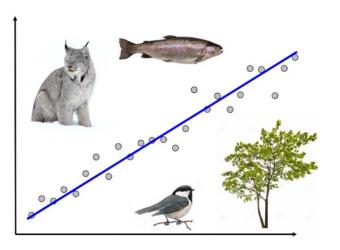
# NRC 290b Introduction to Quantitative Ecology

Week 7 — Tests for differences



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2019 - Fall

#### This week

#### Tuesday (Monday)

- MAP!
- Share our plots with structured feedback
- Finish reports and plots with edits

#### Wednesday

- Difference tests!
  - Test for normally distributed data
    - t-test/paired t-test
  - Tests for skewed data
    - U-test
    - Wilcoxon matched-pairs test

#### Plot feedback

What your group should present (no longer than 3 minutes):

- Your question and your hypotheses
- The year of your data
- What your plot visualizes

#### Feedback:

- What did we learn from the plot (and what about the plot helped us learn it)?
- What one thing could be improved to better visualize the answer to the question?

# Week 7 — Tests for differences

Part II - Wednesday

# Statistical testing

Which statistical test is right if you are looking for differences between two samples where you have less than 30 paired samples and the samples are normally distributed?

- a) Kruskal-Wallis Test for multiple samples
- b) Z-test for matched pairs
- c) T-test for matched pairs
- d) Mann-Whitney U-Test



# Statistical testing

Using a t-test to test the difference between two samples, you calculate a *p*-value of 0.02. If you using a 5% significance (alpha) level – what do you conclude?

- a) You reject the null hypothesis
- b) You accept the null hypothesis
- c) You don't reject the null hypothesis
- d) You reject the alternative hypothesis
- e) You don't reject the alternative hypothesis



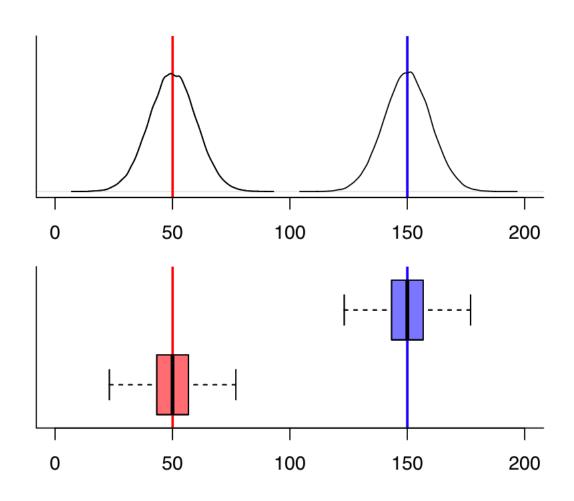
# Why do we conduct difference tests?



# Tests for differences

#### Ask the questions:

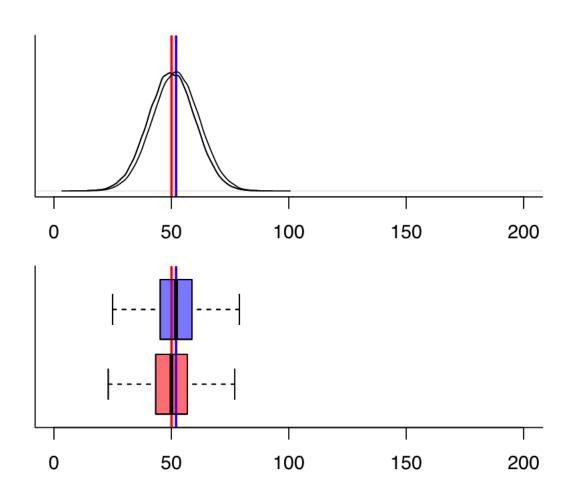
- Are the sample means different?
- Are the sample medians different?
- Are those differences statistically significant?



# Tests for differences

#### Ask the questions:

- Are the sample means different?
- Are the sample medians different?
- Are those differences statistically significant?



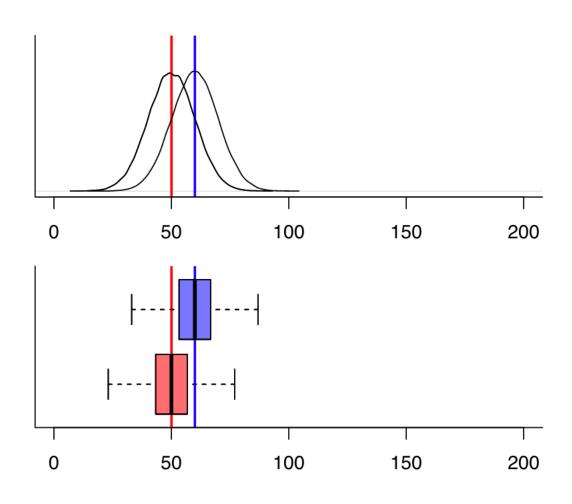
#### Tests for differences

#### Ask the questions:

- Are the sample means different?
- Are the sample medians different?
- Are those differences statistically significant?

#### Statistical tests:

- T-test (normally distributed data)\*
- U-test (skewed data)



### Student's t-test

Compares the means of two samples where  $H_0$ :  $\bar{x}_a = \bar{x}_b$ 

$$t = \frac{|\bar{x}_a - \bar{x}_b|}{\sqrt{\frac{s_a^2}{n_a} + \frac{s_b^2}{n_b}}}$$



#### Assumptions:

- Both samples are randomly collected, normally distributed, and independent
- Both samples have equal variances (i.e. homogeneity of variance)
- BUT the t-test is robust to slight deviations from these assumptions!

Answer the two questions using the t-test equation:

$$t = \frac{|\bar{x}_a - \bar{x}_b|}{\sqrt{\frac{s_a^2}{n_a} + \frac{s_b^2}{n_b}}}$$

If the absolute value of the difference in means is large, then t is...

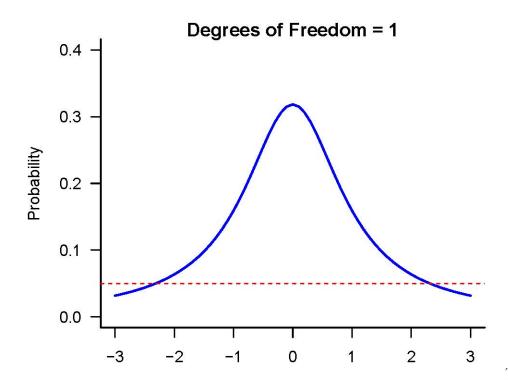
If the sum of the variances is large, then t is...

- a) Large, large
- b) Large, small
- c) Small, large
- d) Small, small



 Critical value of t is determined by your defined alpha (significance value) and degrees of freedom

$$df = (n_a - 1) + (n_b - 1)$$

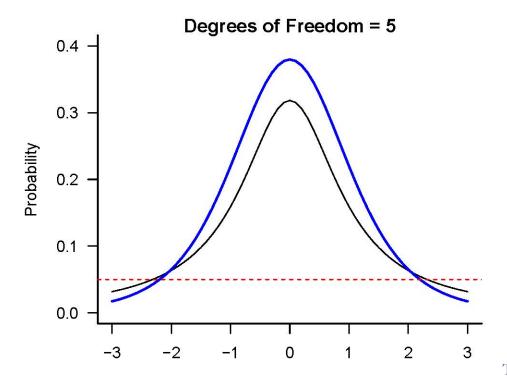


Tests for differences

Table A-	2	The t-Distribution				
df \α	0.10	0.05	0.025	0.01	0.005	
	0.20	0.10	0.05	0.02	0.01	
1	3.08	6.31	12.71	31.82	63.66	
2	1.89	2.92	4.30	6.96	9.92	
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4	1.53	2.13	2.78	3.75	4.60	
5	1.48	2.02	2.57	3.36	4.03	
6	1.44	1.94	2.45	3.14	3.71	
7	1.41	1.89	2.36	3.00	3.50	
8	1.40	1.86	2.31	2.90	3.36	
9	1.38	1.83	2.26	2.82	3.25	
10	1.37	1.81	2.23	2.76	3.17	
11	1.36	1.80	2.20	2.72	3.11	
12	1.36	1.78	2.18	2.68	3.05	
13	1.35	1.77	2.16	2.65	3.01	
14	1.35	1.76	2.14	2.62	2.98	
15	1.34	1.75	2.13	2.60	2.95	
16	1.34	1.75	2.12	2.58	2.92	
17	1.33	1.74	2.11	2.57	2.90	
18	1.33	1.73	2.10	2.55	2.88	
19	1.33	1.73	2.09	2.54	2.86	
20	1.33	1.72	2.09	2.53	2.85	
21	1.32	1.72	2.08	2.52	2.83	
22	1.32	1.72	2.07	2.51	2.82	
23	1.32	1.71	2.07	2.50	2.81	
24	1.32	1.71	2.06	2.49	2.80	
25	1.32	1.71	2.06	2.49	2.79	
26	1.31	1.71	2.06	2.48	2.78	
27	1.31	1.70	2.05	2.47	2.77	
28	1.31	1.70	2.05	2.47	2.76	
29	1.31	1.70	2.05	2.46	2.76	
30	1.31	1.70	2.04	2.46	2.75	
40	1.30	1.68	2.02	2.42	2.70	
60	1.30	1.67	2.00	2.39	2.66	
120	1.29	1.66	1.98	2.36	2.62	
00	1.28	1.64	1.96	2.33	2.58	

 Critical value of t is determined by your defined alpha (significance value) and degrees of freedom

$$df = (n_a - 1) + (n_b - 1)$$

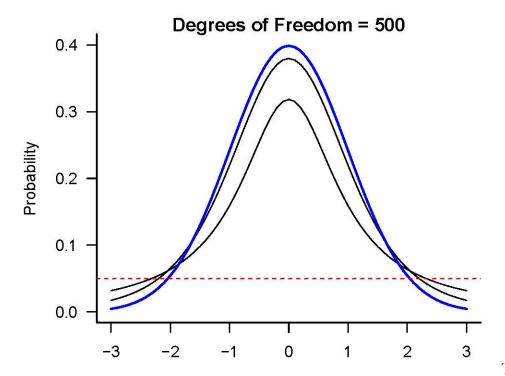


Tests for differences

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14	1.35	1.76	2.14	2.62	2.98	
15	1.34	1.75	2.13	2.60	2.95	
16	1.34	1.75	2.12	2.58	2.92	
17	1.33	1.74	2.11	2.57	2.90	
18	1.33	1.73	2.10	2.55	2.88	
19	1.33	1.73	2.09	2.54	2.86	
20	1.33	1.72	2.09	2.53	2.85	
21	1.32	1.72	2.08	2.52	2.83	
22	1.32	1.72	2.07	2.51	2.82	
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24	1.32	1.71	2.06	2.49	2.80	
25	1.32	1.71	2.06	2.49	2.79	
26	1.31	1.71	2.06	2.48	2.78	
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• Critical value of t is determined by your defined alpha (significance value) and degrees of freedom

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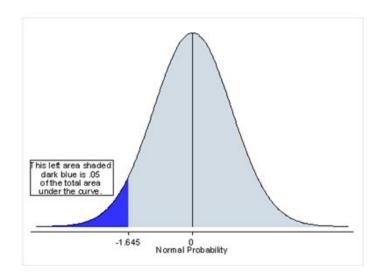


Tests for differences

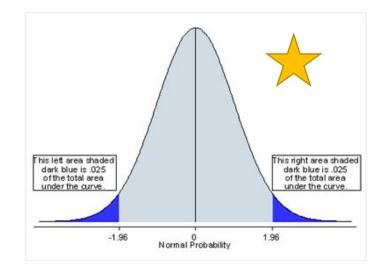
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23	1.32	1.71	2.07	2.50	2.81	
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• Critical value of t is determined by your defined alpha (significance value) and degrees of freedom

$$df = (n_a - 1) + (n_b - 1)$$



A one-tailed test will test either if the mean is significantly greater than x or if the mean is significantly less than x, but not both. The one-tailed test provides more power to detect an effect in one direction by not testing the effect in the other direction.



A two-tailed test will test both if the mean is significantly greater than x and if the mean significantly less than x. The mean is considered significantly different from x if the test statistic is in the top 2.5% or bottom 2.5% of its probability distribution, resulting in a p-value less than 0.05.

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24	1.32	1.71	2.06	2.49	2.80	
25	1.32	1.71	2.06	2.49	2.79	
26	1.31	1.71	2.06	2.48	2.78	
27	1.31	1.70	2.05	2.47	2.77	
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• Critical value of *t* is determined by your defined alpha (significance value) and degrees of freedom

$$df = (n_a - 1) + (n_b - 1)$$

• If *t* is **higher** than the given value in the table at your alpha and df, you **reject** the null hypothesis!

Special case: when you have a large sample size (n > 30) you use a very similar test, the z-test! Not discussed in this book.

Table A	-2	2 The <i>t</i> -Distribution				
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26	1.31	1.71	2.06	2.48	2.78	
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00	1.28	1.64	1.96	2.33	2.58	

### Paired t-test

What if your samples are paired?

Compares the means of two samples that are NOT independent (e.g., before/after study)

$$t = \frac{\overline{D}}{\sqrt{\frac{S_D^2}{n}}}$$

 $\overline{D}$  is the mean of the differences

 $S_D$  is the standard deviation of the differences

n is the number of paired samples

Assumptions:

- Both samples are normally distributed
- Both samples have equal variances/sd (i.e. homogeneity of variance)
- Sample sizes must be exactly the same!

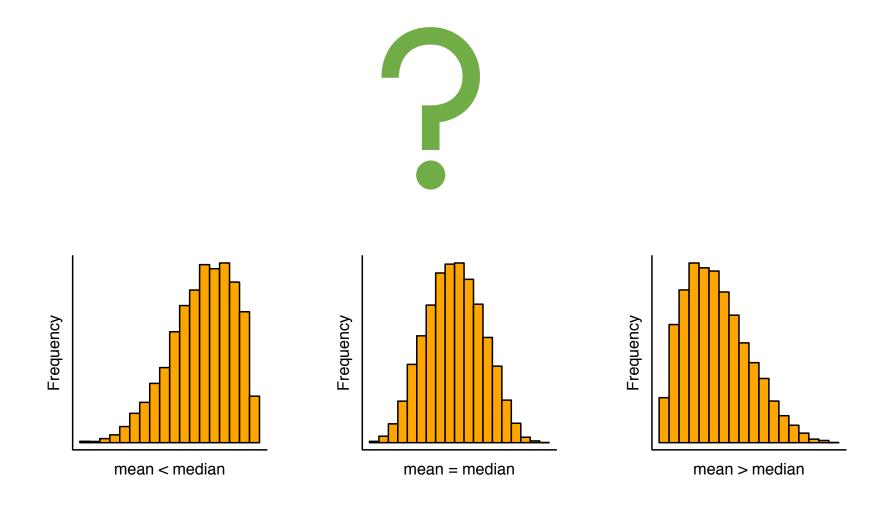
#### Paired t-test

Let's say you have a group of tree frogs you gave a vaccine to and you are trying to see if there is a difference in mucus production of those frogs before and after the vaccine... however, one of your treated frogs was eaten by a predator between your first measurement and your second — what do you do with the initial mucus value you recorded for this frog when conducting your paired t-test?

- a) Get rid of it
- b) Use it, no problem there
- c) Use it and duplicate another similar frogs' second reading
- d) Use it for both before and after treatment



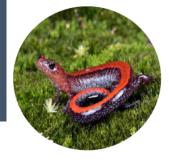
# What happens when our data aren't normally distributed?



#### Difference tests for skewed data

- U-test (for unpaired samples)
  - Like the t-test but uses ranks, median, and range to determine how much overlap there is between samples
  - LOWER U-values are more significant (opposite from t-test)
- Wilcoxon matched-pairs test (for paired samples)
  - Based on ranked differences (calculated differences first, then rank them)
  - Again, lower W-values are more likely to be significant

# Today's Exercise



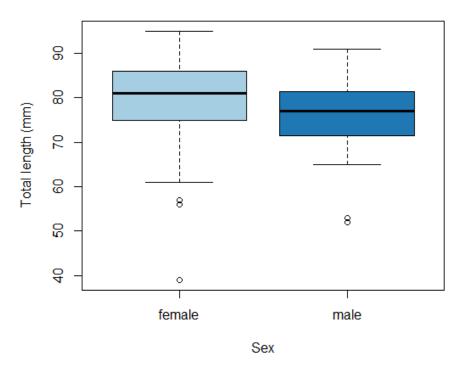
Back to the redbacked salamander (*Plethodon cinereus*) data! We're trying to answer the question:

# Is the average length of salamanders significantly different between salamanders identified as male or female?

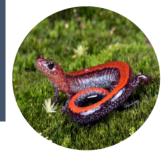
#### To do so:

- 1. Download my code from moodle labeled mander\_sex\_differences.R
- 2. Go through the code and add comments each place you see a hashtag and code where it needs code. The comments should tell me:
  - 1. What each section of of code does
  - 2. Answer the questions outlined in the code
- 3. Upload your *individual* code to moodle

#### Distribution of total salamander lengths by sex



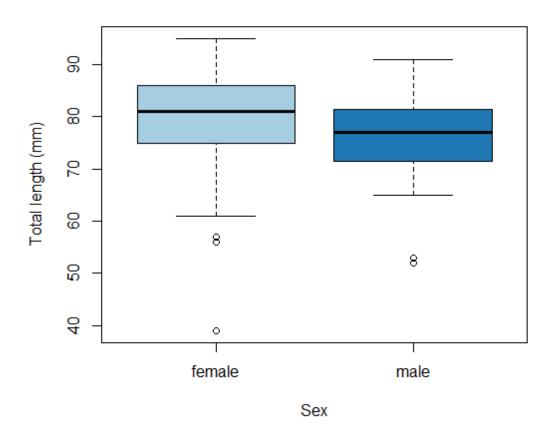
# Quick aside – outliers?



In R (and many other places), data points that are >1.5x the interquartile range (IQR) from the .25 and .75 quartiles are identified as *possible* outliers (something to explore)

- Plotted here as the open circles
- Identifying data points as possible outliers doesn't mean they are "bad data"!

#### Distribution of total salamander lengths by sex



# For Monday:



- 1) Read Ch 10 (Skip 10.2) in Gardner (2017) Differences between more than 2 samples
- 2) Answer the individual evaluation questions on moodle

All before 11:55pm on Sunday