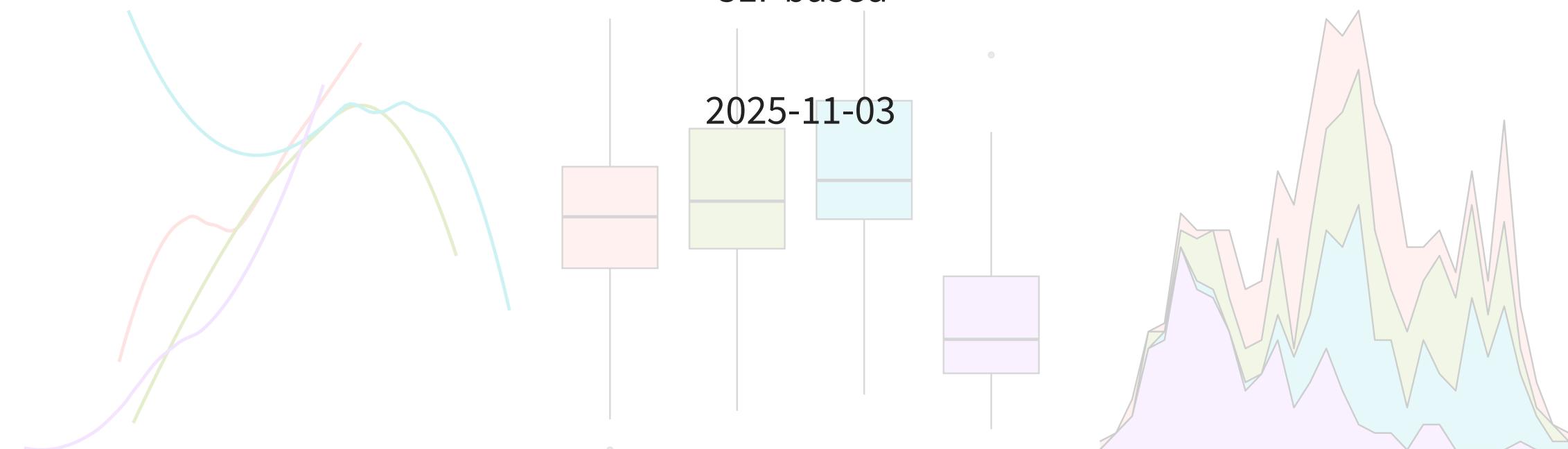
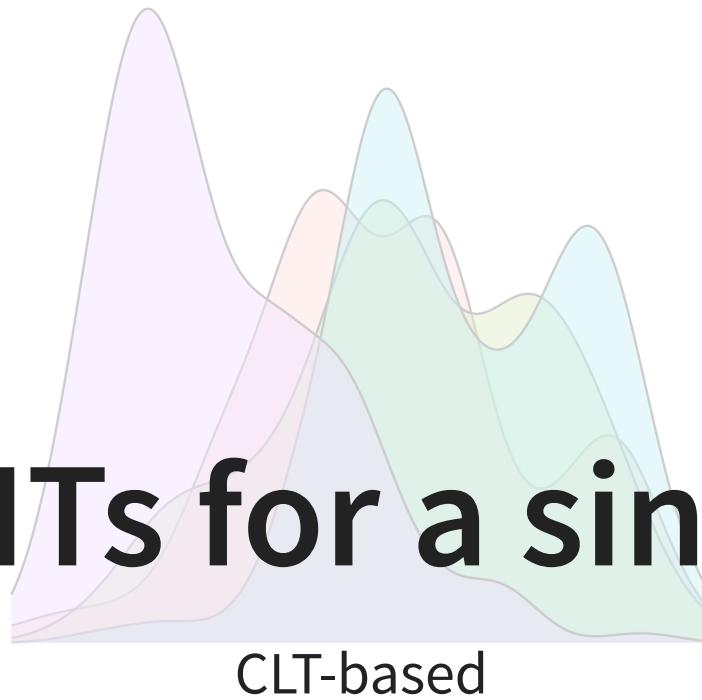


# CLIs and HTs for a single mean



# Housekeeping

- Dessert social today! 3:30-4:30pm in WNS 105!
- Modified office hours today: 1:00-2:00pm
- Homework 7 due tonight
- Project proposals due Wednesday night at 11:59pm

# Recap

- Back to numerical data
- CLT: if we have a “sufficiently large” sample of  $n$  independent observations from a population with mean  $\mu$  and standard deviation  $\sigma$ , then  $\bar{X} \sim N\left(\mu, \frac{\sigma}{\sqrt{n}}\right)$

- To obtain a  $\gamma \times 100\%$  CI via CLT, we use

point estimate  $\pm$  critical value  $\times$  SE

- We *may* need to replace the standard error with an estimate  $\widehat{SE}$

# Checking normality

- Remember, CLT requires a sufficiently large sample size  $n$  or assumption of Normality of the underlying data.
- No perfect way to check Normality, but rule of thumb:
  - If  $n < 30$  small: check that there are no clear outliers
  - If  $n \geq 30$  large: check that there are no particularly extreme outliers

# CI for a single mean

If you'd like supplemental reading on CLT-based inference for a single mean: [Section 19.2](#) of the textbook can be helpful!

# CI for a single mean (known SD)

Suppose we want a  $\gamma \times 100\%$  CI for population mean  $\mu$ . Also suppose that  $\sigma$  is known to us!

- *If* CLT holds, then we know

$$\bar{X} \sim N\left(\mu, \frac{\sigma}{\sqrt{n}}\right)$$

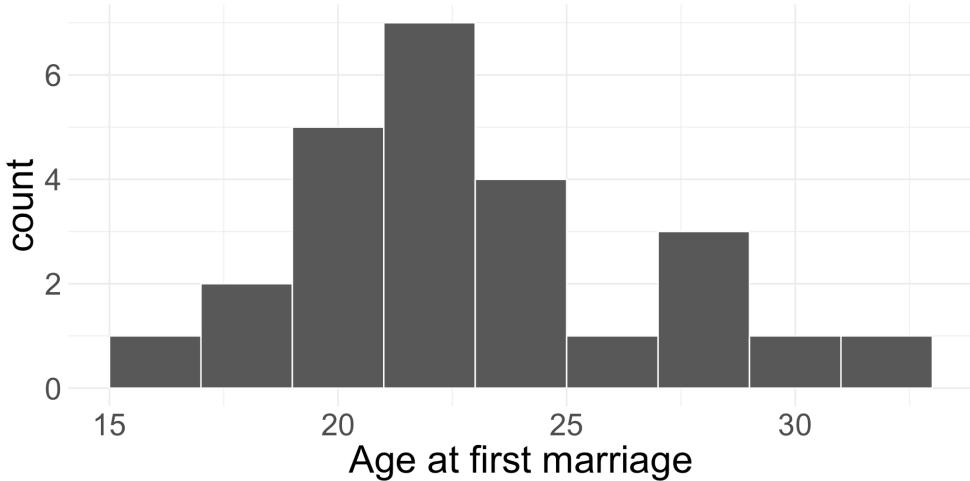
- So our  $\gamma \times 100\%$  CI for  $\mu$  is:

$$\text{point estimate} \pm \text{critical value} \times \text{SE} = \bar{x}_{obs} \pm z_{(1+\gamma)/2}^* \times \frac{\sigma}{\sqrt{n}}$$

- Here, because we assumed  $\sigma$  known, we don't need to (and shouldn't) approximate SE

# Example: age at marriage

In 2006-2010, the CDC conducted a thorough survey asking US women their age at first marriage. Suppose it is known that the standard deviation of the ages at first marriage is 5 years. Suppose we randomly sample 25 US women and ask them their age at first marriage (plotted below). Their average age at marriage was 23.32.



We will obtain an 80% confidence interval for the mean age of US women at first marriage.

- Are conditions of CLT met?
- If so, what does CLT tell us?

What is/are the population parameter(s)? What is the statistic?

# Example: age at marriage (cont.)

Obtain an 80% confidence interval for the mean age of US women at first marriage.

- Because we have a random sample (independence) and there are no outliers in the data (normality condition), we can proceed with CLT!

$$\bar{X} \sim N\left(\mu, \frac{5}{\sqrt{25}}\right) = N(\mu, 1)$$

- Construct your confidence interval and interpret!

1. Point estimate:  $\bar{x}_{obs} = 23.32$

2. Standard error:  $SE = 1$

3. Critical value:  $z_{0.9}^* = qnorm(0.9, 0, 1) = 1.28$

So our 80% confidence interval is  $23.32 \pm 1.28 \times 1 = (22.04, 24.6)$

# Utility of this model

- The previous formula for the confidence interval for  $\mu$  relies on knowing  $\sigma$
- But wait...
  - Want to construct a CI for  $\mu$  because we don't know its value
  - If we don't know  $\mu$ , it seems highly unlikely that we would know  $\sigma$ !
- So in practice, we will have to estimate standard error for  $\bar{X}$ :

$$\widehat{\text{SE}} = \frac{s}{\sqrt{n}}$$

where  $s$  is the observed sample standard deviation

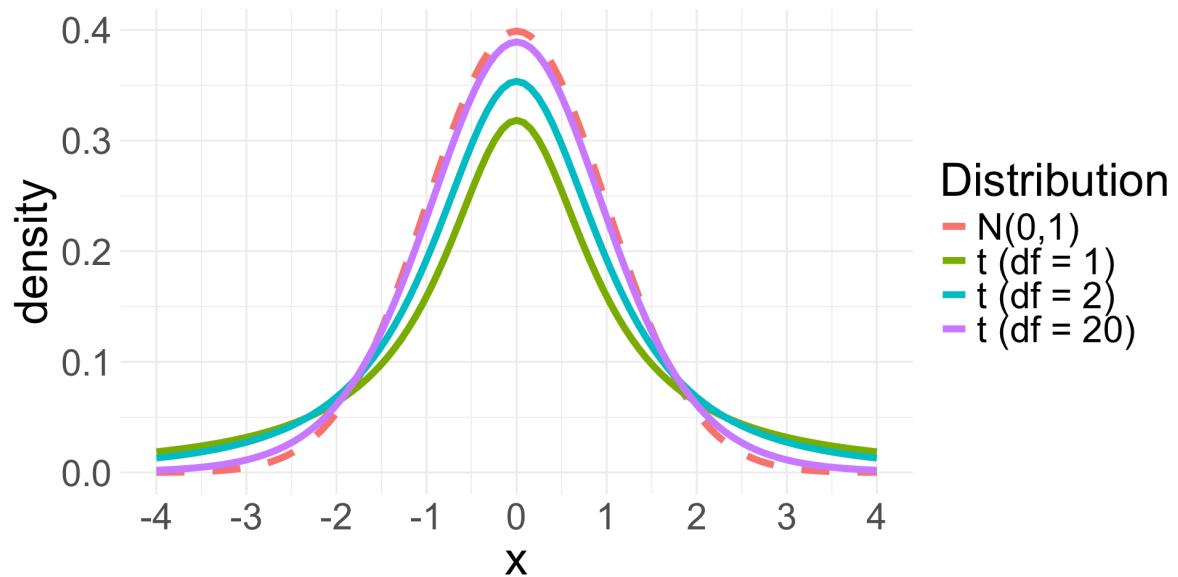
- Recall we did something similar for CI for  $p$ , where we replaced  $p$  with  $\hat{p}_{obs}$

# Variance issue

- Estimating variance is extremely difficult when  $n$  is small, and still not great for large  $n$ 
  - Replacing  $\sigma$  with  $s$  actually invalidates CLT
- **So if  $\sigma$  is unknown, we *cannot* use the Normal approximation to model  $\bar{X}$**  for inferential tasks
- Instead, we will use a different distribution for inference calculations, called the  $t$ -distribution

# $t$ -distribution

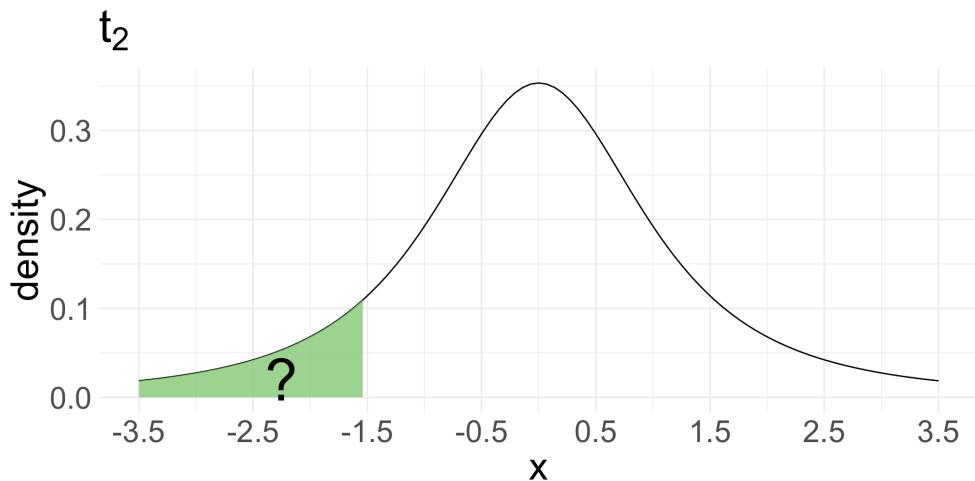
- The  $t$ -distribution is symmetric and bell-curved (like the Normal distribution)
- Has “thicker tails” than the Normal distribution (the tails decay more slowly)



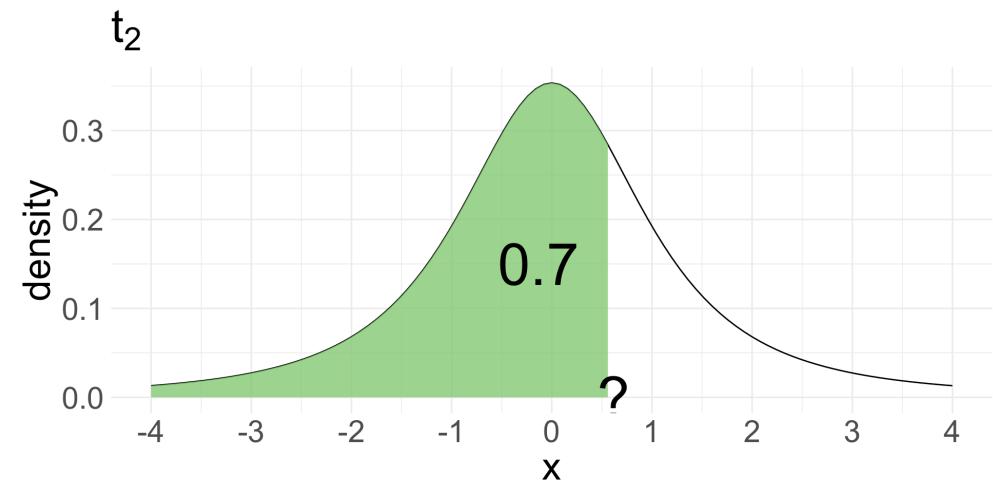
- $t$ -distribution is always centered at 0
- One parameter: **degrees of freedom (df)** defines exact shape of the  $t$ 
  - Denoted  $t_{df}$  (e.g.  $t_1$  or  $t_{20}$ )
- As  $df$  increases,  $t$  resembles the  $N(0, 1)$ . When  $df \geq 30$ , the  $t_{df}$  is nearly identical to  $N(0, 1)$

# *t* distribution in R

- `pnorm(x, mean, sd)` and `qnorm(%, mean, sd)` used to find probabilities and percentiles for the Normal distribution
- Analogous functions for *t*-distribution: `pt(x, df)` and `qt(%, df)`



$$\text{pt}(-1.5, \text{df } = 2) = 0.1361966$$



$$\text{qt}(0.7, \text{df } = 2) = 0.6172134$$

# CI for a single mean (unknown variance)

- Still require independent observations and the Normality condition for CLT
- General formula for  $\gamma \times 100\%$  CI is the same, but we simply change what goes into the margin of error.

$$\text{point estimate} \pm t_{df,(1+\gamma)/2}^* \times \widehat{\text{SE}} = \bar{x}_{obs} \pm t_{df,(1+\gamma)/2}^* \times \frac{s}{\sqrt{n}}$$

- $df = n - 1$  (always for this CI)
- critical value  $t_{df,(1+\gamma)/2}^* = (1 + \gamma)/2$  percentile of the  $t_{df}$  distribution

# Example: age at marriage (cont.)

Let's return to the age at marriage example. Once again, obtain an 80% CI for the average age of first marriage for US women, but now suppose we **don't know**  $\sigma$ .

In our sample of  $n = 25$  women, we observed a sample mean of 23.32 years and a sample standard deviation of  $s = 4.03$  years.

1. Point estimate:  $\bar{x}_{obs} = 23.32$

2. Standard error:  $\widehat{SE} = \frac{s}{\sqrt{n}} = \frac{4.03}{\sqrt{25}} = 0.806$

3. Critical value:

- $df = n - 1 = 24$
- $t_{24,0.9}^* = \text{qt}(0.9, df = 24) = 1.32$

So our 80% confidence interval for  $\mu$  is:

$$23.32 \pm 1.32 \times 0.806 = (22.26, 24.38)$$

# Remarks

- Interpretation of CI does not change even if we use a different model!
- If you have access to both  $\sigma$  and  $s$ , which should you use?
  - You should use  $\sigma$ !

# Test for a single mean

# Hypothesis test recap

1. Set hypotheses
2. Collect and summarise data, set  $\alpha$
3. Obtain null distribution and p-value
  - For CLT-based method, obtain *test statistic*
4. Decision and conclusion

# Hypotheses and null distribution

Want to conduct a hypothesis test for the mean  $\mu$  of a population.

- Hypotheses:  $H_0 : \mu = \mu_0$  versus  $H_A : \mu \neq \mu_0$  (or  $\mu > \mu_0$  or  $\mu < \mu_0$ )
- Verify conditions for CLT
  1. Independence
  2. Approximate normality or large sample size
- Then from population with mean  $\mu$  and standard deviation  $\sigma$ , we have
$$\bar{X} \sim N\left(\mu, \frac{\sigma}{\sqrt{n}}\right)$$
- **What does the (approximate) null distribution for  $\bar{X}$  look like?**

$$\bar{X} \sim N\left(\mu_0, \frac{\sigma}{\sqrt{n}}\right)$$

# z-test and *t*-test statistics

Our test statistic is always of the form:

$$\frac{\text{observed} - \text{null}}{\text{SE}}$$

or

$$\frac{\text{observed} - \text{null}}{\widehat{\text{SE}}}$$

- If  $\sigma$  known and CLT met, we perform a **z-test** where our test-statistic is:
- If  $\sigma$  unknown and CLT met, we perform a ***t*-test** by estimating  $\sigma$  with  $s$ . Our test statistic is:

$$z = \frac{\bar{x}_{obs} - \mu_0}{\frac{\sigma}{\sqrt{n}}} \sim N(0, 1)$$

and we obtain our p-value using `pnorm()`

- Everything else proceeds as usual!

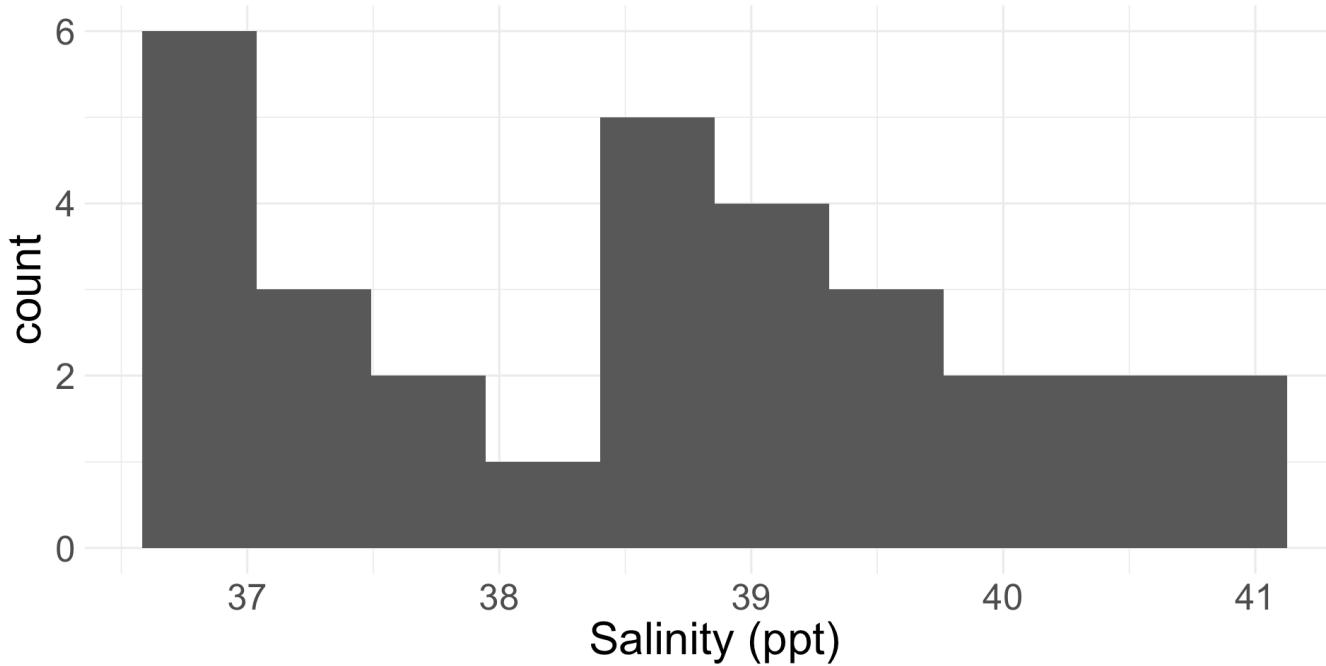
$$t = \frac{\bar{x}_{obs} - \mu_0}{\frac{s}{\sqrt{n}}} \sim t_{df} \quad df = n - 1$$

and we obtain our p-value using `pt()`

# Example: salinity

The salinity level in a body of water is important for ecosystem function.

We have 30 salinity level measurements (ppt) collected from a random sample of water masses in the Bimini Lagoon, Bahamas.



- We want to test if the average salinity level in Bimini Lagoon is different from 38 ppm at the  $\alpha = 0.05$  level.

# Example: salinity (cont.)

## 1. Set hypotheses (define parameters as necessary).

- $\bar{x}_{obs} = 38.6$

- Let  $\mu$  be the average salinity level in Bimini Lagoon in ppt.
- $H_0 : \mu = 38$  versus  $H_A : \mu \neq 38$
- $s = 1.29$
- $n = 30$

## 2. Collect summary information, set $\alpha$ .

- $\alpha = 0.05$

# Example: salinity (cont.)

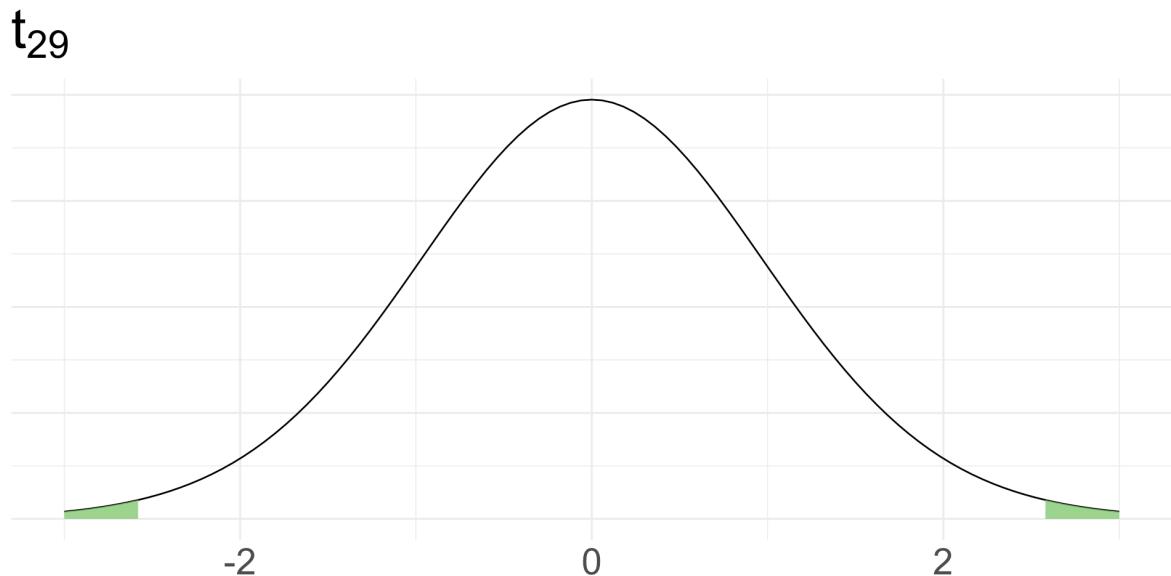
3. Obtain null distribution, test statistic, and p-value
  - i. Check conditions for CLT
  - ii. If conditions met, obtain null distribution and test-statistic, and determine distribution of test-statistic

- Conditions:
  - Independence: random sample
  - Approximate normality:  $n = 30$ , but no clear outliers
- So by CLT, null dist. is  $\bar{X} \sim N\left(38, \frac{\sigma}{\sqrt{30}}\right)$
- Since we don't know  $\sigma$ , we perform a  $t$ -test and obtain the following test-statistic:
  - $t = \frac{\bar{x}_{obs} - \mu_0}{\widehat{SE}} = \frac{38.6 - 38}{1.29/\sqrt{30}} = 2.543$
  - This test-statistic follows a  $t_{29}$  distribution



# Example: salinity (cont.)

iii. Use test-statistic to obtain p-value (draw picture and/or write code using appropriate distribution)



Want

$P(T \geq 2.54) + P(T \leq -2.54)$   
because  $H_A$  is two-sided!

```
1 p_val <- 2 * (1 - pt(t, df = n-1))  
2 p_val
```

[1] 0.01658569

# Example: salinity (cont.)

## 4. Decision and conclusion

- Since our p-value 0.017 is less than 0.05, we reject  $H_0$ .
- The data do provide sufficient evidence to suggest that the average salinity level in Bimini Lagoon is different from 38 ppt.

Let's code it up together!