

Introduction to Statistical analysis

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Outline

Introduction

Hypothesis Testing

Tests for continuous variables: T-tests

One-sample t-test

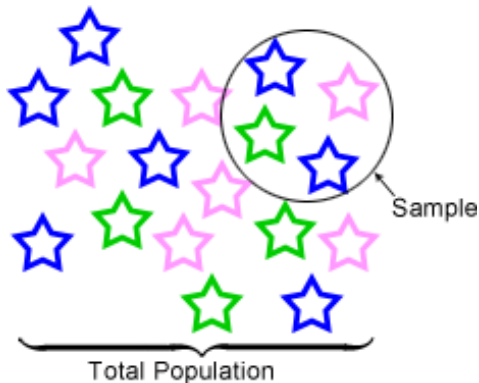
Two-sample t-test

Independent two-sample t-test

Paired two-sample t-test

The point of statistics

- ▶ Rarely feasible to study the whole population that we are interested in, so we take a sample instead
- ▶ Assume that data collected represents a larger population
- ▶ Use sample data to make conclusions about the overall population



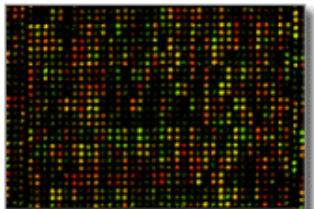
Data type - example

- Success / failure of achieving a task for a mouse which may be wild-type or knock-out, male or female, 2, 4 or 6 months old



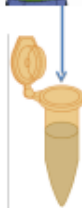
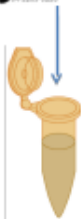
Data type - example

- ▶ Gene expression in each cell sample which may be one of five cell-types (A,B,C,D,E)



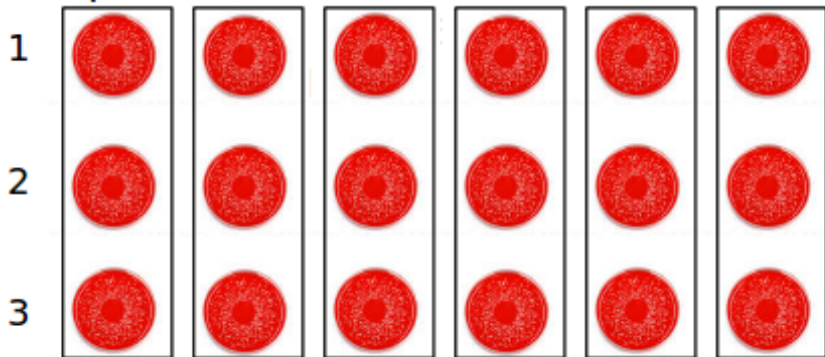
Data type - example

- ▶ The number of bacteria for each subject which may be a cancer patient or a normal



Measurements: Dependent / Independent?

- ▶ 18 measurements; from repeating an experiment 6 times, each time in triplicate



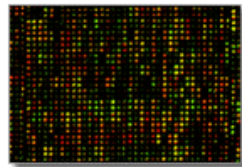
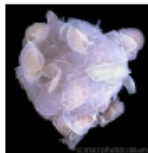
Measurements: Dependent / Independent?

- ▶ Measuring blood pressure before and after treatment for 30 patients

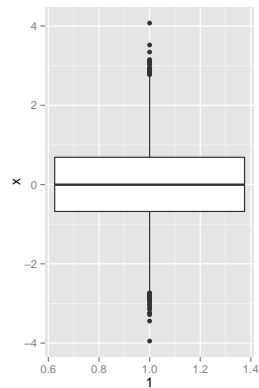
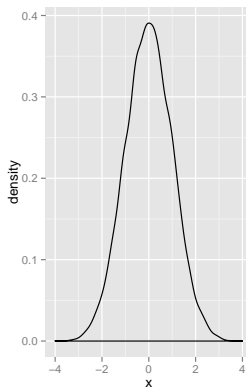
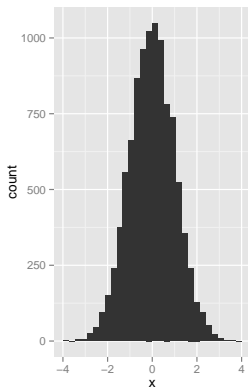


Measurements: Dependent / Independent?

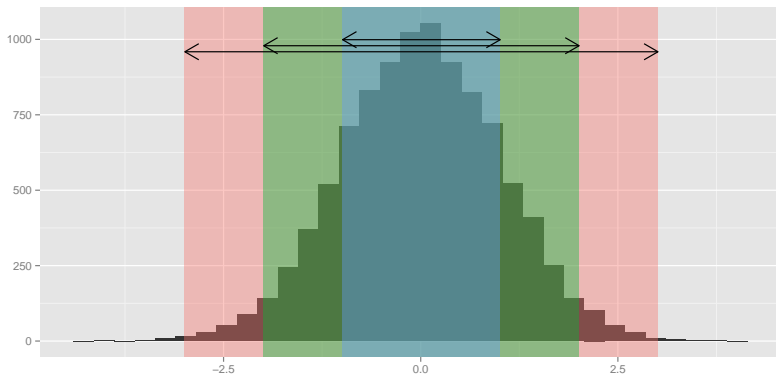
- ▶ Measuring gene expression in each cell sample, which may be one of five cell-types from cancer patients or normal individuals



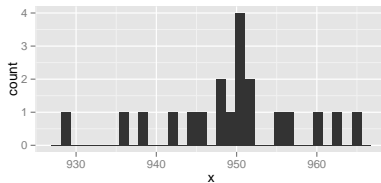
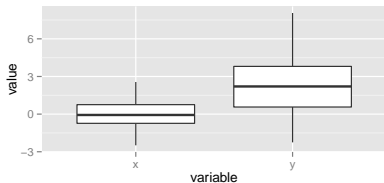
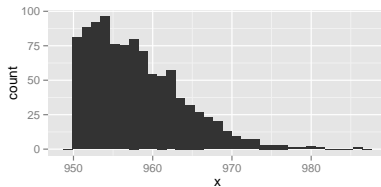
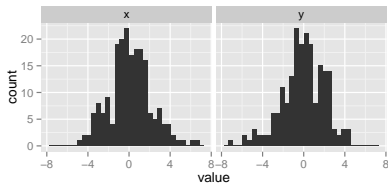
Continuous Data - Distribution



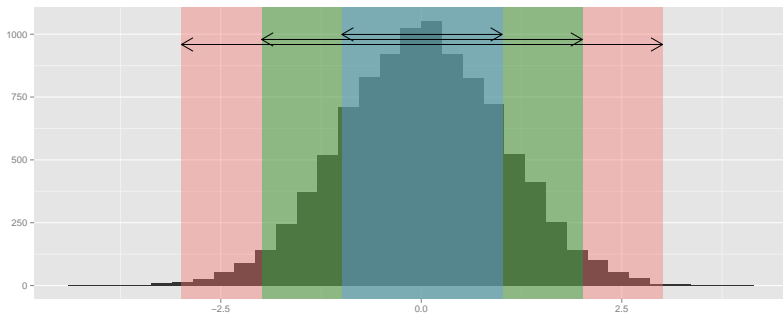
Continuous Data - Distribution



Continuous or not??



Continuous Data - Descriptive Statistics

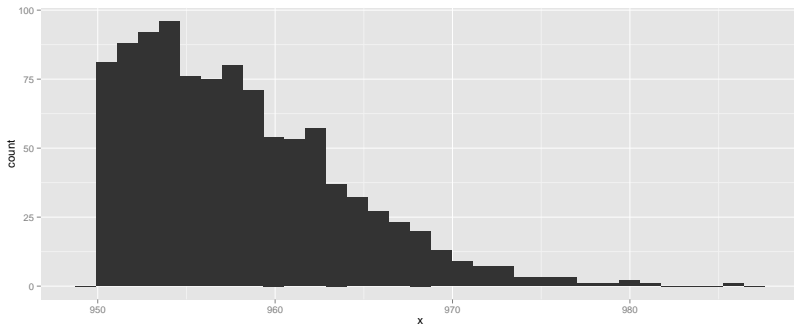


- ▶ Measures of location and spread
- ▶ Mean and standard deviation

- ▶ $\bar{X} = \frac{X_1 + X_2 + \dots + X_n}{n}$

- ▶ $s.d = \sqrt{\frac{1}{N-1} \sum_{i=1}^N (x_i - \bar{x})^2}$

Continuous Data - Descriptive Statistics



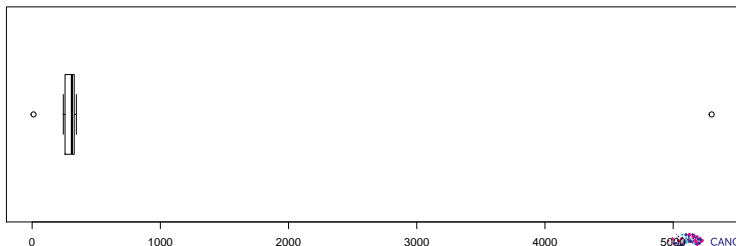
- ▶ Median: middle value
- ▶ Lower quartile: median bottom half of data
- ▶ Upper quartile: median top half of data

Continuous Data - Descriptive Statistics Example

E.g. No. of facebook friends for 7 colleagues

311, 345, 270, 310, 243, 5300, 11

- ▶ Measures of location and spread
- ▶ Mean and standard deviation
 - ▶ $\bar{X} = \frac{X_1 + X_2 + \dots + X_n}{n} = 970$
 - ▶ $s.d = \sqrt{\frac{1}{N-1} \sum_{i=1}^N (x_i - \bar{x})^2} = 1912.57$
- ▶ Median and interquartile range
 - ▶ 11, 243, 270, 310, 311, 345, 5300



Continuous Data - Descriptive Statistics Example

E.g. No. of facebook friends for 7 colleagues

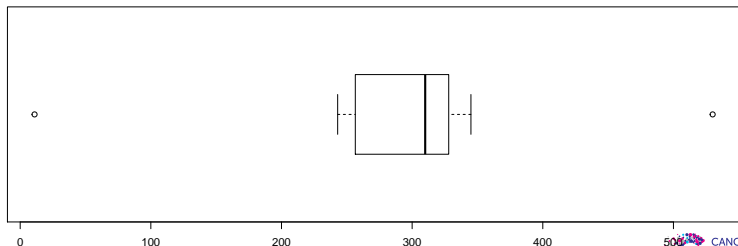
311, 345, 270, 310, 243, 530, 11

- ▶ Measures of location and spread
- ▶ Mean and standard deviation
 - ▶ $\bar{X} = \frac{X_1 + X_2 + \dots + X_n}{n} = 289$
 - ▶ $s.d = \sqrt{\frac{1}{N-1} \sum_{i=1}^N (x_i - \bar{x})^2} = 153.79$
- ▶ Median and interquartile range
 - ▶ 11, 243, 270, 310, 311, 345, 530

Continuous Data - Descriptive Statistics Example

E.g. No. of facebook friends for 7 colleagues
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- ▶ Median and interquartile range
 - ▶ 11, 243, 270, 310, 311, 345, 530



Categorical Data

- ▶ Summarised by counts and percentages
- ▶ Examples
 - ▶ 19 / 82 (23%) subjects had Grade IV tumour
 - ▶ 48 / 82 (58%) subjects had Diarrhoea as a Adverse Event

Standard Deviation and Standard Error

- ▶ Commonly confused
- ▶ Standard deviation
 - ▶ Measure of spread of data
 - ▶ Used for describing **population**
- ▶ Standard error
 - ▶ Variability of the mean from repeated sampling
 - ▶ Precision of the mean
 - ▶ Used to calculate confidence interval
- ▶ SD: How widely scattered measurements are
- ▶ SE: Uncertainty in **estimate** of sample mean

Confidence intervals for the mean

- ▶ Confidence Interval (CI) is a random interval
- ▶ In repeated experiments
 - ▶ 95% of time cover the mean
- ▶ Looser interpretation 95% of time 95% CI:
 $(\bar{X} - 1.96 \times se, \bar{X} + 1.96 \times se)$

For facebook friends data:

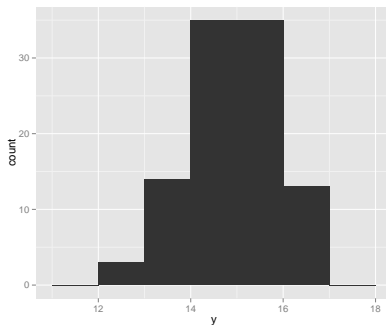
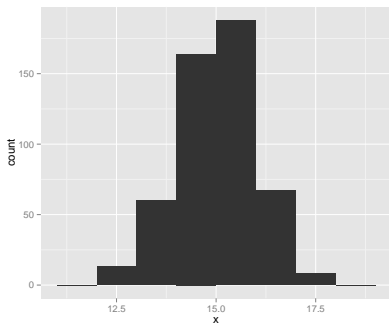
$$se = \frac{sd}{\sqrt{(n)}} = \frac{154}{\sqrt{7}} = 58$$

Mean 289, 95% CI (289 - (1.96 × 58), 289 + (1.96 × 58))

Mean 289, 95% CI (175, 402)

Confidence intervals

- ▶ As number of observations goes up....
- ▶ Standard deviation stays **the same**...
- ▶ But standard error **goes down**....



Basic set-up

- ▶ Formulate a *null hypothesis* - H_0
- ▶ e.g. "*The difference in treatment before and after treatment = 0*"
- ▶ Calculate a test statistic from the data under the null hypothesis
- ▶ Determine whether the test statistic is more extreme than expected under the null hypothesis
- ▶ Reject or do not reject the null hypothesis
- ▶ Absence of evidence is not evidence of absence (Bland and Altman, 1995)

Example

Lady Tasting Tea (Randomised experiment by Fisher)

- ▶ Randomly-ordered 8 cups of tea
 - ▶ 4 were prepared by adding milk first
 - ▶ 4 were prepared by adding tea first
- ▶ Task: Lady had to select the 4 cups of one particular method
- ▶ H_0 Lady no such ability
- ▶ Test Statistic: number of successes in selecting 4 cups
- ▶ Result: Lady got all 4 cups correct
- ▶ **Reject the null hypothesis**

Errors

	Null hypothesis does not hold	Null hypothesis holds
Reject null hypothesis	Correct True positive	Wrong False positive
Do not reject null hypothesis	Wrong False negative	Correct True negative

Significance

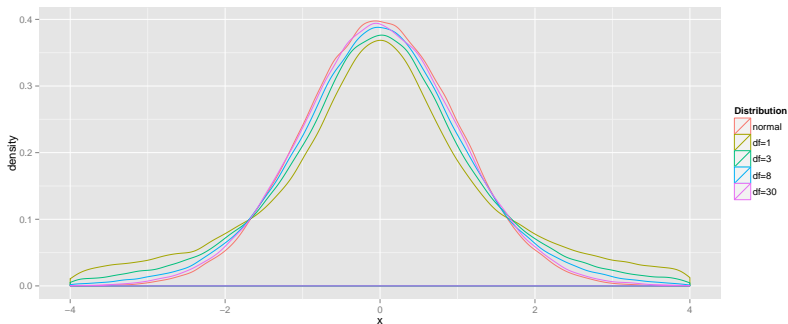
level, sample size, different of interest, variability of the observations

Be aware of issues of multiple testing

Various flavours

- ▶ One-sample t-test: e.g. H_0 : mean = 5
- ▶ Independent two-sample t-test: e.g. H_0 : mean of sample 1 = mean of sample 2
- ▶ Paired two-sample t-test: e.g. H_0 : mean difference between pairs = 0

T-distributions



Does mean = X?

- ▶ **Research question:** Published data suggests that the microarray failure rate for a particular supplier is 2.1%
- ▶ Genomics want to know if this holds true in their own lab

Formulating the question

- ▶ Null hypothesis, H_0 Mean monthly failure rate = 2.1%
- ▶ Alternative hypothesis, H_1 Mean monthly failure rate \neq 2.1%
- ▶ Tails: two-tailed
- ▶ Either *reject* or *do not reject* the null hypothesis - **never accept the alternative hypothesis**

The Data

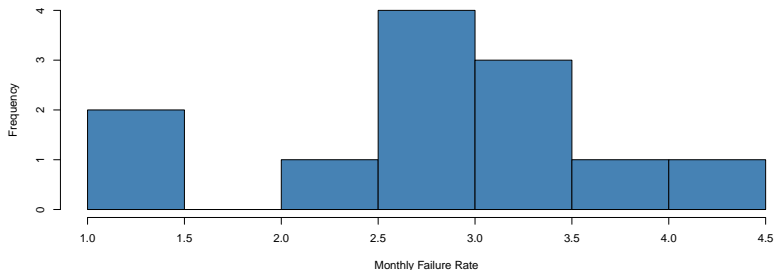
	Month	Monthly.failure.rate
1	January	2.90
2	February	2.99
3	March	2.48
4	April	1.48
5	May	2.71
6	June	4.17
7	July	3.74
8	August	3.04
9	September	1.23
10	October	2.72
11	November	3.23
12	December	3.40

Summary Statistics

mean = $(2.9 + \dots + 3.40)/12 = 2.841$

Standard deviation = 0.837

- ▶ Observations must be independent
- ▶ Observations must be normally distributed

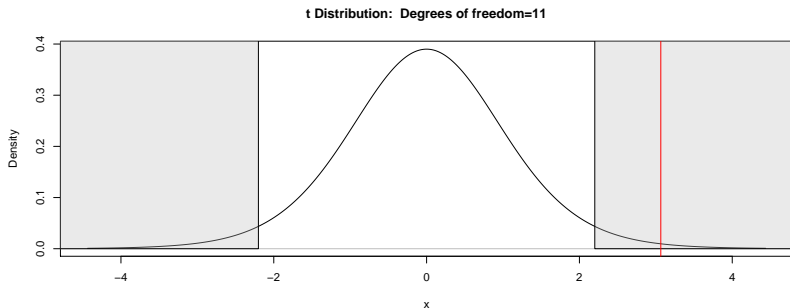


Results

$$\text{Test statistic } t_{n-1} = t_{11} = \frac{\bar{x} - \mu_0}{s.d./\sqrt{n}} = \frac{2.84 - 2.10}{s.e.(\bar{x})} = 3.065$$

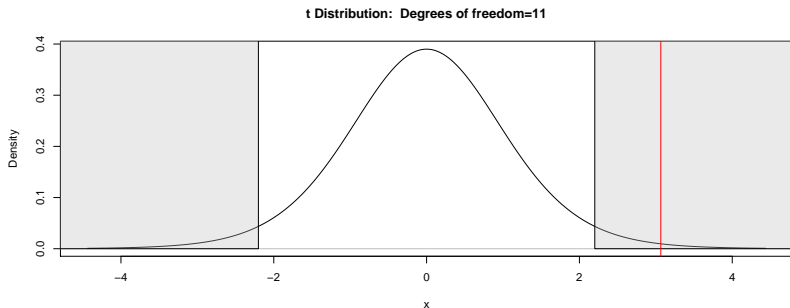
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p:value = 0.011 Reject H_0 and conclude that mean monthly failure rate in Genomics is not 2.1%

Independent two-sample t-test

Does mean of group A = mean of group B?

e.g. **Research question:** 40 male mice (20 of breed A and 20 of breed B) were weighed at 4 weeks old.

Does the weight of 4 week old male mice depend on breed?

The data

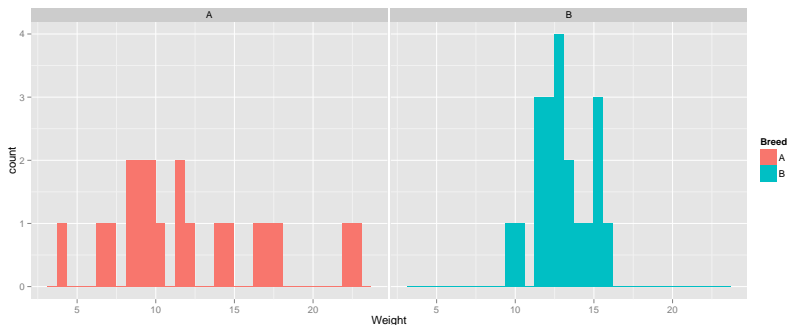
	BreedAMouse	WeightA	BreedBMouse	WeightB
1	1	22.77	21	15.51
2	2	9.08	22	12.93
3	3	9.80	23	11.50
4	4	8.13	24	16.07
5	5	16.54	25	15.51
6	6	11.36	26	15.16
7	7	11.47	27	11.25
8	8	22.25	28	13.65
9	9	14.04	29	14.28
10	10	17.12	30	13.21
11	11	6.32	31	10.28
12	12	17.51	32	12.41
13	13	9.87	33	9.63
14	14	12.41	34	14.75
15	15	7.39	35	12.56
16	16	9.23	36	13.02
17	17	4.06	37	12.33

Data summary

Mean of breed A: 12.12 **Mean** of breed B: 12.99

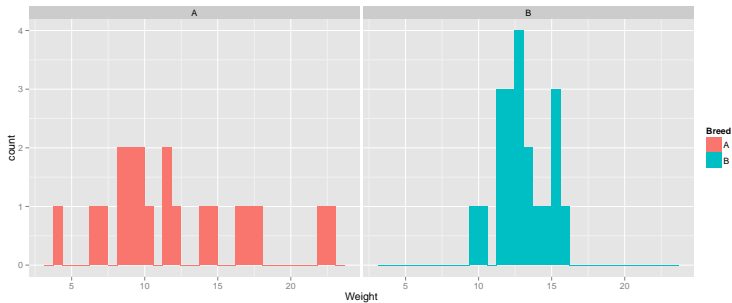
Standard Deviation of breed A: 5.05

Standard Deviation of breed B: 1.8



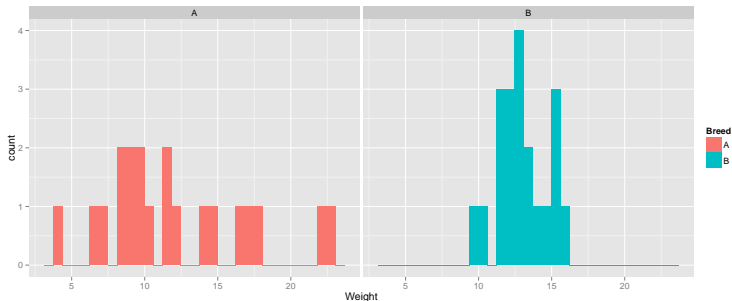
Checking assumptions

- ▶ Observations are independent
- ▶ Observations are normally distributed



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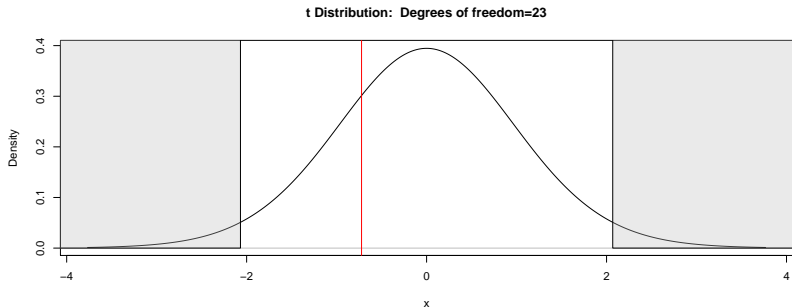


- ▶ Use Welch's correction if variances are different (alters the t-statistic and degrees of freedom)

Test results

$$t_{df} = \frac{\bar{X}_A - \bar{X}_B}{s.e.(\bar{X}_A - \bar{X}_B)} = -0.7229$$

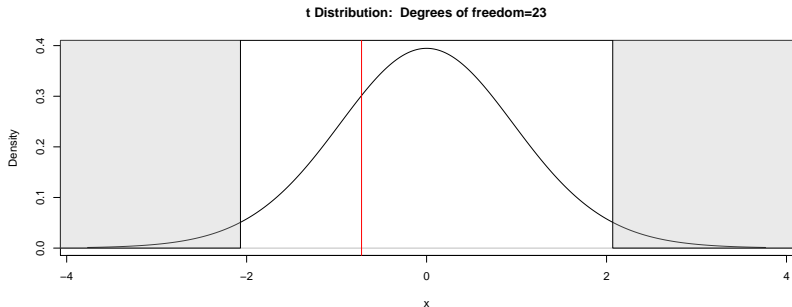
df = 23 (with Welch's correction)



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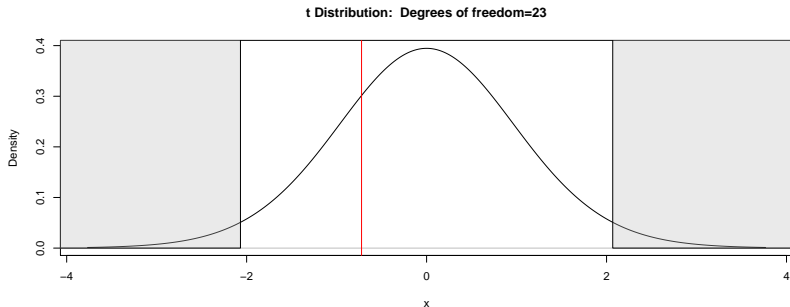


p-value: 0.4768

Test results

$$t_{df} = \frac{\bar{X}_A - \bar{X}_B}{s.e.(\bar{X}_A - \bar{X}_B)} = -0.7229$$

df = 23 (with Welch's correction)



***p*-value: 0.4768 Do not reject H_0 . There is no evidence for a difference in weight between breeds**

- ▶ Null hypothesis, H_0 Cellularity at site A = Cellularity at site B
- ▶ Alternative hypothesis, H_1 Cellularity at site A \neq Cellularity at site B
- ▶ Tails: two-tailed
- ▶ Either *reject* or *do not reject* the null hypothesis - **never accept the alternative hypothesis**

Null hypothesis

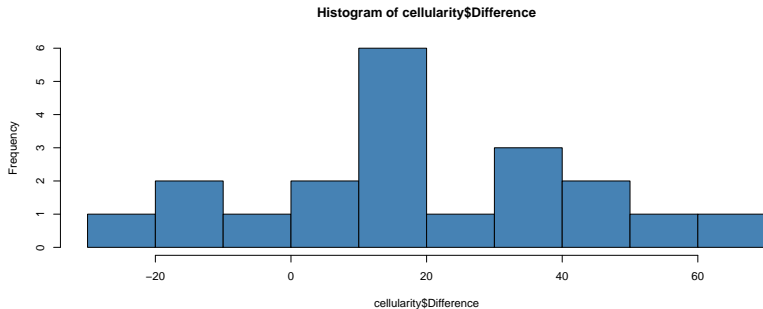
- ▶ H_0 Cellularity at site A = Cellularity at site B
OR
- ▶ H_0 Cellularity at site A - Cellularity at site B = 0

The data

	Ovarian	Peritoneal	Difference
1	1201.33	1155.98	45.35
2	1029.64	1020.82	8.82
3	895.57	881.21	14.36
4	842.14	830.78	11.36
5	903.07	897.06	6.01
6	1311.57	1262.73	48.84
7	833.52	823.06	10.46
8	1007.66	951.01	56.65
9	1465.51	1450.98	14.53
10	967.82	978.15	-10.33
11	812.72	778.26	34.46
12	884.08	823.57	60.51
13	1358.56	1335.78	22.78
14	1280.10	1293.91	-13.81
15	942.38	925.75	16.63
16	884.33	891.34	-7.01
17	930.09	892.02	38.07

Mean difference: 19.139

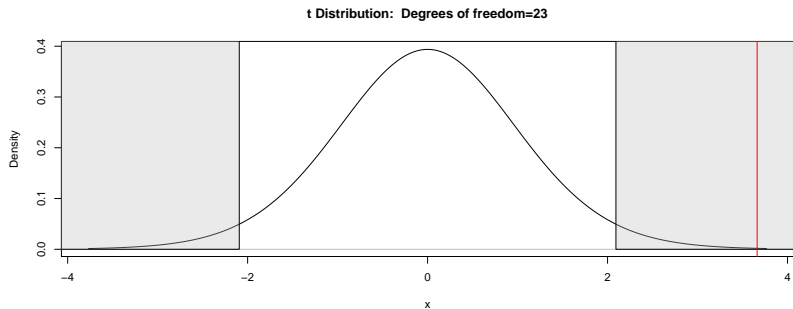
- ▶ Observations are independent
- ▶ The paired differences are normally distributed



Test results

$$t_{n-1} = t_{19} = \frac{\bar{X}_{A-B}}{s.e.(\bar{X}_{A-B})} = 3.6624$$

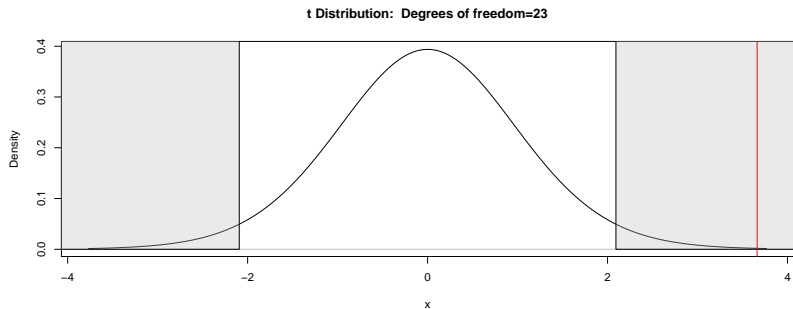
df = 19



Test results

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$$df = 19$$

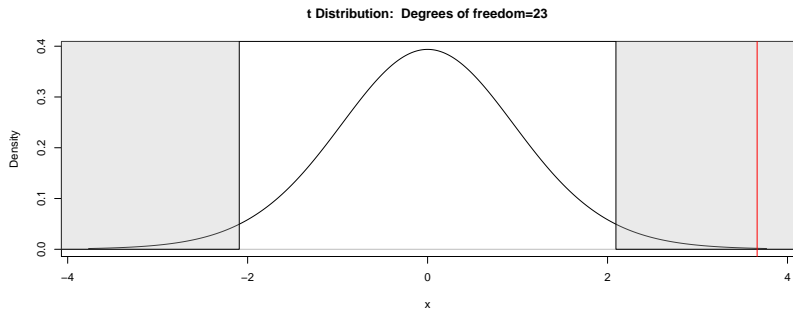


p-value: 0.0017

Test results

$$t_{n-1} = t_{19} = \frac{\bar{X}_{A-B}}{s.e.(\bar{X}_{A-B})} = 3.6624$$

df = 19



p-value: 0.0017 **Reject H_0** . There is evidence that cellularity at site A \neq cellularity at site B

Summary - Continuous variables

- ▶ One-sample t-test Use when we have *one group*
- ▶ Independent two-sample t-test Use when we have two independent groups. A Welch correction may be needed if the two groups have different spread.
- ▶ Paired two-sample t-test Use when we have *two non-independent groups*
- ▶ Non-parametric tests or transformations Use when we *cannot assume normality*