be a function of the observed data
- it must be a function of the parameter θ
- its distribution must not depend on θ

Then, assuming the null hypothesis is true, the test statistic T follows a particular distribution which we call the null distribution





Next we calculate t, the observed value of the test statistic, by substituting the observed data and θ_0 into the expression for T.

Note that expressions for *T* typically incorporate terms of the form:

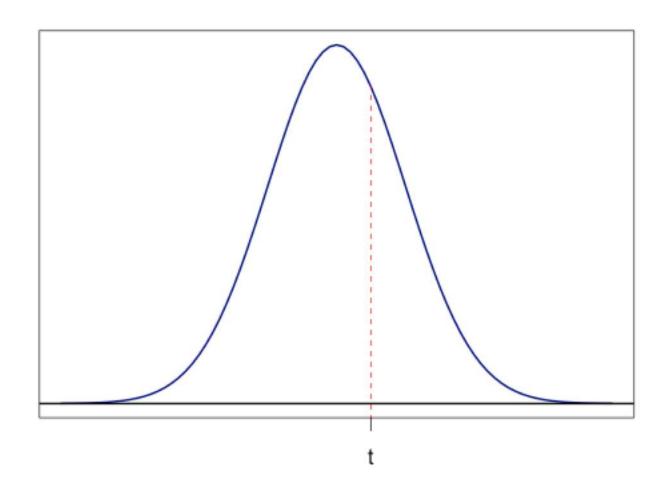
$$\hat{\theta} - \theta_0$$
 or $\hat{\theta}/\theta_0$

Then we evaluate the extremity of *t* relative to the null distribution:

- If t seems very extreme, this provides evidence against H_0
- If t seems reasonable, this provides evidence in favor of H_0

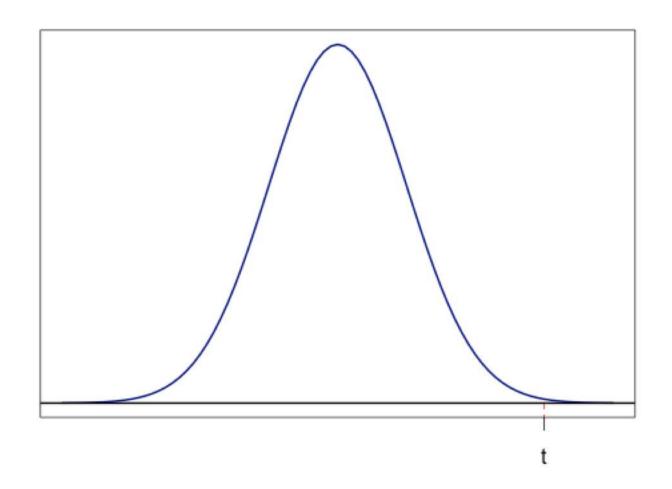
















We formalize the extremity of *t* using the p-value of the test.

The p-value is defined to be:

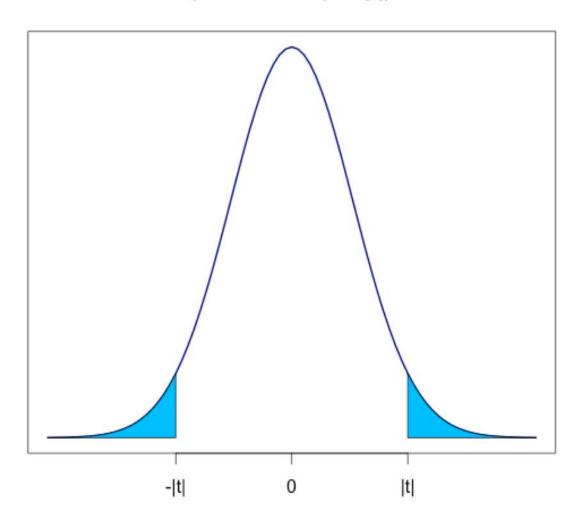
The probability of observing a value of the test statistic at least as extreme as the value we observed, if the null hypothesis is true.

Whether large values of t, small values of t, or both, are to be considered extreme depends on whether H_A is one- or two-sided





$$H_0$$
: $\theta = \theta_0$ versus H_A : $\theta \neq \theta_0$
p-value = 2P(T > |t|)

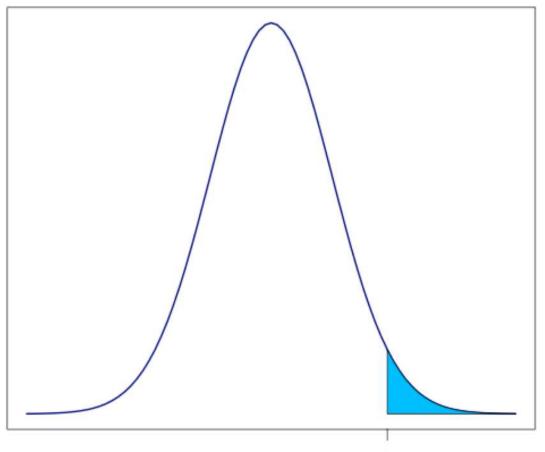






 $H_0: \theta \leq \theta_0 \text{ versus } H_A: \theta > \theta_0$

p-value = P(T > t)

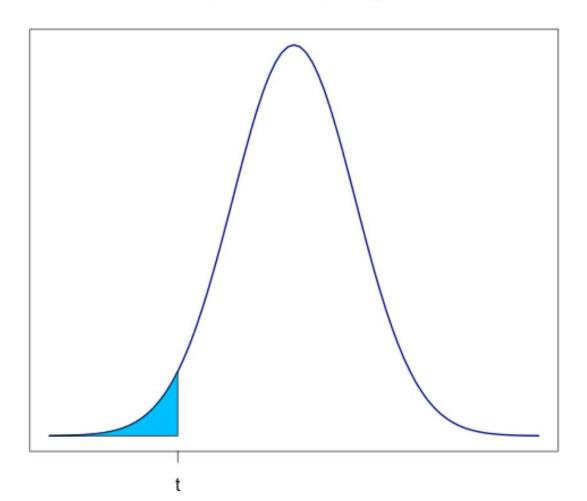






 $H_0: \theta \ge \theta_0 \text{ versus } H_A: \theta < \theta_0$

p-value = P(T < t)







How "extreme" t must be, and hence how small the p-value must be, to reject H_0 is determined by the significance level of the test, which we denote by α .

In particular, if

- p-value $\leq \alpha$ we reject H_0 in favor of H_A
- p-value $> \alpha$ we do not reject H_0

Note that $\alpha = 0.01$ and 0.05 are common choices.





In drawing such a conclusion, it is possible that we make one of two types of errors:

- Type I Error: based on the observed data we reject H_0 when it is in fact true
- Type II Error: based on the observed data we fail to reject H_0 when it is in fact false.

It is important to note that there is typically an imbalance in the consequences associated with these two types of error.





Courtroom Analogy

 H_0 : defendant is innocent vs. H_A : defendant is guilty

- Type I Error: the defendant is truly innocent, but the evidence leads the jury to find the defendant guilty
- Type II Error: the defendant is truly guilty, but the evidence leads the jury to find the defendant innocent

In any hypothesis testing setting, both types of errors lead to negative outcomes, but these negative outcomes may be prioritized differently





Clearly we would like to reduce the likelihood of either type of error happening.

We define

$$\alpha = P(\text{Type I Error}) \text{ and } \beta = P(\text{Type II Error})$$

which reflect the chances that a Type I or Type II error will occur.

We call α the significance level of the test and $1-\beta$ the power of the test.

Thus a test with a **small significance level** and **large power** is desirable.





In practice we choose α and β based on how often we are comfortable allowing Type I and Type II errors to occur.

Suppose we can tolerate a Type I error 1% of the time and a Type II error 5% of the time.

In this case we would choose $\alpha=0.01$ and $\beta=0.05$ and the significance level of the test would be 1% and the power of the test would be 95%

Common choices for significance level and power are respectively 5% and 80%, corresponding to $\alpha=0.05$ and $\beta=0.2$





Thus the significance level α (i.e., the probability of making a Type I error), determines how small a p-value must be in order to reject a null hypothesis.

Next week we will see that for a fixed value of α the desired power determines the required sample size.

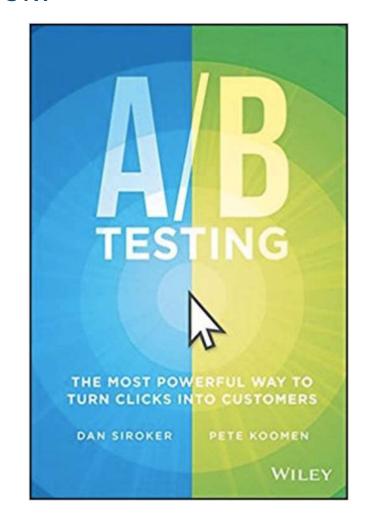
The choices for α and β should be made prior to testing the hypothesis and in fact prior to collecting any data.





Take Home Task

Read this book!







See you next week!



