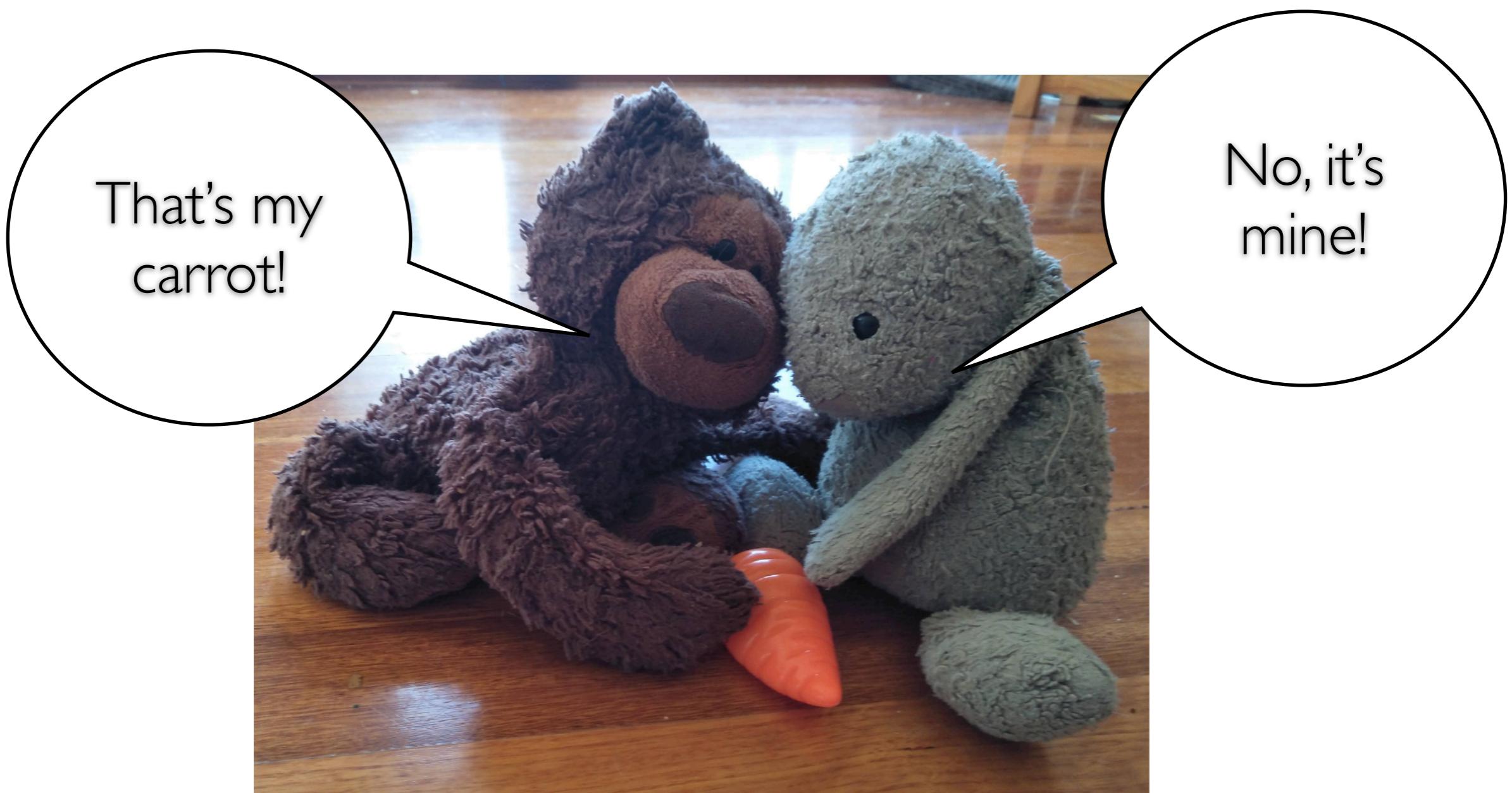


# **T-tests: One sample theory**

Research Methods for Human Inquiry  
Andrew Perfors

# Today's story...

Everyone in Bunnyland is going hungry, and they are starting to fight about it



# Today's story...

Most people think they need to take some sort of aggressive action against the Others



# Today's story...

But some advise caution - there's no evidence the  
Others are responsible

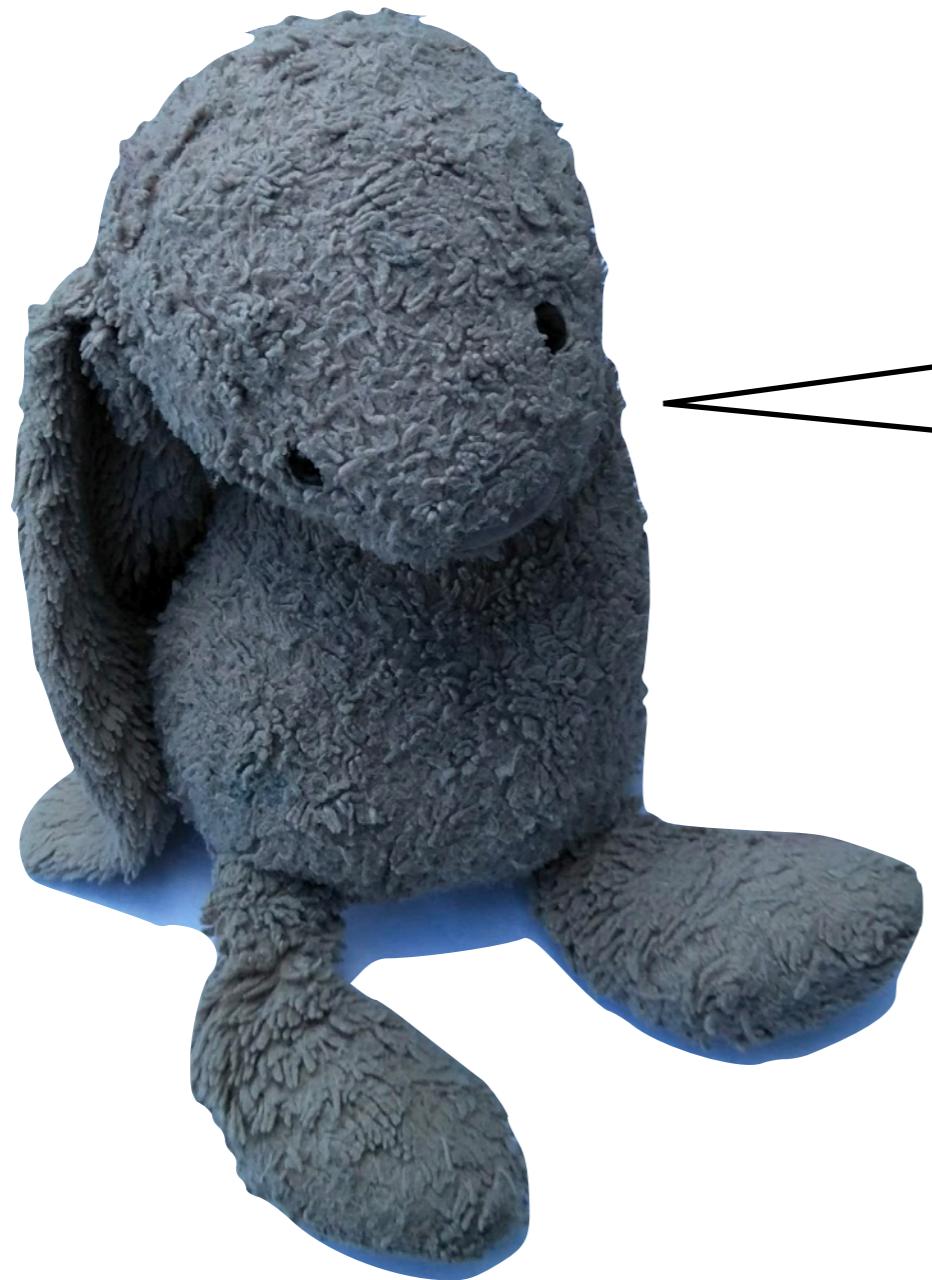


# Today's story...

You just don't  
want to attack the  
Others because  
you're scared



# Today's story...



I'm scared all right,  
but I'm more scared of  
us running out of food or  
fighting with each other  
when that's not even  
the real problem

# Today's story...



And you are the one  
that's acting irrationally:  
you're blinded by anger

# Today's story...

I'm not angry at all! I just don't want to abandon LFB and Foxy.



# Today's story...

Hey... I have an idea.  
Let's each take the CR test  
to make sure we're not  
acting emotionally



# Shadow's CR test



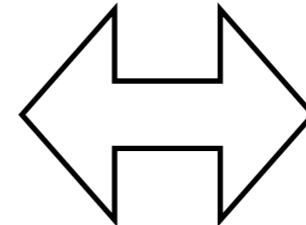
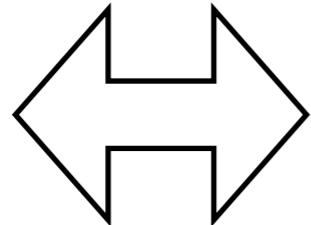
Measures Calmness  
and Rationality. If  
everyone takes it, we  
can figure out who is  
acting irrationally and  
being emotion driven!

# But how do you compare things like this?

Bunny's CR score: 70



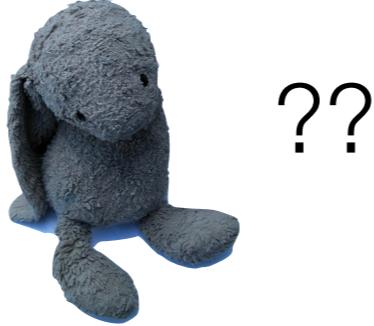
??



Sometimes you want to know how single scores  
compare in a somewhat objective way

# But how do you compare things like this?

Bunny's CR score: 70

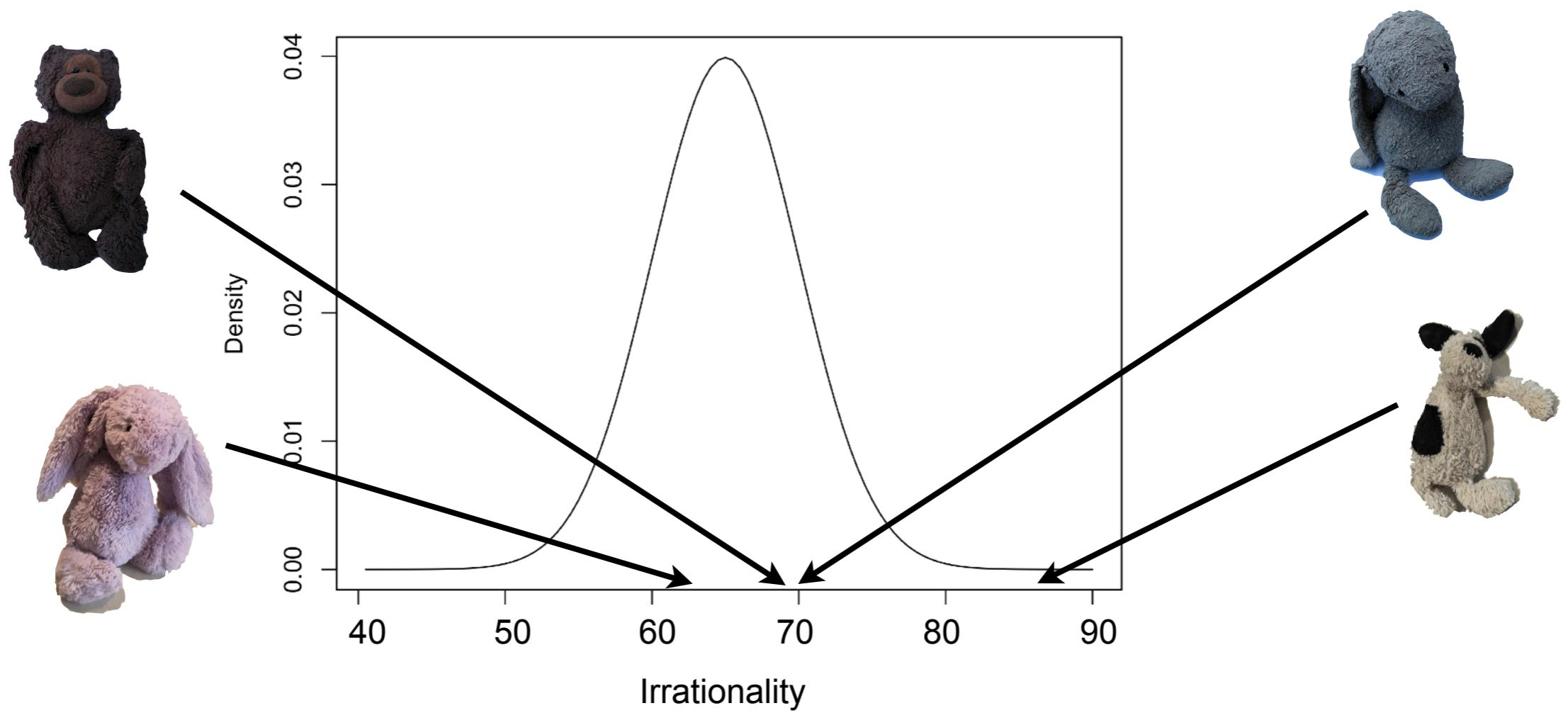


- Even knowing the scale doesn't help much...
  - Bunny answered 70 of 100 questions in an "irrational" way
  - Does that answer the question satisfactorily?
- For her CR score to make sense, we need to know what everyone else's CR scores were!

Sometimes you want to know how single scores compare in a somewhat objective way

# This helps a lot

- Bunny's score: 70
- Population mean:  $\mu = 64$
- Population std dev:  $\sigma = 12$



# This helps a lot

- Bunny's score: 70
- Population mean:  $\mu = 64$
- Population std dev:  $\sigma = 12$
- Much more informative, right?
  - Bunny is 0.5 standard deviations above the mean CR score in the population
  - Or, she has a standardised CR score of  $z = 0.5$

# Standard scores

$$z = \frac{X - \mu}{\sigma}$$

my standard score      my raw score      the population mean

the population standard deviation

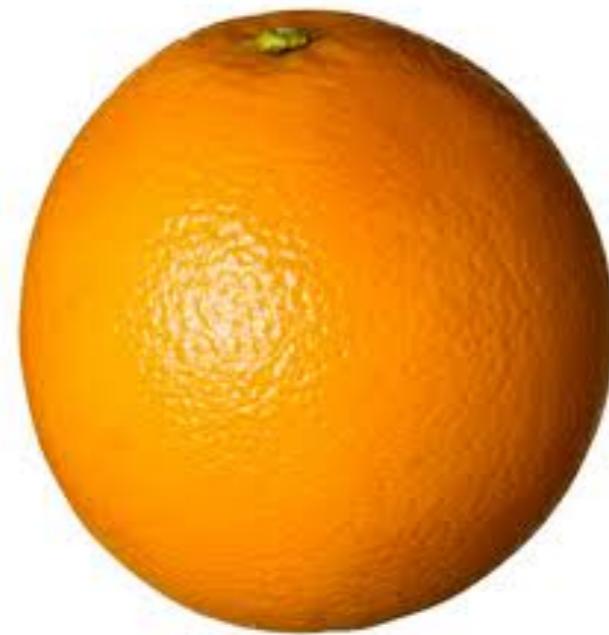
The diagram illustrates the components of a z-score formula. At the center is the formula  $z = \frac{X - \mu}{\sigma}$ . Above the formula, three labels point to its parts: 'my standard score' points to the top term  $X - \mu$ , 'my raw score' points to the variable  $X$ , and 'the population mean' points to the symbol  $\mu$ . Below the formula, an upward-pointing arrow indicates the position of 'the population standard deviation', which corresponds to the denominator  $\sigma$ .

*All z-scores have mean 0 and standard deviation 1, by definition*

# We can use this to compare two things too...

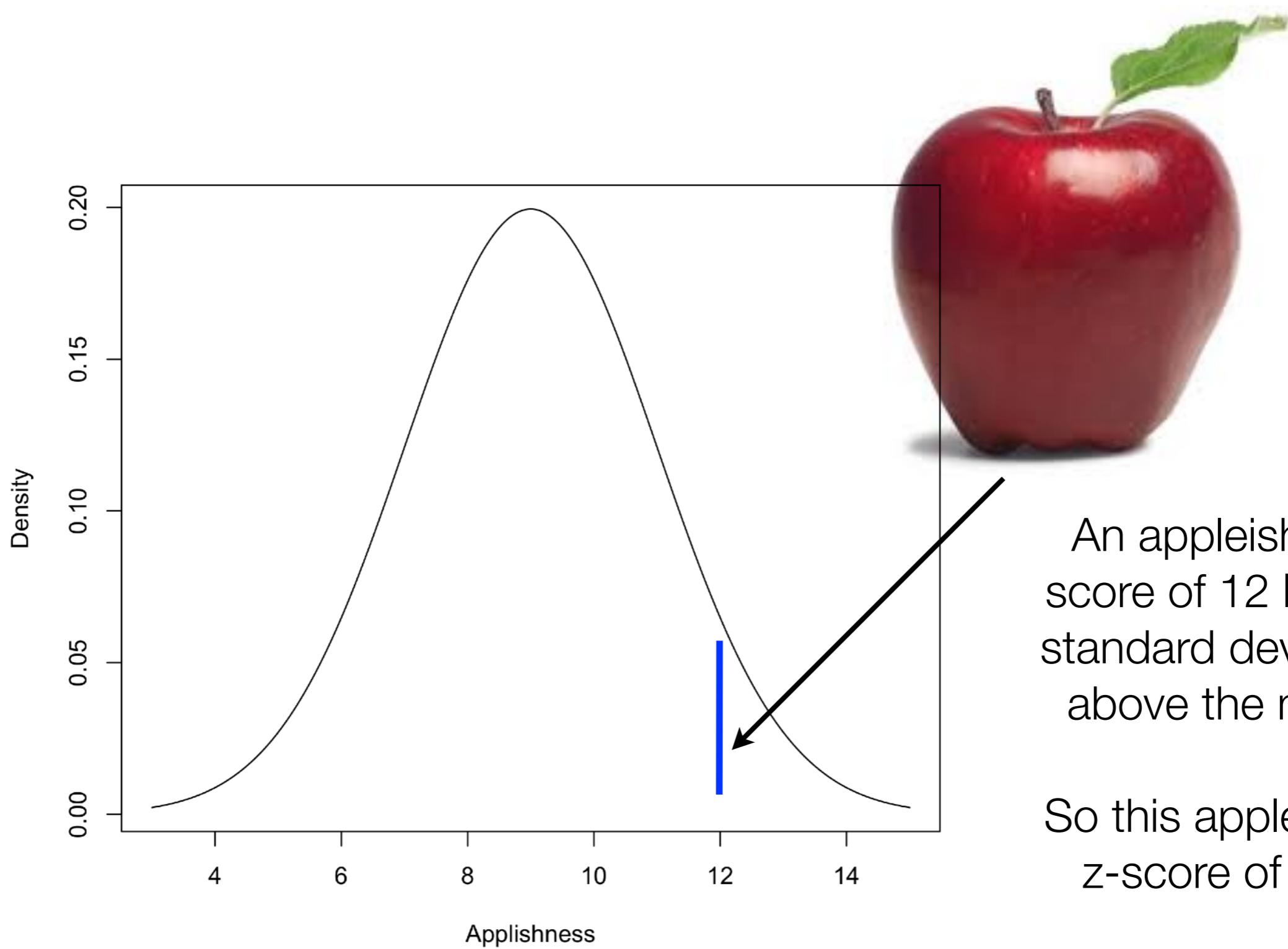


this apple has an  
apple-ness of 12

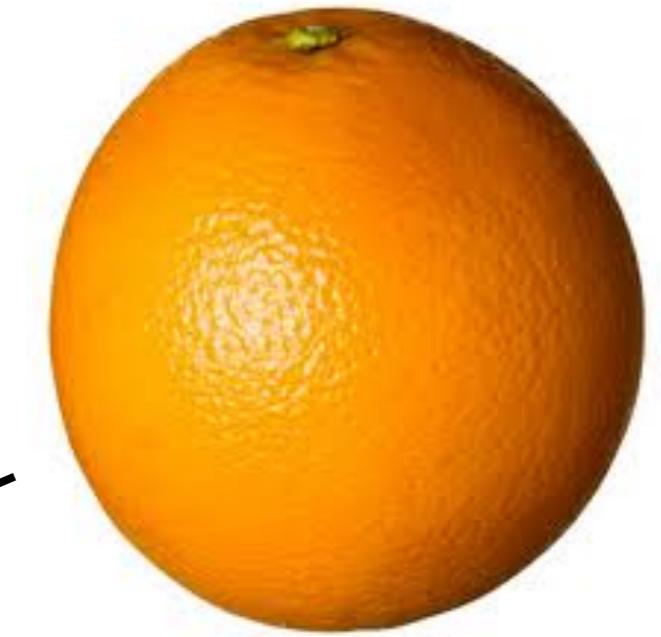
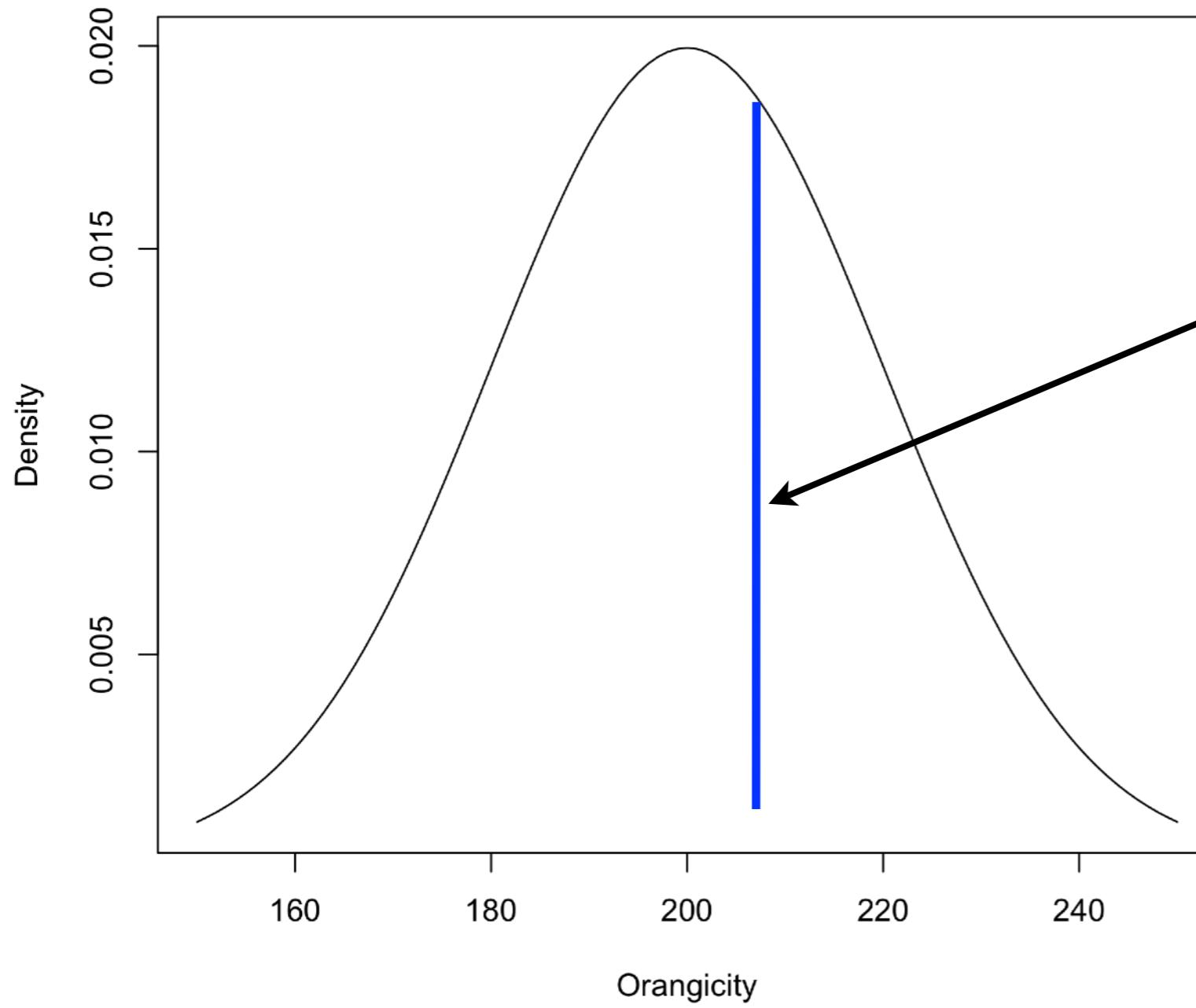


this orange has an  
orange-ness of 205

So... we begin by comparing our apple  
to other apples...



# And then we compare our orange to other oranges

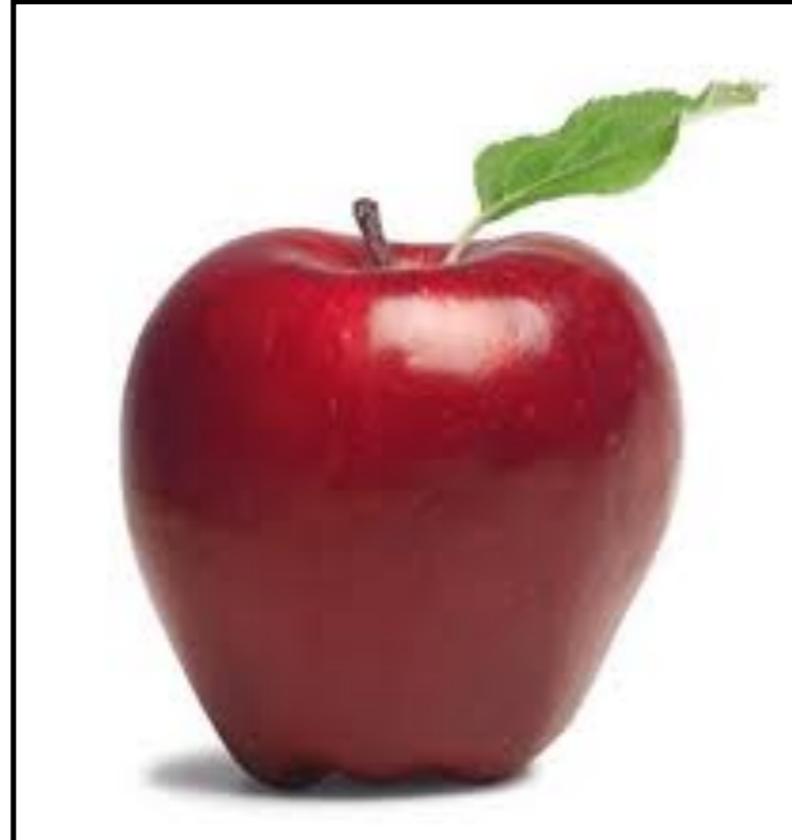


An orangicity of 205 is only 0.25 standard deviations above the mean

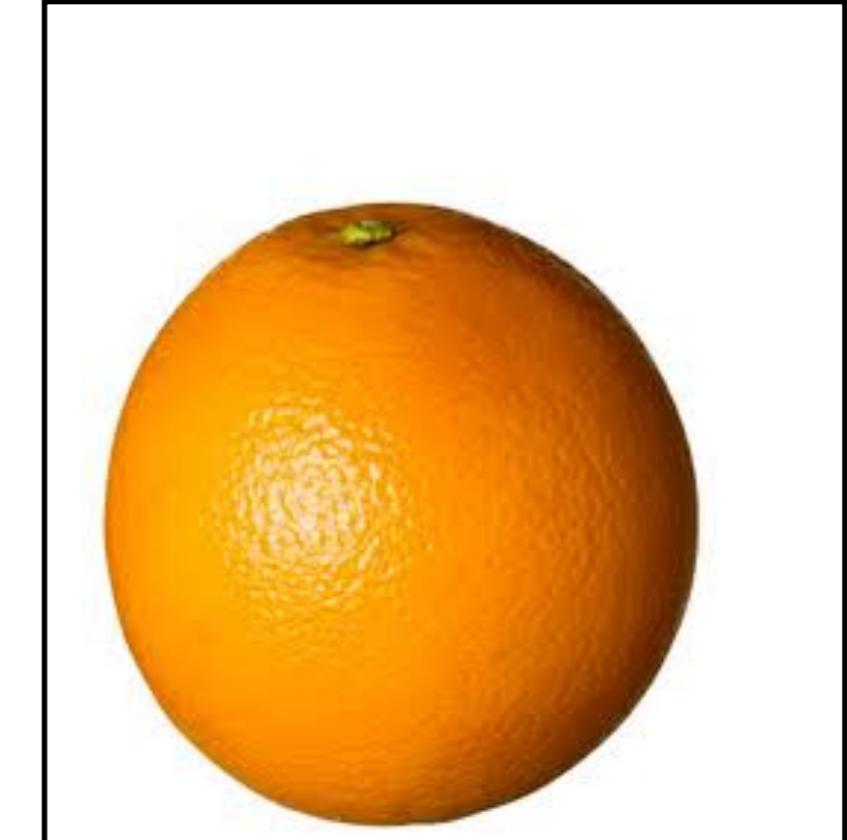
So this orange has a standard score of  
 $z = +0.25$

# Comparing apples to oranges using standard scores actually makes sense!

The apple is more applish than the orange is orangish



this apple has a z-score of +2.5



this orange has an z-score of +0.25

# This is great in psychology...

- Psychology is filled with situations when we want to make apples-to-oranges comparisons
- Examples:
  - I'm less introverted ( $z = -1.0$ ) than nerdy ( $z = +2.5$ )
  - Voldemort is more evil ( $z = +8$ ) than ugly ( $z = +4.5$ )



# (This is also what adjusted residuals reveal)

- In Week 6 Day 2 we talked about adjusted residuals for chi-squared tests — they are conceptually equivalent to z-scores (different calculation though since based on E and O, not  $\mu$ )

```
> ctResult$stdres
```

	box1	box2
doggie	-0.6654653	0.6654653
gladly	0.4077840	-0.4077840
shadow	0.4968698	-0.4968698

Returns the  
adjusted  
residuals

These are the raw residuals divided by the standard error  $\sqrt{E(1 - O_i/N)(1 - O_j/N)}$

Essentially what this does is allow us to interpret these as *kind of* indicating how many “standard deviations” away individual cells are.

A (weak) rule of thumb some use is that if you have an overall significant test and any adjusted residuals are more extreme than  $+/- 1.96$  or so, those individual items are significant as well

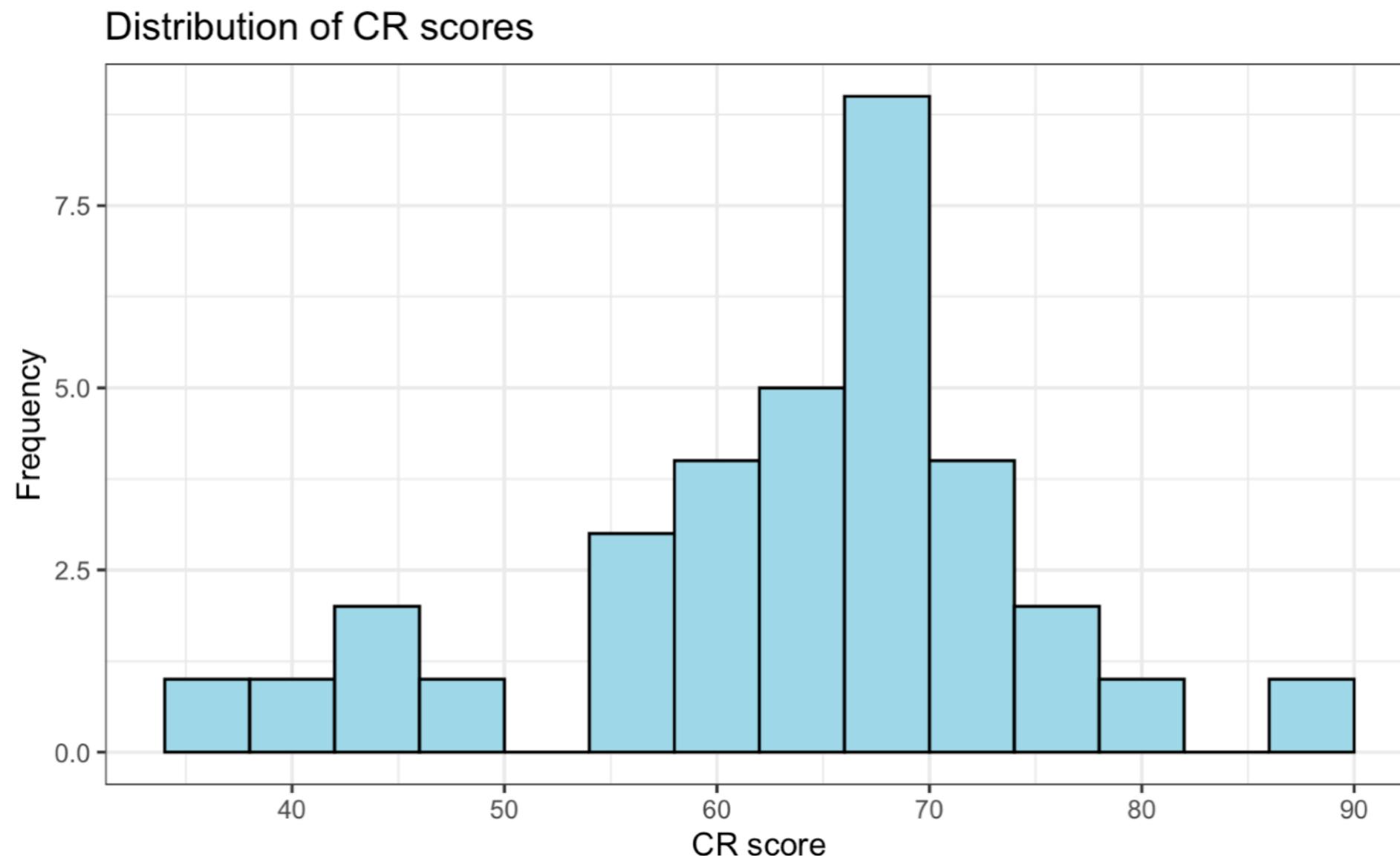
Comparing your data against a  
theoretically predicted mean using a  
one sample t-test

# Some data

```
> dcr <- read_csv(file=here("crscores.csv"))
> dcr
# A tibble: 34 x 2
  name          cr
  <chr>        <dbl>
1 bunny         70
2 gladly        69
3 flopsy        63
4 doggie        87
5 nosey         67
6 cuddly paws   56
7 shadow         56
8 pink bunny     59
9 purple bunny    68
10 blue bunny      55
# ... with 24 more rows
```

# Always have a look at it first!!

The CR test is designed to have a mean of 50 and sd of 10...  
Does this describe Bunnyland, or are they more generally irrational  
as a population right now?



# Are these two means the same???

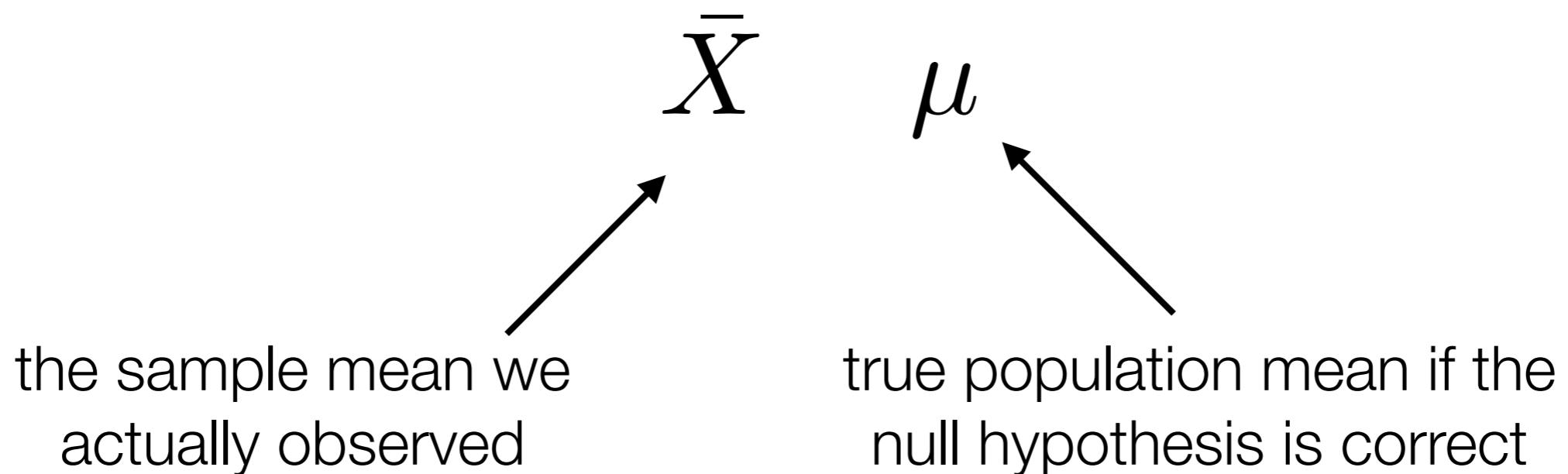
	sample	theoretical population
our mean	63.6	50
our standard deviation	11.6	10

# Let's construct a test....

- 1) A diagnostic test statistic,  $T$
- 2) Sampling distribution of  $T$  if the null is true
- 3) The observed  $T$  in your data
- 4) A rule that maps every value of  $T$  onto a decision (accept or reject  $H_0$ )

# A diagnostic test statistic

the two obvious quantities  
of interest...



# A diagnostic test statistic

the difference between them should be pretty close to zero if the null hypothesis is actually true

$$\frac{\bar{X} - \mu}{\sigma}$$

ideally we could just convert this into a standard z score, by dividing by the true population standard deviation if the null hypothesis is correct...

but we don't know what that is!

# A diagnostic test statistic

$$t = \frac{\bar{X} - \mu}{\hat{\sigma} / \sqrt{N}}$$

this is called  $t$ , and it is  
our test statistic

instead we convert  
it to something *like*  
a standard score,  
by dividing it by the  
SEM (which we  
can calculate)

this has mean of about  
zero and standard  
deviation of about one

# Let's construct a test....

t ✓ 1) A diagnostic test statistic,  $T$

6.86 ✓ 2) Sampling distribution of  $T$  if the null is true

3) The observed  $T$  in your data

4) A rule that maps every value of  $T$  onto a decision (accept or reject  $H_0$ )

$$t = \frac{\bar{X} - \mu}{\hat{\sigma}/\sqrt{N}}$$
$$= \frac{63.6 - 50}{11.6/\sqrt{34}} = 6.86$$

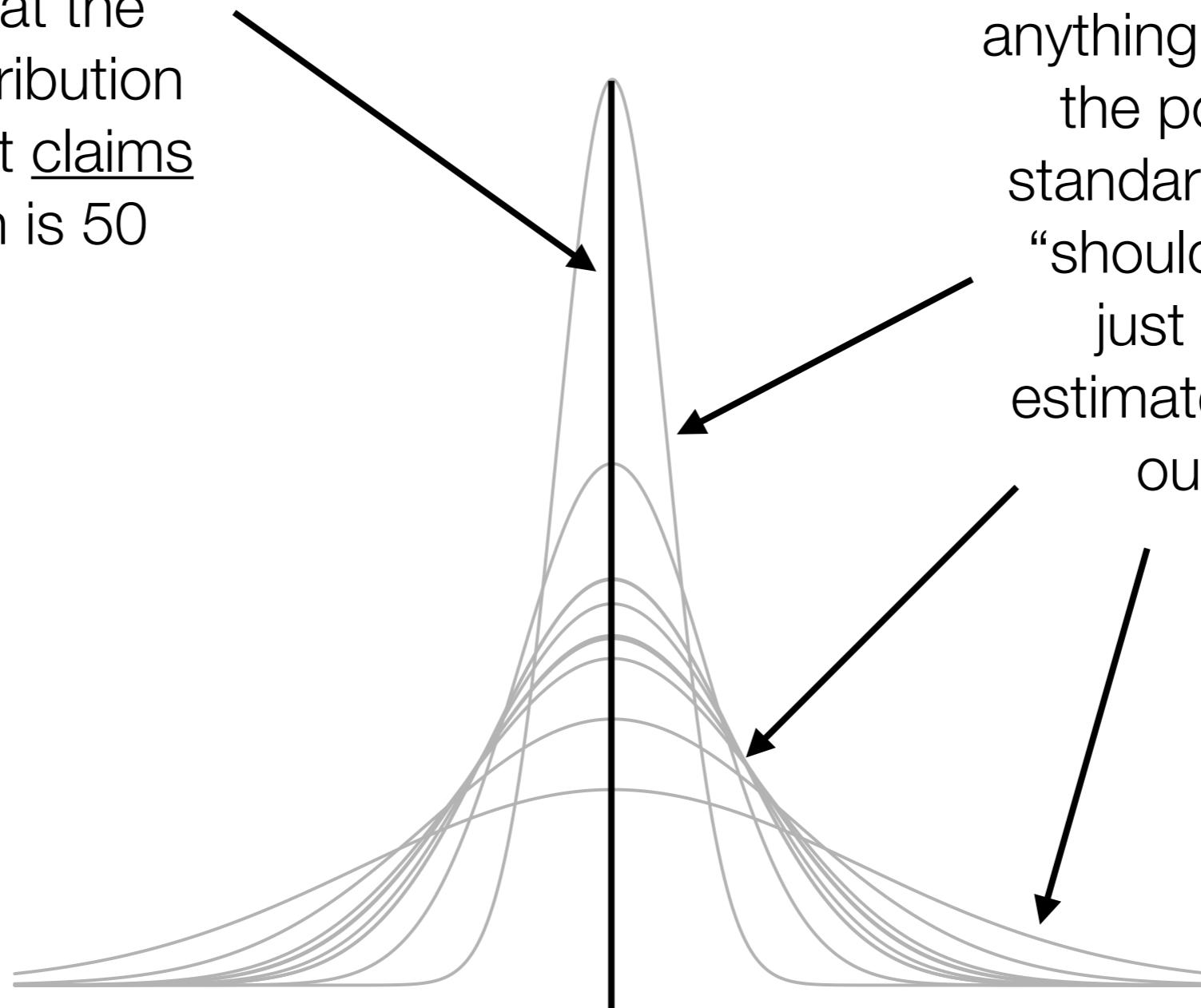
`(mean(dcr$cr) - 50) / (sd(dcr$cr)/sqrt(nrow(dcr)))`

# Sampling distribution of t

What does our null hypothesis actually say?

It assumes that the population distribution is normal, and it claims that the mean is 50

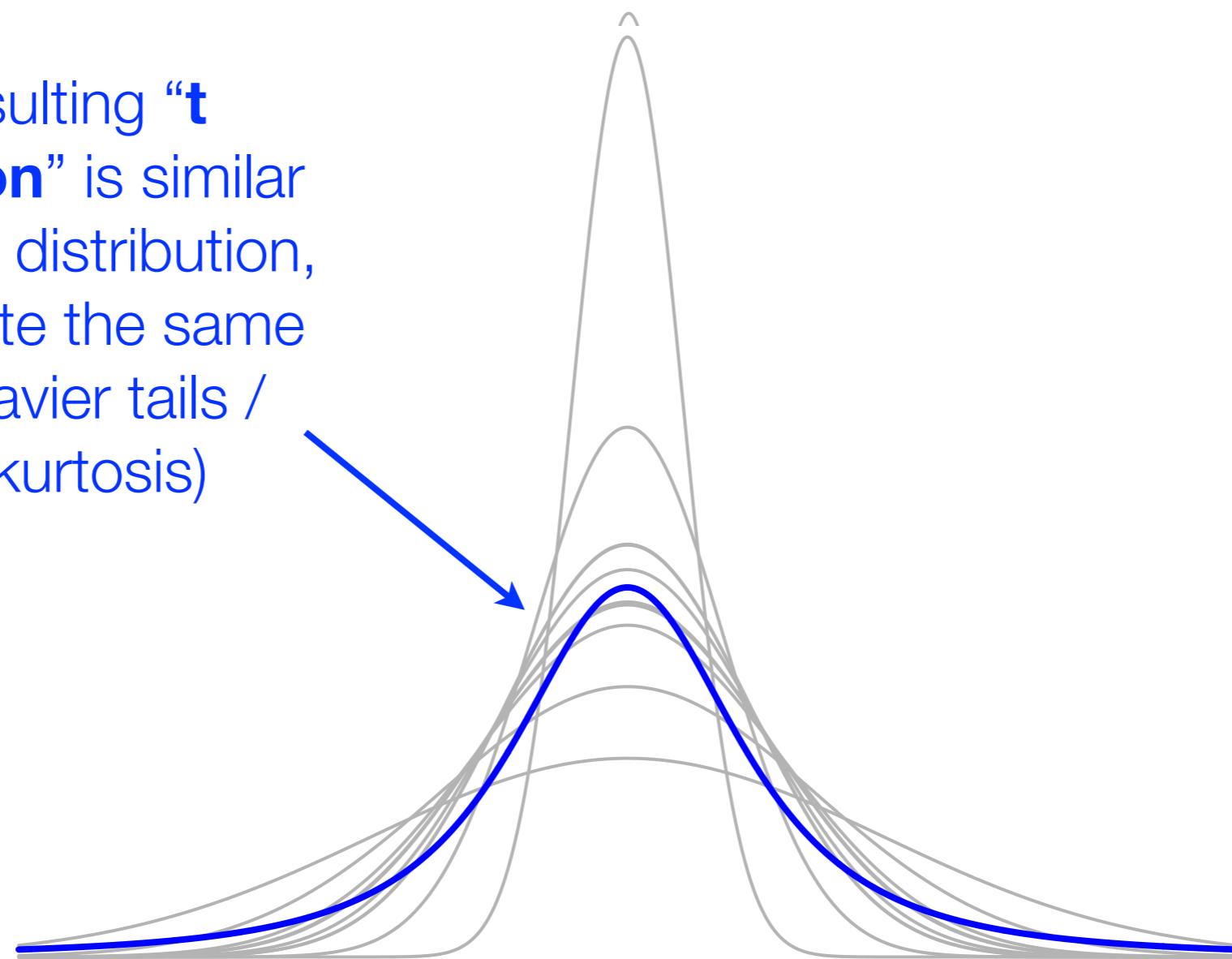
It doesn't say anything about what the population standard deviation "should" be. We just have to estimate that from our data



# Sampling distribution of t

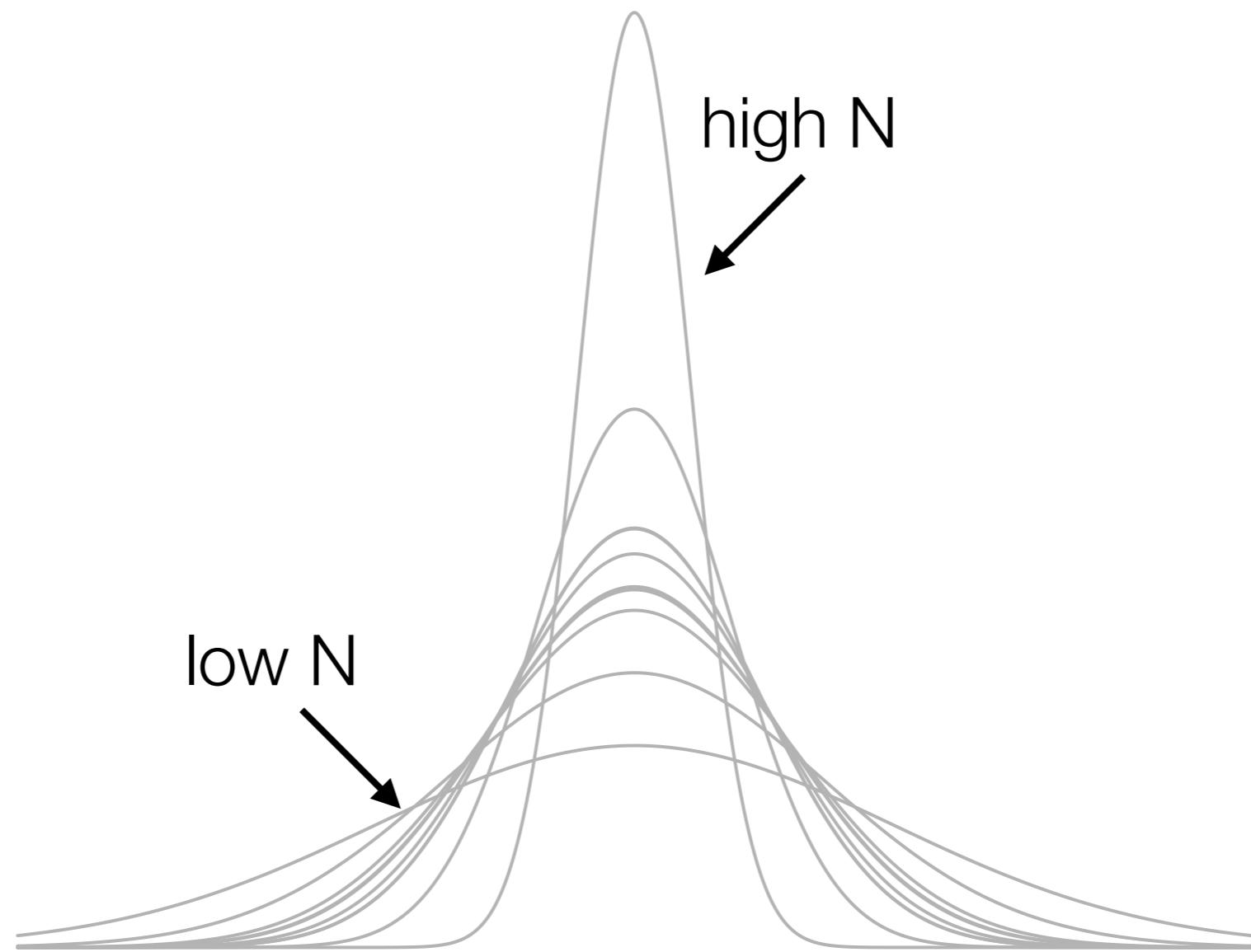
Since we're unsure about what the true standard deviation is, the overall sampling distribution for the t statistic is obtained by *averaging over lots of possible choices* for the population standard deviation

The resulting “**t distribution**” is similar to a normal distribution, but not quite the same  
(it has heavier tails / higher kurtosis)



# Sampling distribution of t

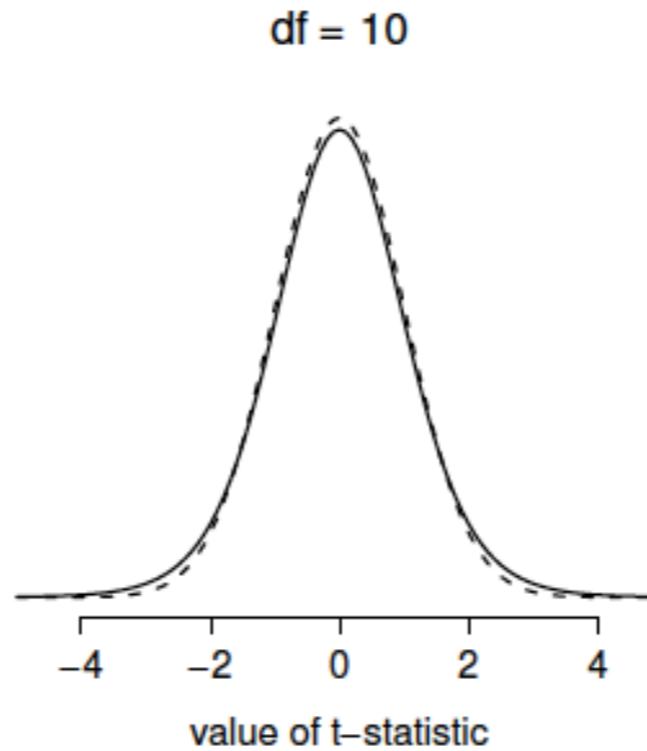
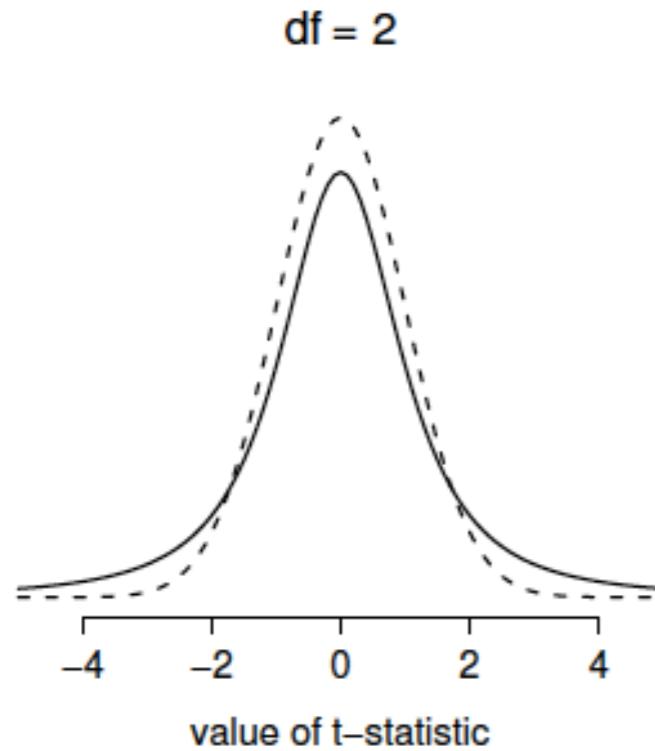
However, as sample size (N) gets larger, our estimate of the standard deviation is more precise, so the t-distribution grows more similar to normal



# Sampling distribution of t

However, as sample size (N) gets larger, our estimate of the standard deviation is more precise, so the t-distribution grows more similar to normal

This is reflected in the *degrees of freedom*, which for a t distribution is  $N-1$ .  
( $N = \#$  of data points; 1 = # of constraints, i.e., the mean)



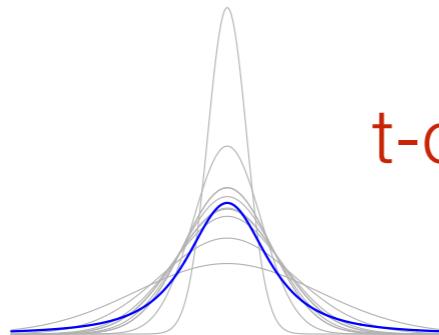
higher degrees of freedom / larger sample means the t-distribution looks more normal

# Let's construct a test....

- ✓ 1) A diagnostic test statistic,  $T$

$$t = \frac{\bar{X} - \mu}{\hat{\sigma}/\sqrt{N}}$$

- ✓ 2) Sampling distribution of  $T$  if the null is true



t-distribution with  $N-1$  degrees of freedom

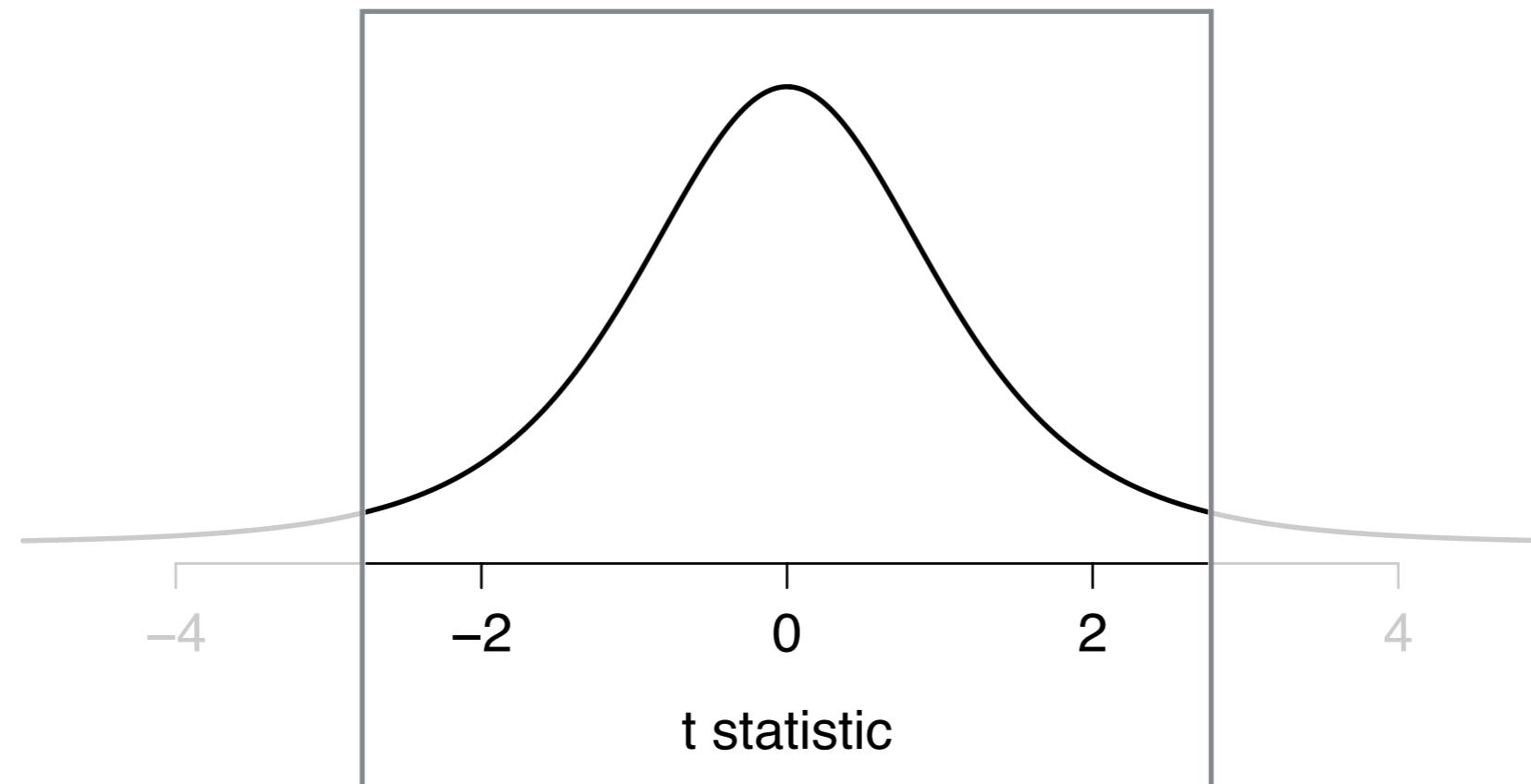
- ✓ 3) The observed  $T$  in your data

6.86

- 4) A rule that maps every value of  $T$  onto a decision (accept or reject  $H_0$ )

# A rule for mapping values of t onto a decision

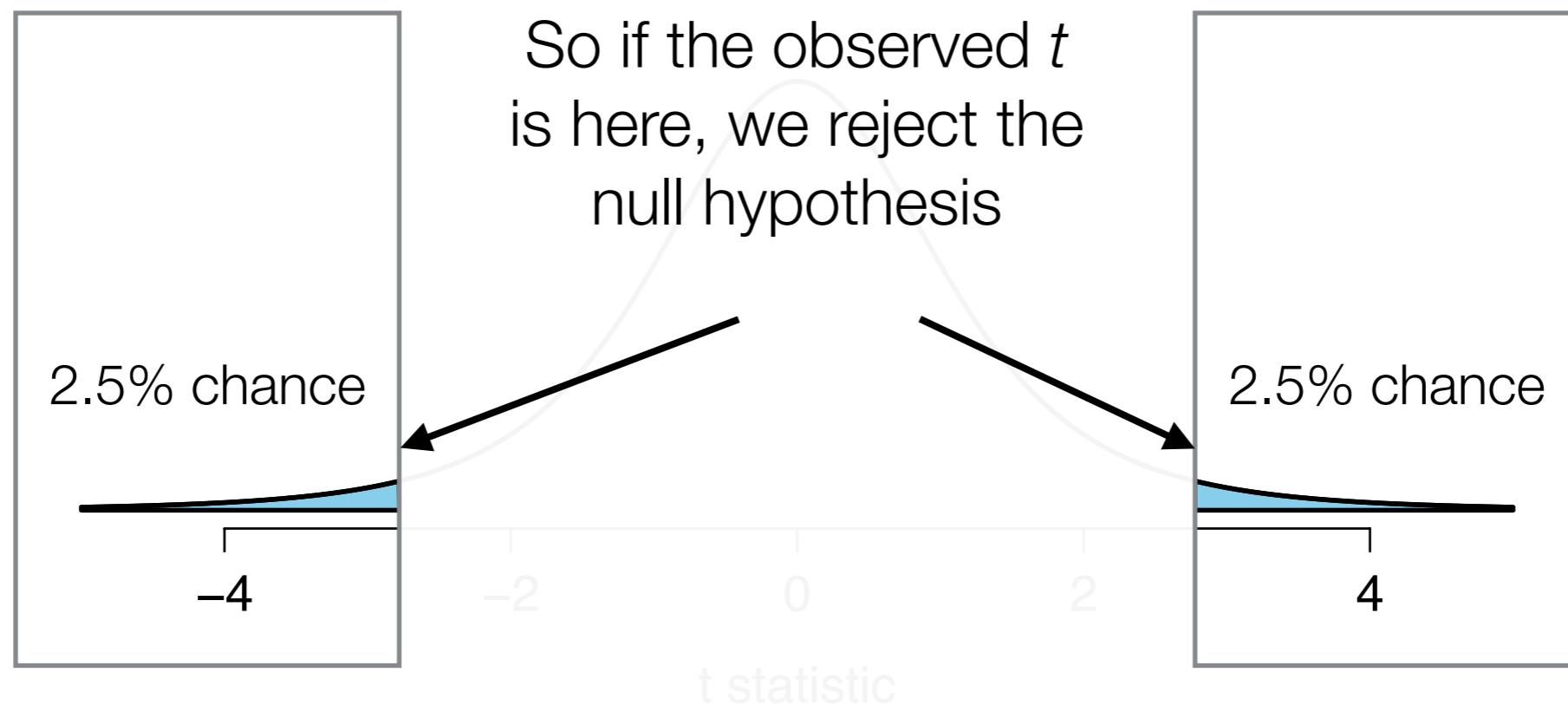
If the null hypothesis is true, there's a 95% chance that the t-statistic falls in this range



If the null hypothesis is true, we expect the t-statistic to have come from this sampling distribution...

# A rule for mapping values of $t$ onto a decision

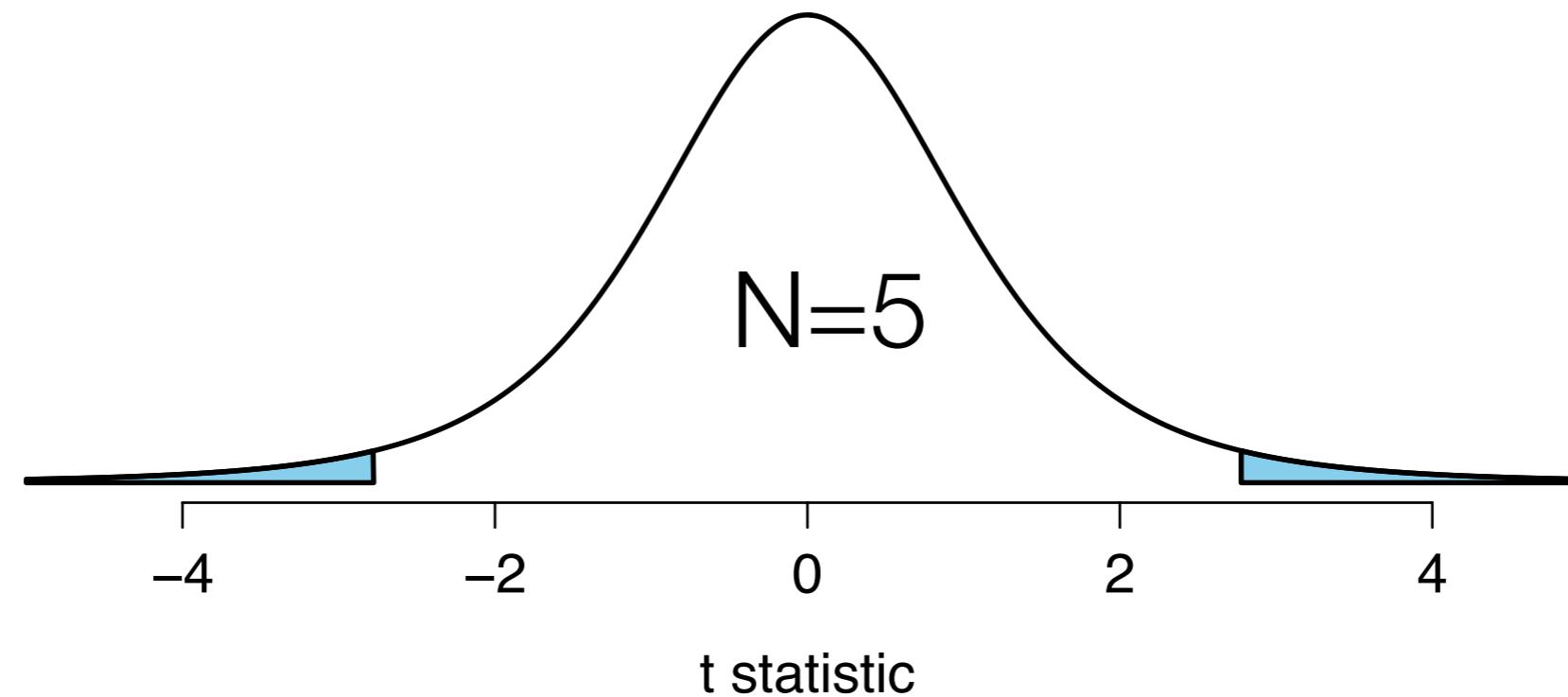
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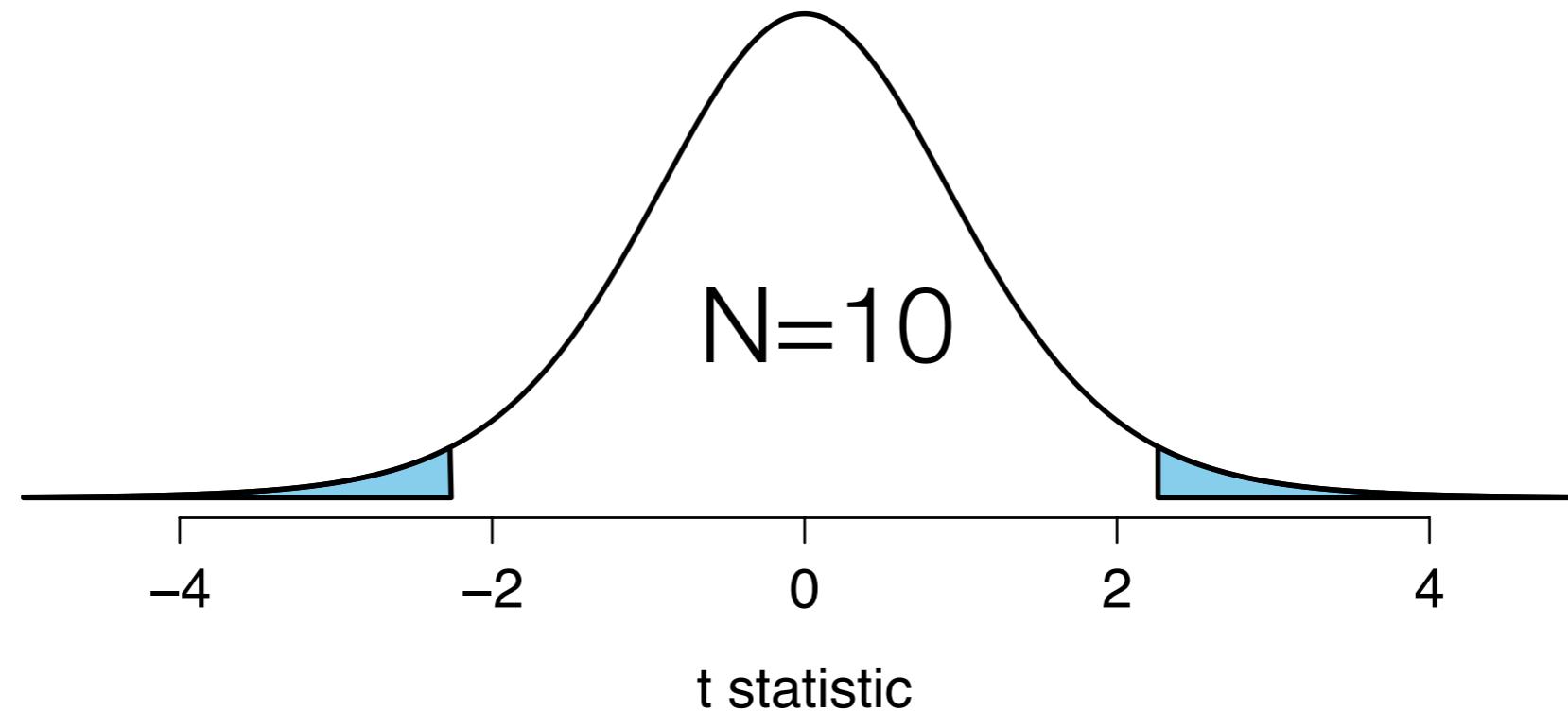
# A rule for mapping values of t onto a decision

Because the width of the t distribution depends on sample size, the size of the rejection region changes as N increases...



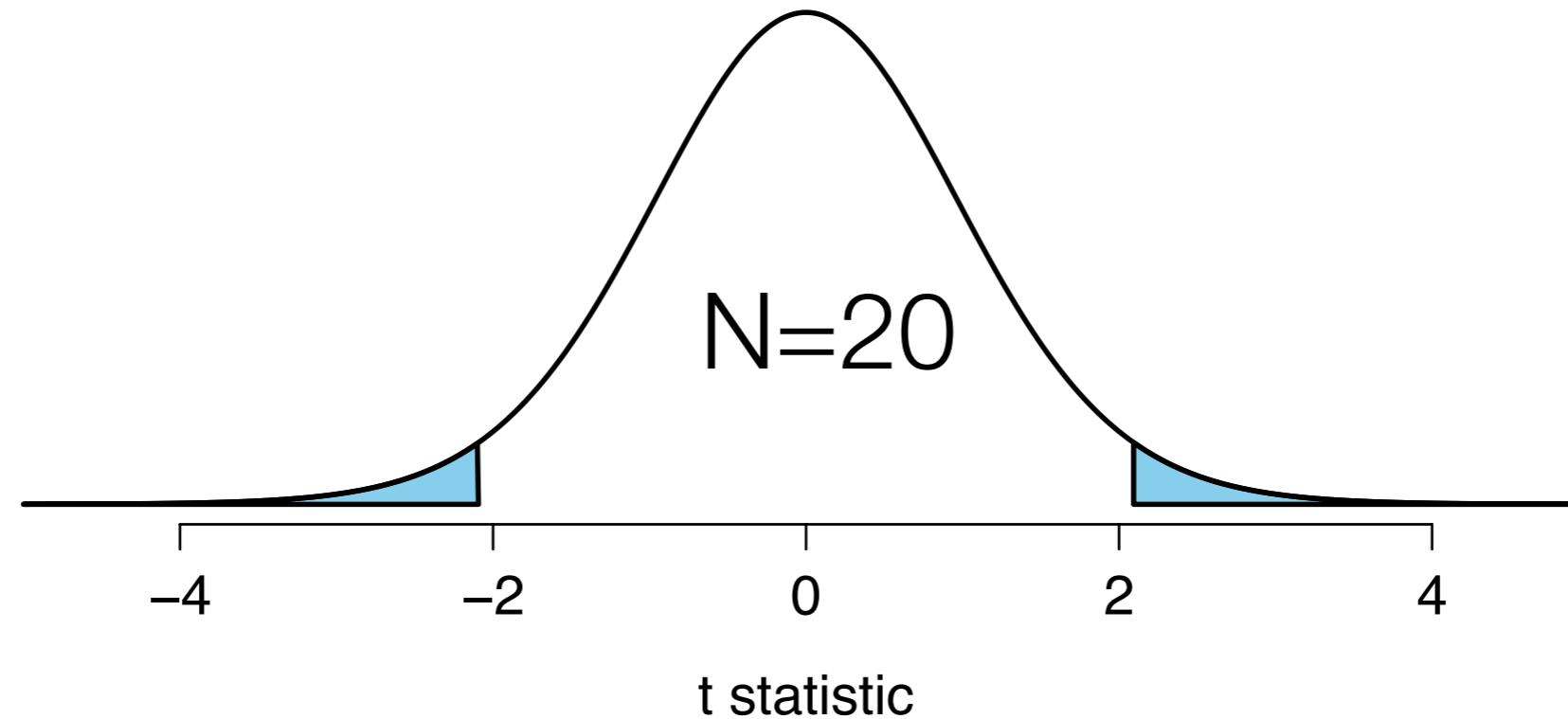
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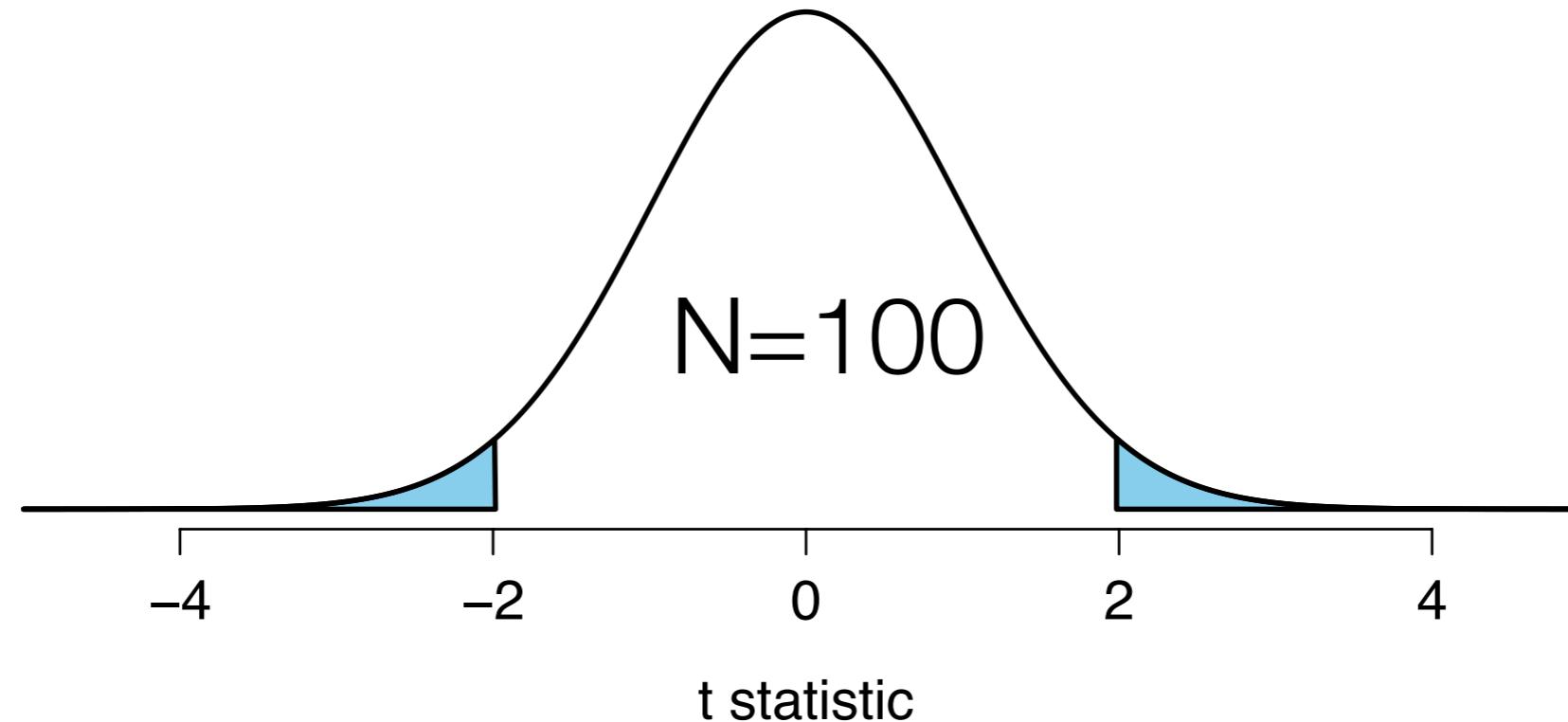
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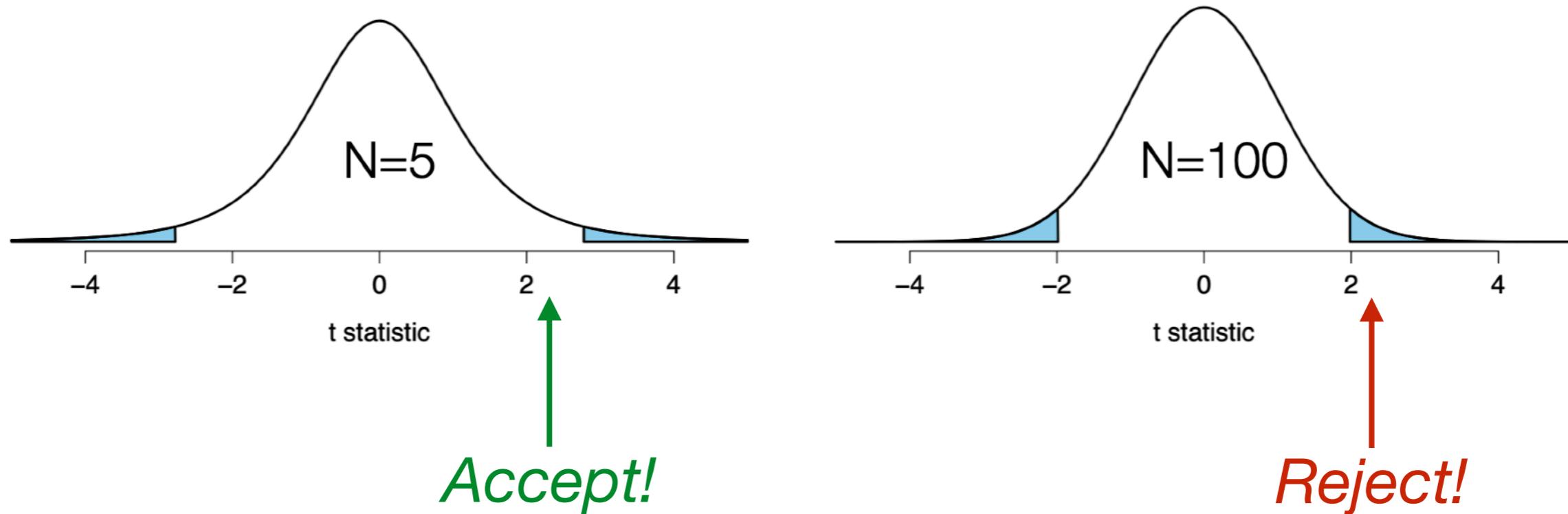
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# A rule for mapping values of t onto a decision

Because the width of the t distribution depends on sample size, the size of the rejection region changes as N increases...



This means that whether a given t-statistic is significant depends on sample size!

Exercises are in w7dayexercises.Rmd