



Denotes  
He-man



Denotes  
practical

# Power and p-values



**SHE-RA: PRINCESS OF POWER**  
Members of the He-Man, She-Man, Princess of Power, She-Ra, and the Judo Kids.  
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**FILMATION**  
GROUP W  
PRODUCTIONS

Benjamin Neale  
March 10<sup>th</sup>, 2016  
International Twin Workshop, Boulder, CO



SCIENTIFIC METHOD | 10:23 AM | MAR 7, 2016



# **Statisticians Found One Thing They Can Agree On: It's Time To Stop Misusing P-Values**

By CHRISTIE ASCHWANDEN

**p-values are in the press**

In February, 2014, George Cobb, Professor Emeritus of Mathematics and Statistics at Mount Holyoke College, posed these questions to an ASA discussion forum:

Q: Why do so many colleges and grad schools teach  $p = .05$ ?

A: Because that's still what the scientific community and journal editors use.

Q: Why do so many people still use  $p = 0.05$ ?

A: Because that's what they were taught in college or grad school.

**What we've been teaching**

- A way to ask if the data are consistent with a null model

**What exactly is a p-value?**

- The baseline model for comparison, usually no effect [e.g. no heritability]

**What's a null model?**



- Distrust of his aunt's claims of being able to discriminate between milk in first or tea in first

**Whose fault is it anyway?**



alternative vs      null  
some effect      no effect

**Hypothesis testing**

## Statistics

	Reject $H_0$	Fail to reject $H_0$
Truth		
$H_0$ is true	$\alpha$	$1-\alpha$
$H_a$ is true	$1-\beta$	$\beta$

$\alpha$  = type 1 error rate

$\beta$  = type 2 error rate

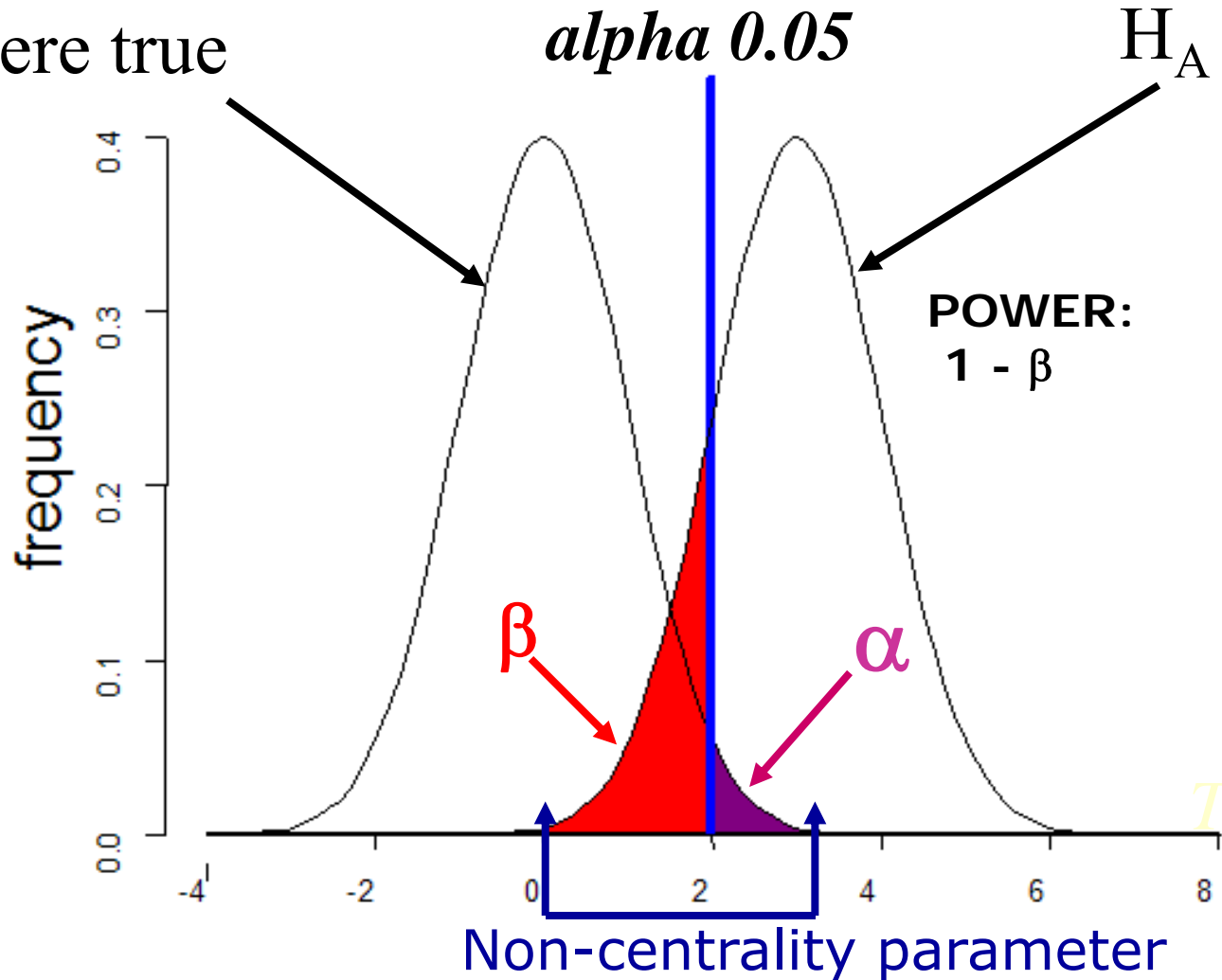
$1-\beta$  = statistical power

## Possible scenarios

# Standard Case

Sampling distribution if  $H_0$  were true

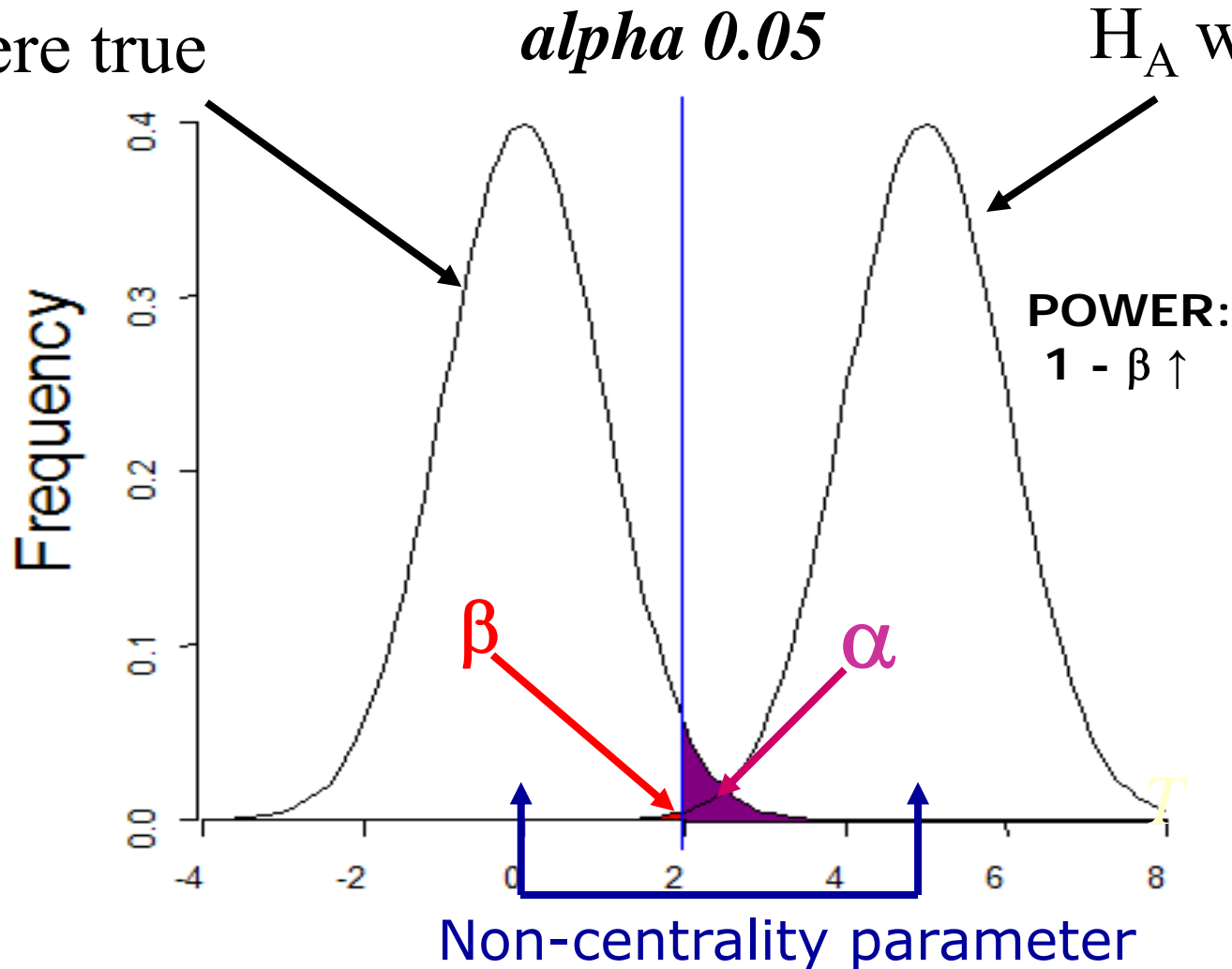
Sampling distribution if  $H_A$  were true



# Increased effect size

Sampling distribution if  $H_0$  were true

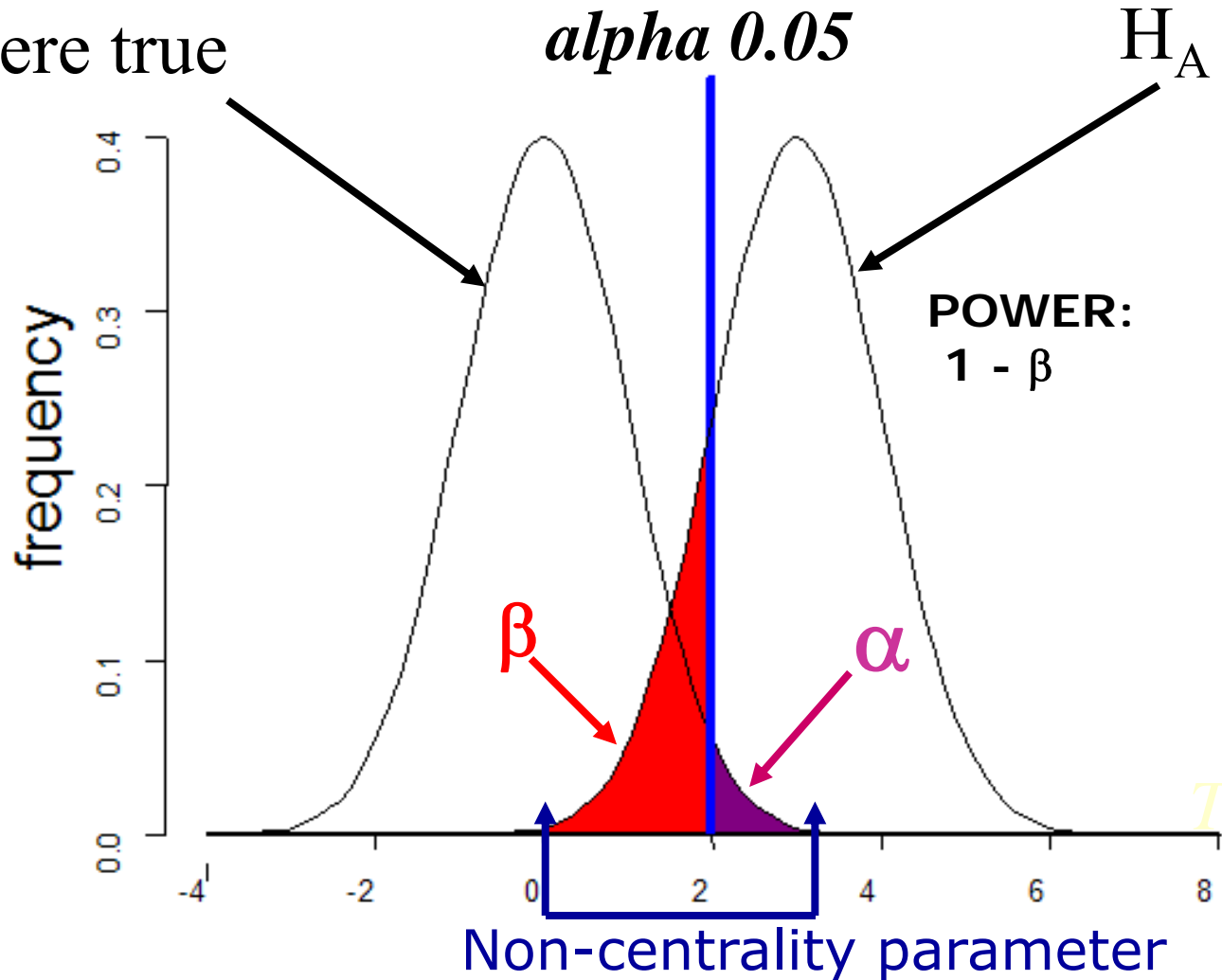
Sampling distribution if  $H_A$  were true



# Standard Case

Sampling distribution if  $H_0$  were true

Sampling distribution if  $H_A$  were true

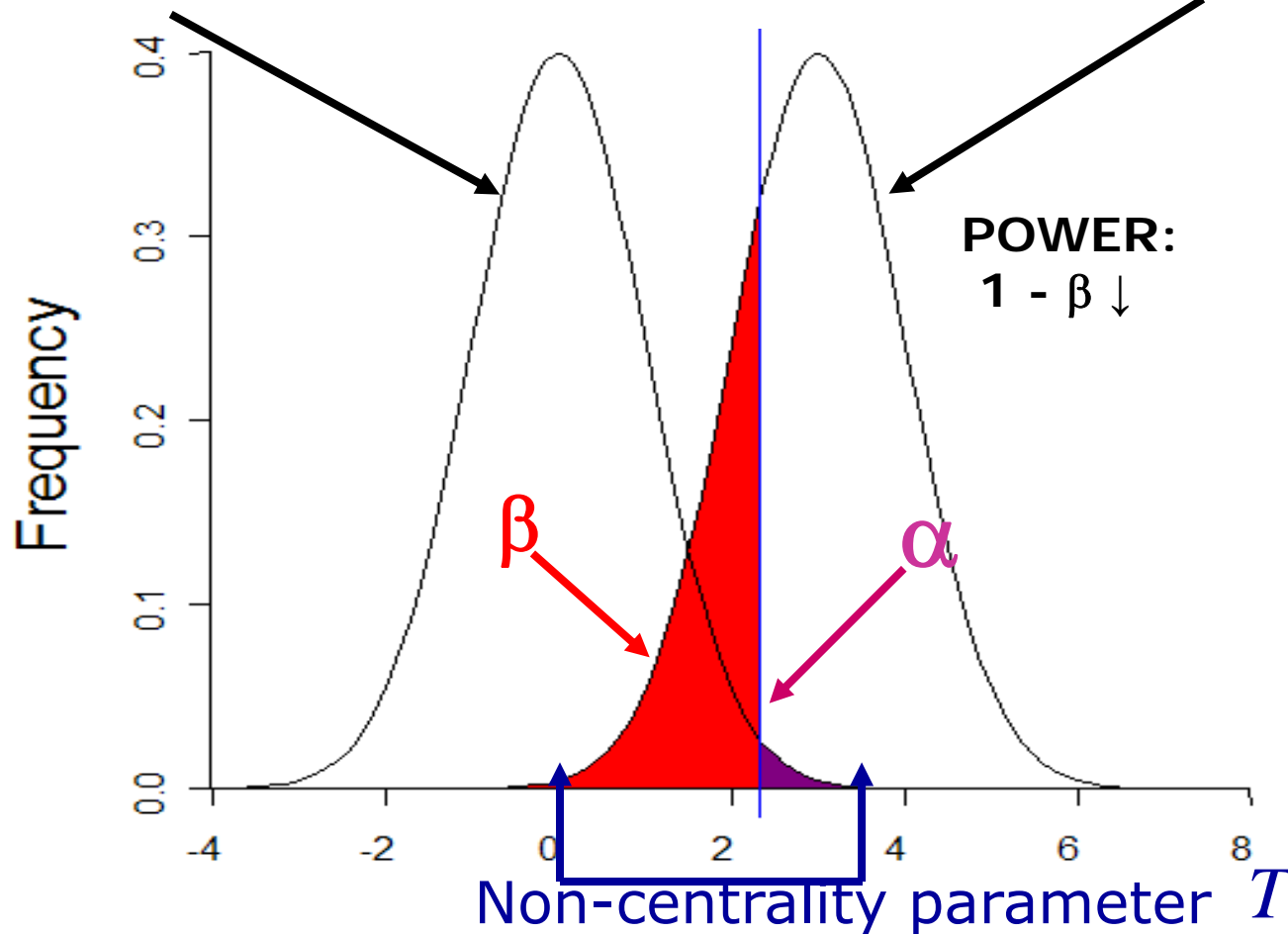


# More conservative $\alpha$

Sampling distribution if  $H_0$  were true

Sampling distribution if  $H_A$  were true

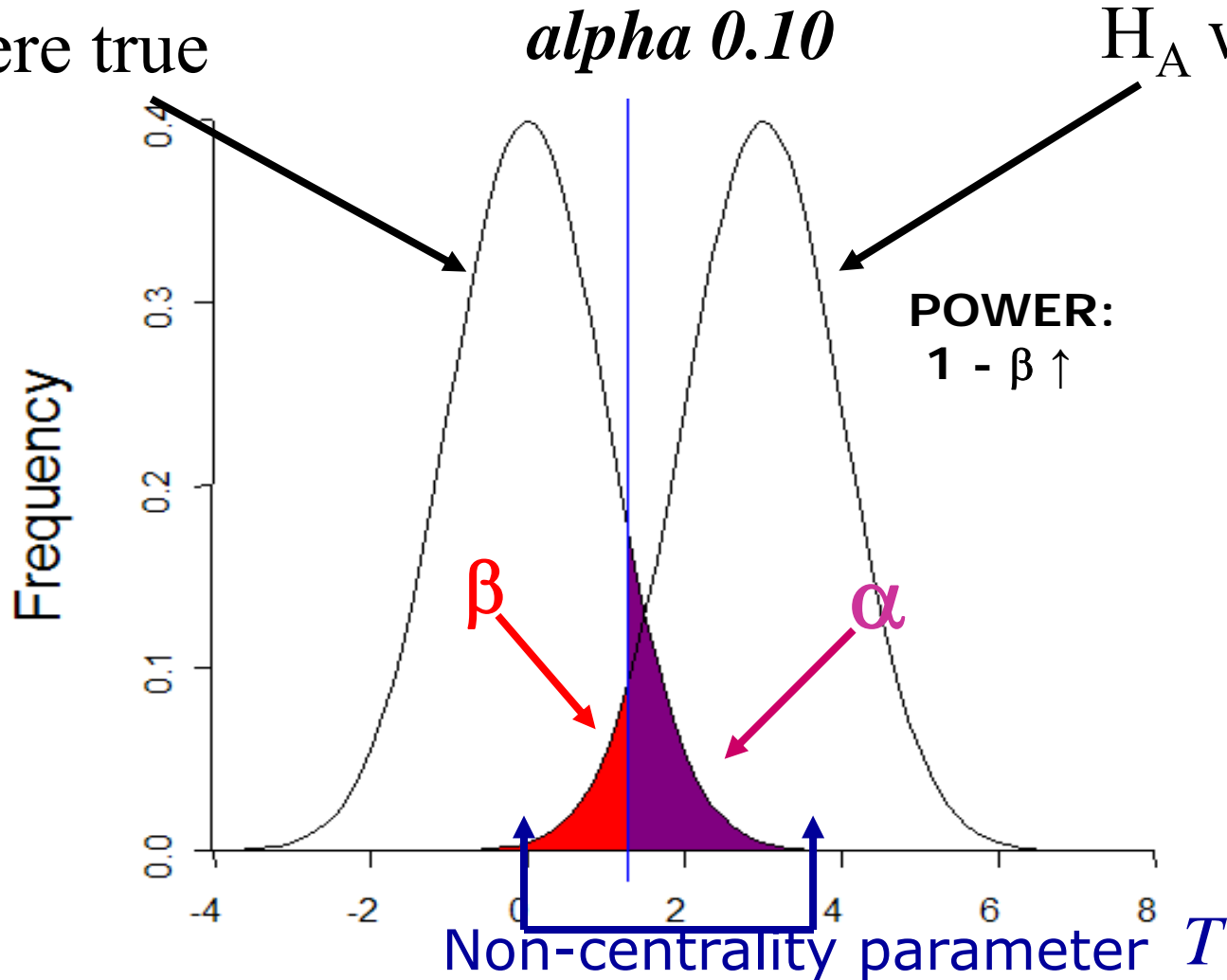
*alpha 0.01*



# Less conservative $\alpha$

Sampling distribution if  $H_0$  were true

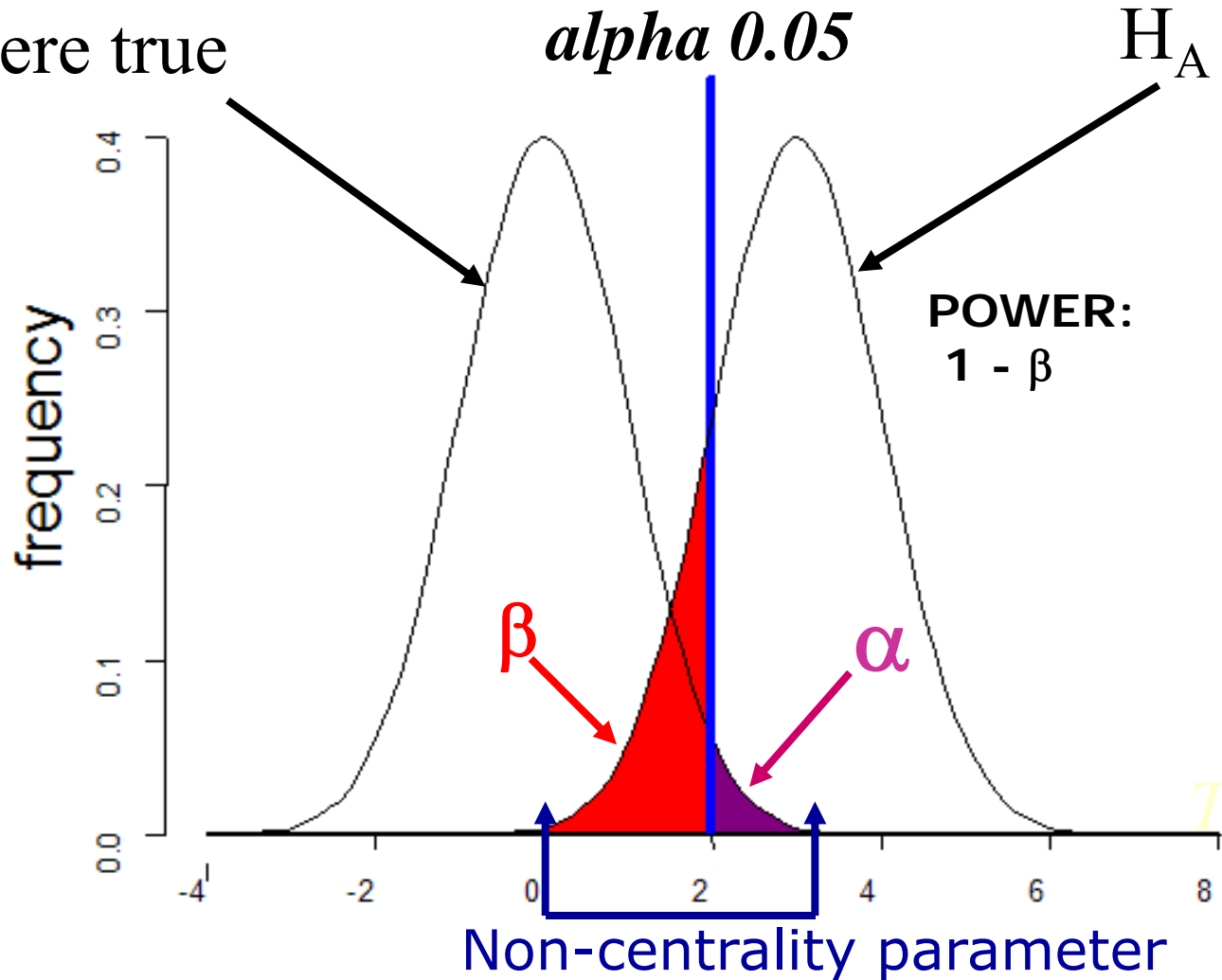
Sampling distribution if  $H_A$  were true



# Standard Case

Sampling distribution if  $H_0$  were true

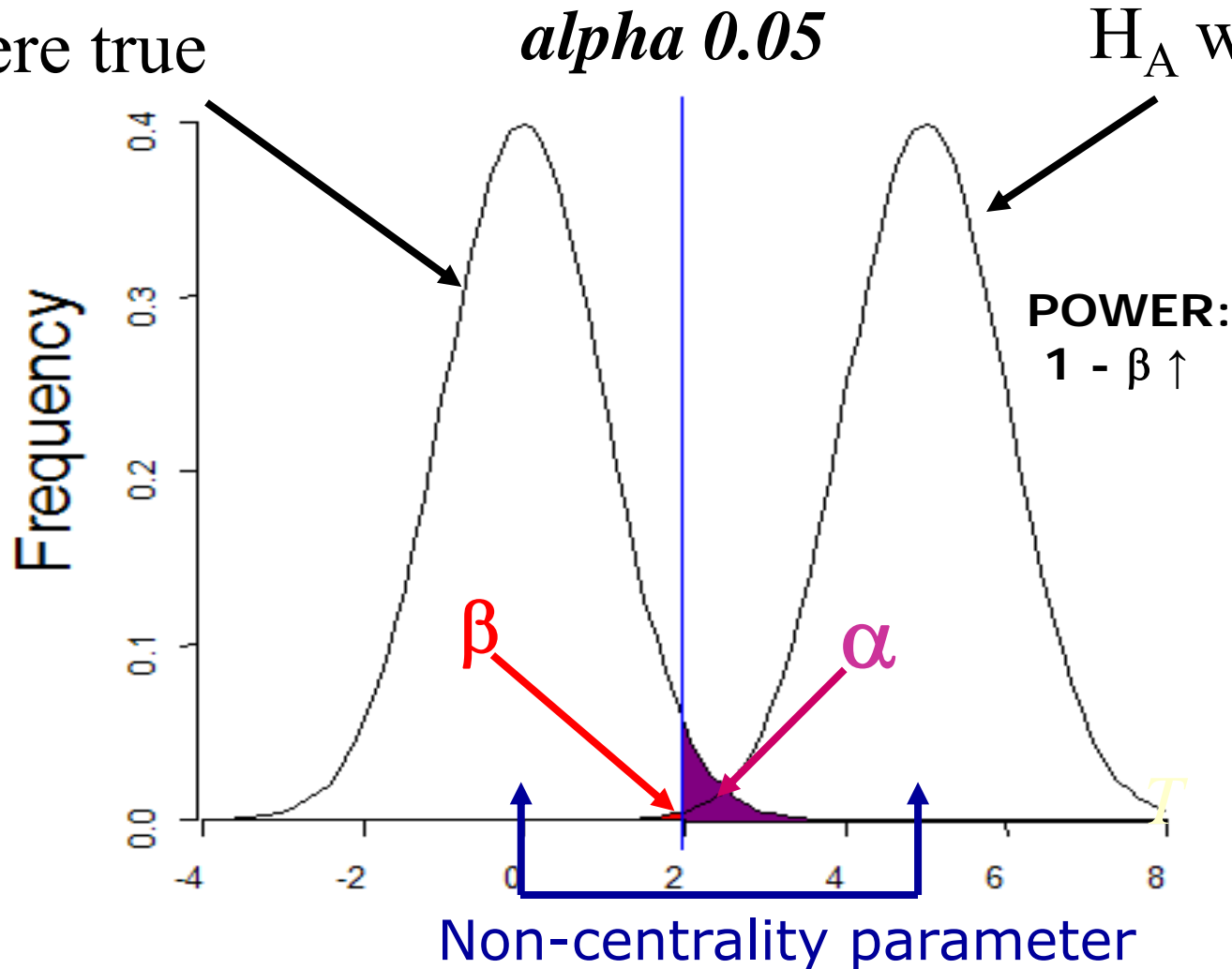
Sampling distribution if  $H_A$  were true



# Increased sample size

Sampling  
distribution if  
 $H_0$  were true

Sampling  
distribution if  
 $H_A$  were true

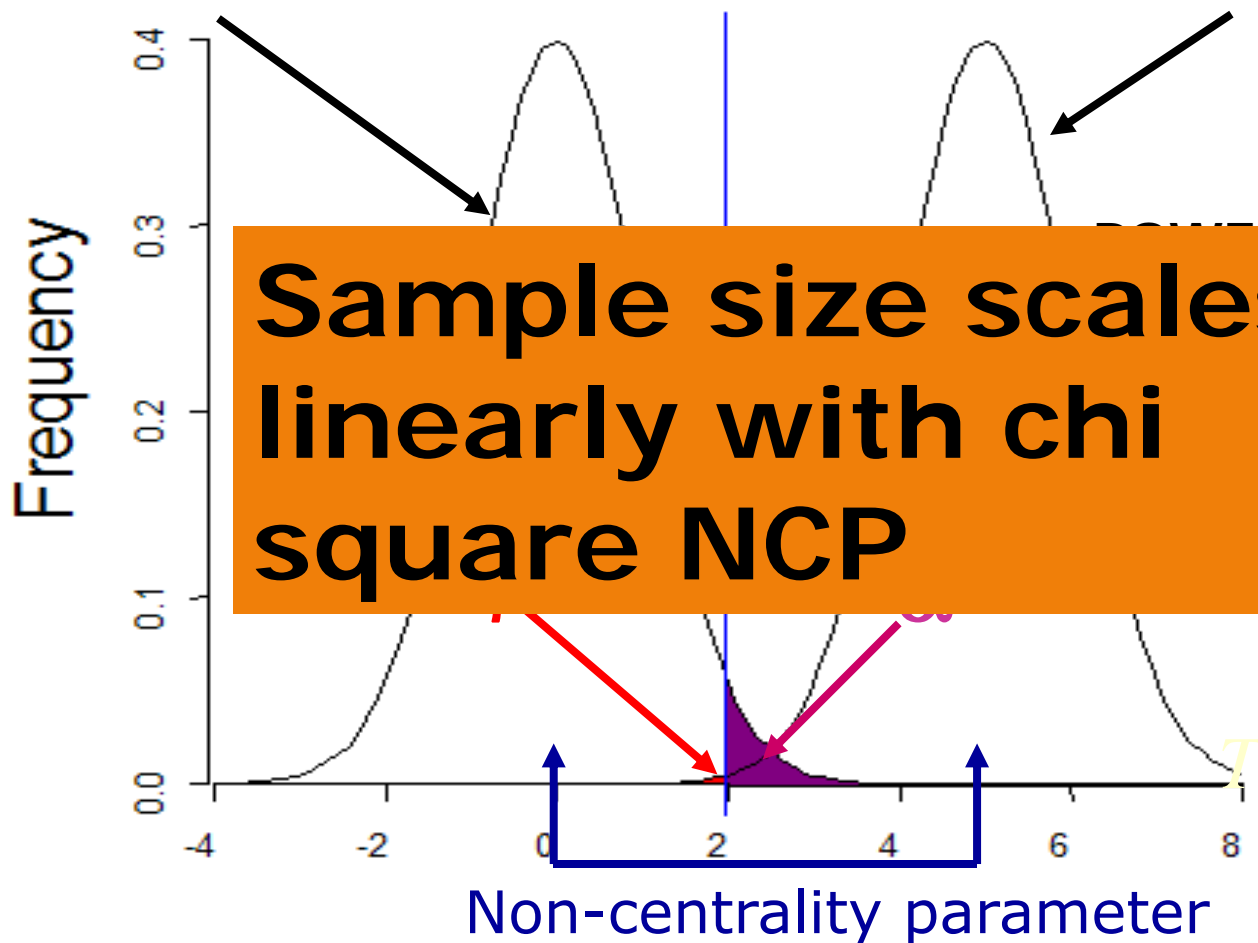


# Increased sample size

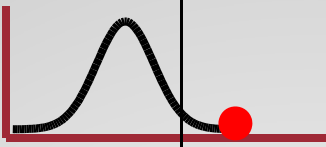
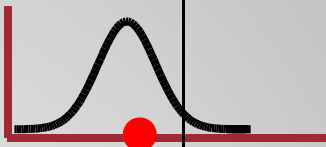
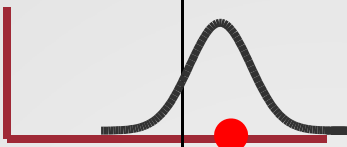
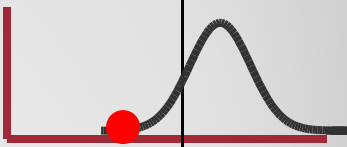
Sampling  
distribution if  
 $H_0$  were true

Sampling  
distribution if  
 $H_A$  were true

*alpha 0.05*



# Statistical Analysis

		Rejection of $H_0$	Non-rejection of $H_0$
Truth	$H_0$ true	<p>Type I error at rate <math>\alpha</math></p> 	<p>Nonsignificant result (<math>1 - \alpha</math>)</p> 
	$H_A$ true	<p>Significant result (<math>1 - \beta</math>)</p> 	<p>Type II error at rate <math>\beta</math></p> 

- Definitions of power
- The probability that the test will reject the null hypothesis if the alternative hypothesis is true
- The chance the your statistical test will yield a significant result when the effect you are testing exists

**What is power?**

- We are going to simulate a normal distribution using R
- We can do this with a single line of code, but let's break it up

## Practical 1

- R has functions for many distributions
- Normal,  $\chi^2$ , gamma, beta (others)
- Let's start by looking at the random normal function: `rnorm()`

**Simulation functions**

The image shows a screenshot of an R Help window. The window title is "R Help". At the top, there are navigation buttons: a left arrow, a right arrow, and a "Print" button. On the right side, there is a search bar labeled "Help Search". The main content area displays the following text:

Normal {stats} R Documentation

**The Normal Distribution**

**Description**

Density, distribution function, quantile function and random generation for the normal distribution with mean equal to `mean` and standard deviation equal to `sd`.

**Usage**

```
dnorm(x, mean = 0, sd = 1, log = FALSE)
pnorm(q, mean = 0, sd = 1, lower.tail = TRUE, log.p = FALSE)
qnorm(p, mean = 0, sd = 1, lower.tail = TRUE, log.p = FALSE)
rnorm(n, mean = 0, sd = 1)
```

**Arguments**

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations. If <code>length(n) &gt; 1</code> , the length is taken to be the number required.
<code>mean</code>	vector of means.
<code>sd</code>	vector of standard deviations.

In R: ?rnorm

# rnorm Documentation

- `rnorm(n, mean = 0, sd = 1)`



Function name



Number of observations to simulate



Mean of distribution  
with default value



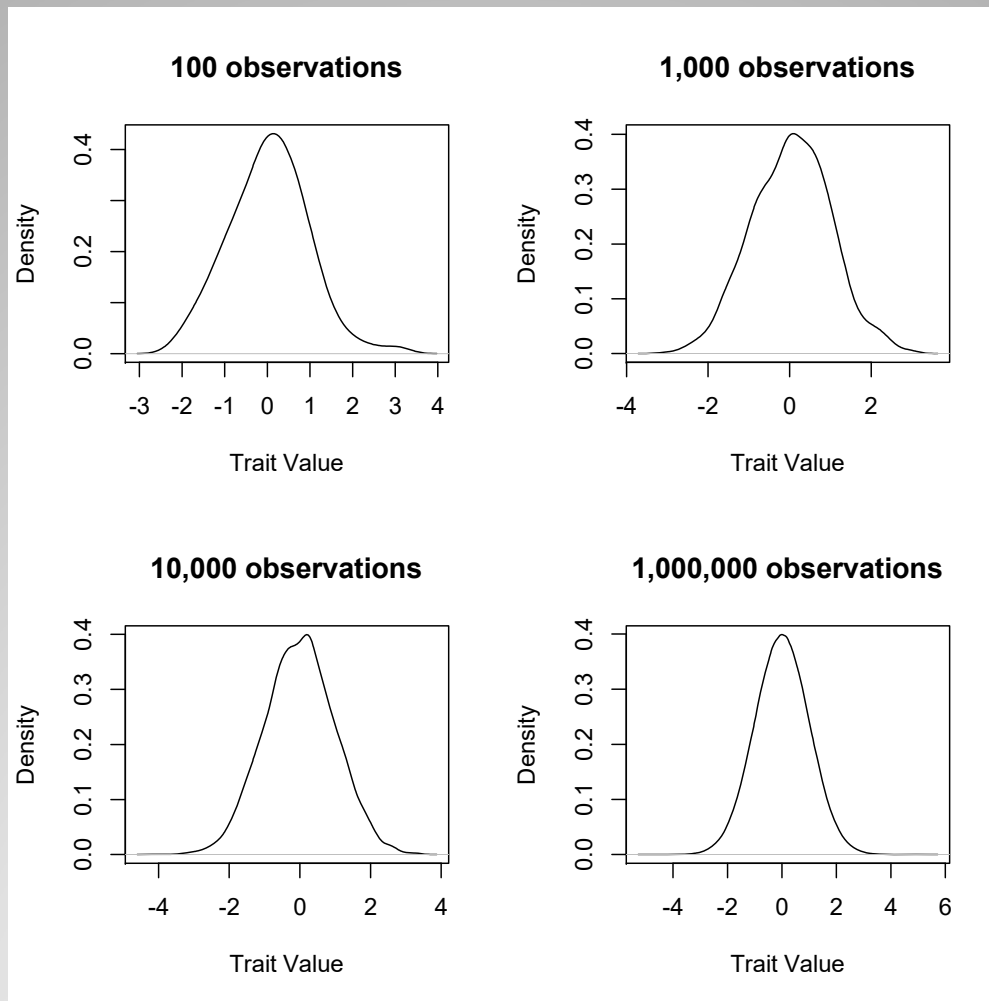
Standard deviation of distribution  
with default value

**rnorm syntax**

- This script will plot 4 samples from the normal distribution
- Look for changes in shape
- Thoughts?



**R script: Norm\_dist\_sim.R**



One I made earlier

- Sampling variance
  - We saw that the 'normal' distribution from 100 observations looks stranger than for 1,000,000 observations
- Where else may this sampling variance happen?
- How certain are we that we have created a good distribution?

Concepts

- Rather than just simulating the normal distribution, let's simulate what our estimate of a mean looks like as a function of sample size
- We will run the R script `mean_estimate_sim.R`

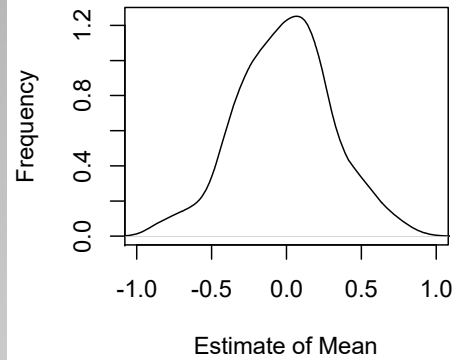
**Mean estimation**

- This script will plot 4 samples from the normal distribution
- Look for changes in shape
- Thoughts?

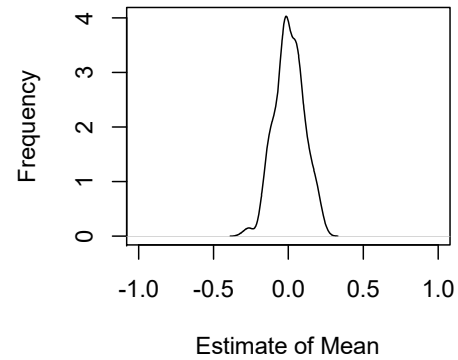


**R script: mean\_estimate\_sim.R**

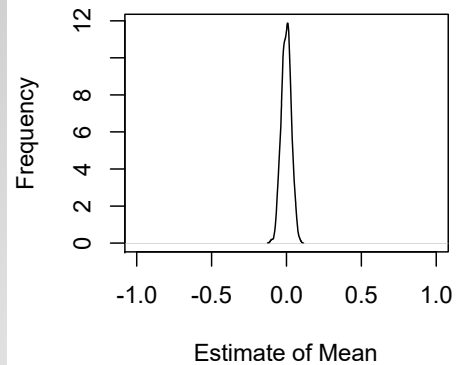
**10 sample size mean estimate**



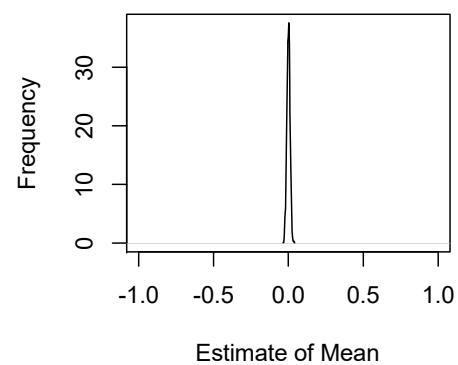
**100 sample size mean estimate**



**1,000 sample size mean estimate**



**10,000 sample size mean estimate**



**One I made earlier**

- We see an inverse relationship between sample size and the variance of the estimate
- This variability in the estimate can be calculated from theory
- $SE_x = s/\sqrt{n}$
- $SE_x$  is the standard error,  $s$  is the sample standard deviation, and  $n$  is the sample size

## Standard Error

The sampling variability in my estimate affects my ability to declare a parameter as significant (or significantly different)

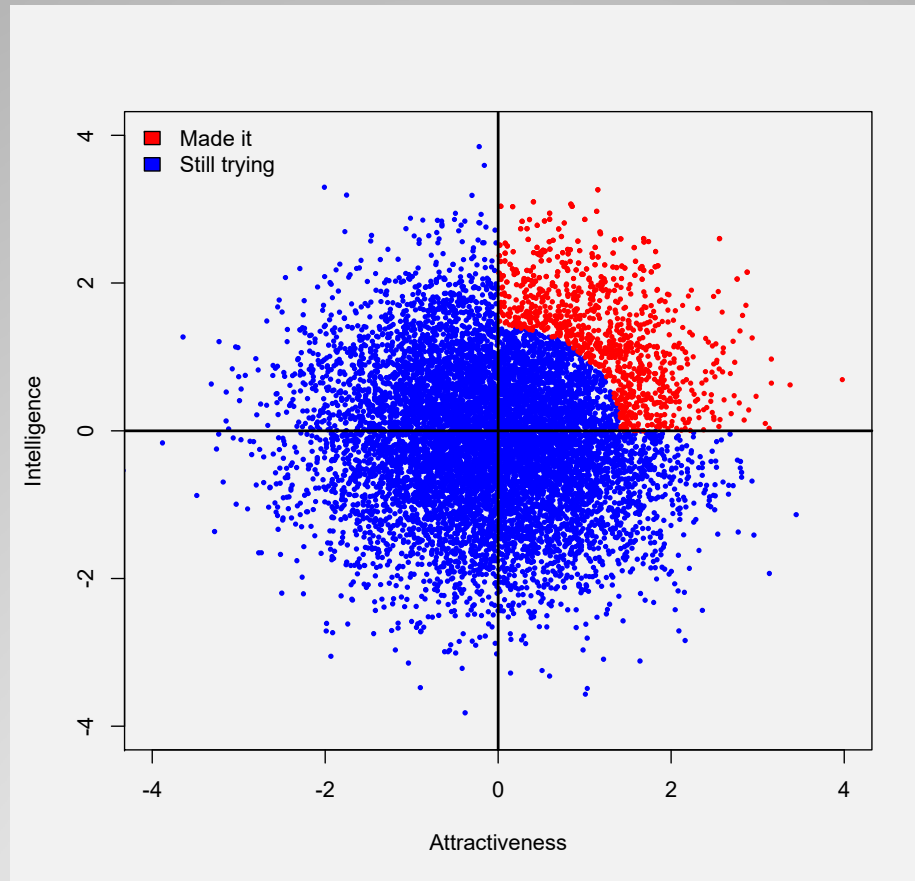
**Key Concept 1**

The probability that the test will reject the null hypothesis if the alternative hypothesis is true

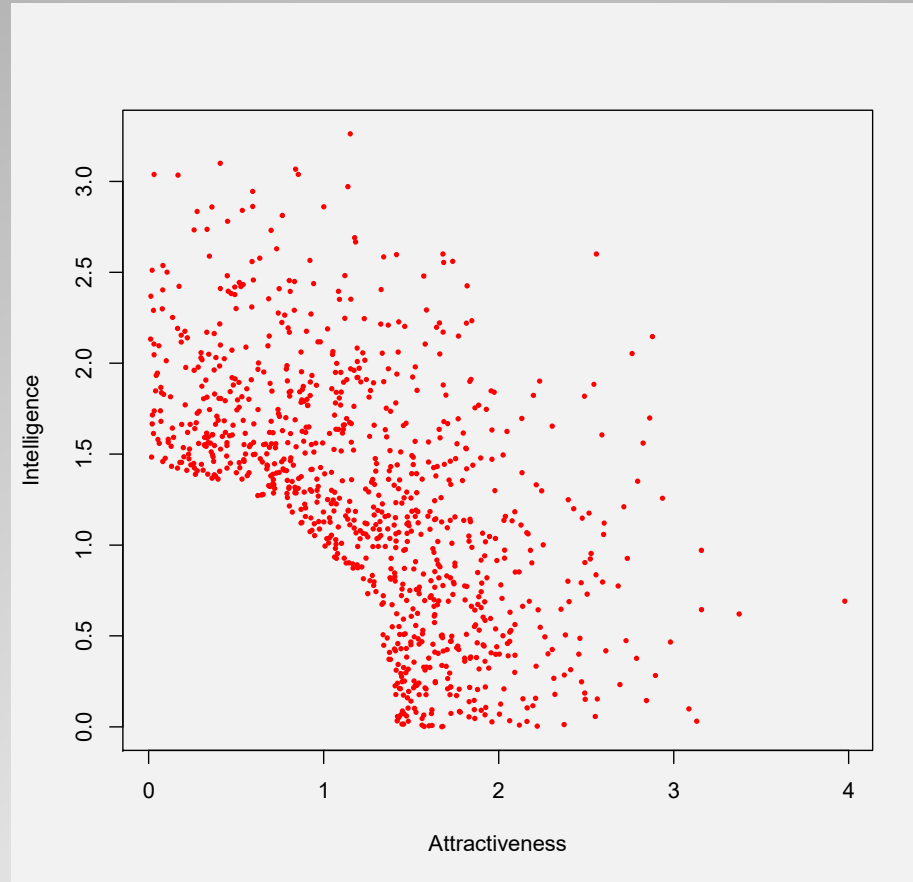
**Power definition again**

# Ascertainment

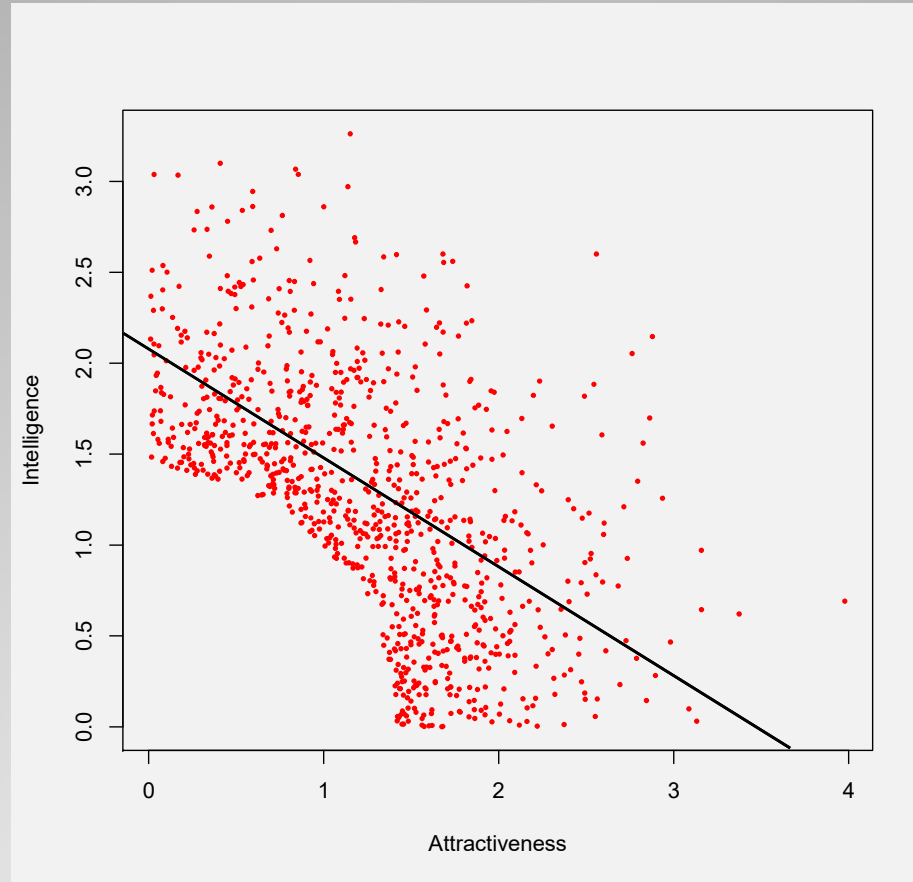
Why being picky can be good and bad



**Bivariate plot for actors in  
Hollywood**



**Bivariate plot for actors who  
"made it" in Hollywood**



$P < 2e-16$

**Bivariate plot for actors who  
"made it" in Hollywood**

- Again – you're meant to say something
- I'm waiting...

**What happened?**

- Bias in your parameter estimates
  - Bias is a difference between the “true value” and the estimated value
- Can apply across a range of scenarios
  - Bias estimates of means, variances, covariances, betas etc.

**Ascertainment**

- For testing means, ascertainment increases power
- For characterizing variance:covariance structure, ascertainment can lead to bias

**When might we want to ascertain?**

- For testing means, ascertainment increases power
- For characterizing variance:covariance structure, ascertainment can lead to bias

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## Mapping Mendelian Factors Underlying Quantitative Traits Using RFLP Linkage Maps

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**When might we want to ascertain?**

- Power calculations using NCP
- We create the model
  - specifying our effect sizes
- We then simulate data
  - empirical = T means that the simulated data matches the specifications [within some error]
- The chi square can then be used to generate power



**Practical!**