Inference for numerical data

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Central Limit Theorem Recap

If:

- Samples are independent,
- Sample size is bigger or equal to 30,
- Population distribution is not strongly skewed.

Then:

Point estimate distribution can be approximated by Normal Distribution with mean equal to population mean and standard deviation equal to Standard Error.

Confidence Intervals

95% confidence interval

point estimate
$$\pm~1.96\cdot SE$$

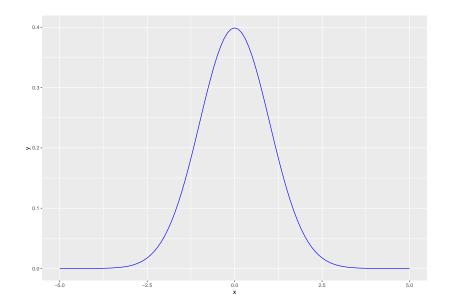
99% confidence interval

point estimate \pm 2.58 \cdot SE

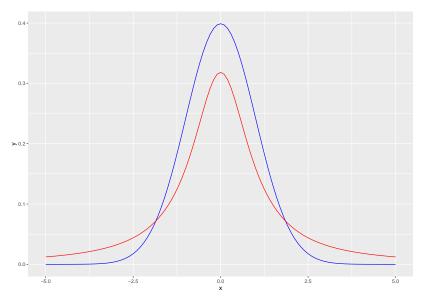
Normal distribution vs Student-t distribution

What if the sample is not even 30?

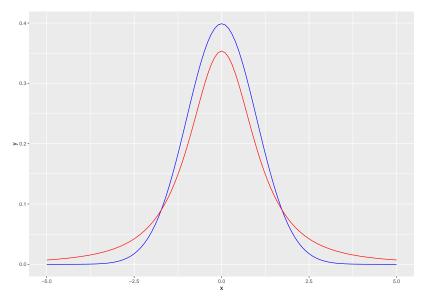
Normal distribution vs Student-t distribution



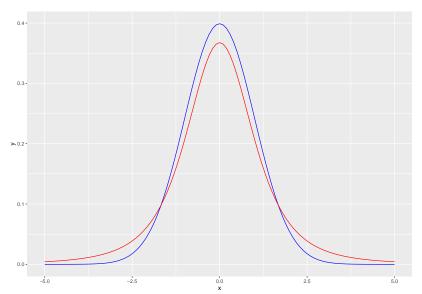
Normal distribution vs Student-t distribution N(0,1) T(1)



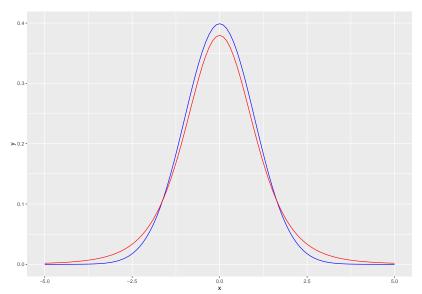
Normal distribution vs Student-t distribution N(0,1) T(2)



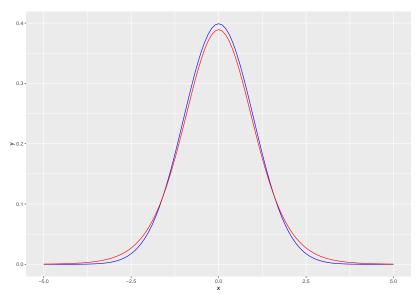
Normal distribution vs Student-t distribution N(0,1) T(3)



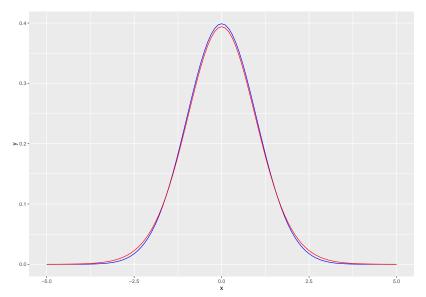
Normal distribution vs Student-t distribution N(0,1) T(5)



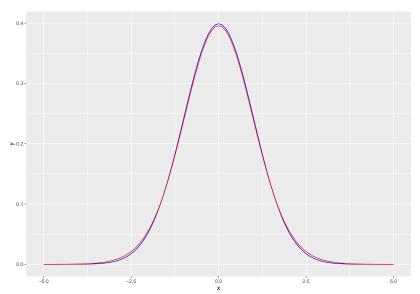
Normal distribution vs Student-t distribution N(0,1) T(10)



Normal distribution vs Student-t distribution N(0,1) T(20)



Normal distribution vs Student-t distribution N(0,1) T(30)



Normal distribution vs Student-t distribution

$$Z$$
-score = $\frac{x-\bar{x}}{s}$ \longrightarrow T -score = $\frac{x-\bar{x}}{s}$

To read out probability:

$$mean = 0$$
, $SD = 1 \longrightarrow df = n - 1$

T-score is sometimes also called T-statistic.

Central Limit Theorem Recap

If:

- Samples are independent,
- Sample size is bigger or equal to 30,
- Population distribution is not strongly skewed.

Then:

Normalized point estimate distribution can be approximated by Student-T distribution with degrees of freedom equal to number of cases minus 1.

- 1) Set-up the hypothesis:
- $ightharpoonup H_0$: Position of no change
- \blacktriangleright H_A : Alternative to H_0 One-sided and two-sided hypothesis test.
- 2) Assume threshold values:
- ightharpoonup α -significance level typicaly 0.05
- 2.1) Check CLT conditions
 - 3) Calculate the Results

p-value - probability of something as different as point estimate existing within H_0 sampling distribution.

Draw conclusions

- 1) Set-up the hypothesis:
- $ightharpoonup H_0$: Position of no change
- $ightharpoonup H_A$: Alternative to H_0

One-sided and two-sided hypothesis test.

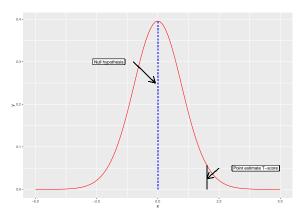
- 2) Assume threshold values:
- ightharpoonup α -significance level typicaly 0.05

- 3) Calculate the Results
- ightharpoonup p-value probability of something as different as point estimate existing within H_0 sampling distribution.

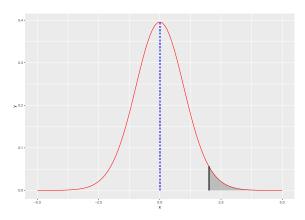
First calculate point estimate and SE, then:

$$\textit{T-score} = \frac{\text{point estimate} - \text{null hypothesis}}{\textit{SE}}$$

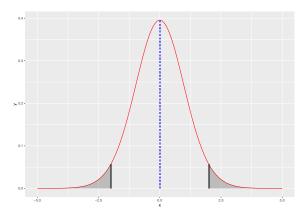
- 3) Calculate the Results
- **P**-value probability of something as different as point estimate existing within H_0 sampling distribution.



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- 3) Calculate the Results
- **P**-value probability of something as different as point estimate existing within H_0 sampling distribution.



4) Draw conclusions

$$p$$
 – value $> \alpha$ – significance level

We keep the Null Hypothesis and reject the Alternative (We failed to reject null hypothesis). The difference is not significant.

$$p-$$
 value $< \alpha -$ significance level

We reject Null Hypothesis in favour of Alternative. The difference is significant.

Type 1 and Type 2 hypothesis errors.

	H ₀ is True	H_A is True
Reject H_0	Type I Error	correct
Accept H ₀	correct	Type II error

Within Hypothesis we compare a sample point estimate to a **set value**.

The sample must fulfill conditions of CLT.

$$H_0: \mu = \mathsf{set} \ \mathsf{value}$$

$$H_A: \mu \neq \text{set value}$$

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

Kiwi dataset.

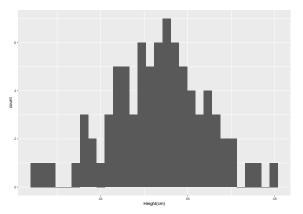
We want to check, if mean male Great Spotted height is 45 cm.

 H_0 : population mean height of male kiwis is 45 cm

 H_A : population mean height of male kiwis is not 45 cm

CLT Conditions?

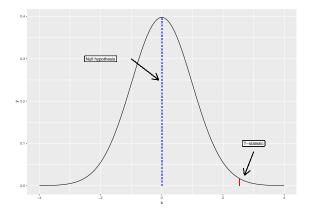
Number of male Great Spotted Kiwis is 82.



```
(point estimate <- mean(kiwi GS M$`Height(cm)`))
## [1] 45.30854
(SE <- sd(kiwi_GS_M$`Height(cm)`)/sqrt(nrow(kiwi_GS_M)))
## [1] 0.1213097
(df <- nrow(kiwi GS M) - 1)
## [1] 81
```

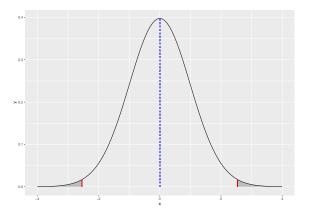
```
(t_statistic <- (point_estimate - 45)/SE)
```

[1] 2.543379



```
(p_value \leftarrow 2*(1- pt(t_statistic, df = df)))
```

[1] 0.01288087



We reject null hypothesis in favour of Alternative.

Mean Male Great Spotted Kiwi is not 45 cm tall.

Other way of conducting the t test:

```
t.test(kiwi_GS_M$`Height(cm)`, mu = 45)
##
##
    One Sample t-test
##
## data: kiwi GS M$'Height(cm)'
## t = 2.5434, df = 81, p-value = 0.01288
## alternative hypothesis: true mean is not equal to 45
## 95 percent confidence interval:
## 45.06717 45.54990
## sample estimates:
## mean of x
## 45.30854
```

Within Hypothesis we compare a sample point estimate to a set value.

The sample must fulfill conditions of CLT.

Difference of means t-test

Used to compare two different groups.

Category or sample.

Both samples must fulfill conditions of CLT.

Difference of means t-test

- ► Male and Females
- ► Regular American Males and NBA players
- ► Great Spotted Kiwis and Tokoeka Kiwis
- SDU and UCL students

$$H_0: \mu_1 = \mu_2 \longrightarrow \mu_1 - \mu_2 = 0$$

$$H_A: \mu_1 \neq \mu_2 \longrightarrow \mu_1 - \mu_2 \neq 0$$

$$SE = \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$$

Degrees of freedom

Welch-Satterthwaite equation

$$df = \frac{\left(\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}\right)^2}{\left(\frac{s_1^2}{n_1}\right)^2 + \left(\frac{s_2^2}{n_2}\right)^2}$$

or... smaller of the two.

Kiwi dataset.

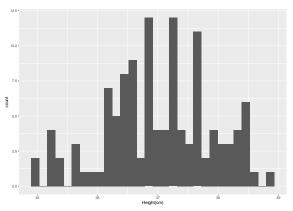
We want to check, if mean male Great Spotted are same height as mean male Tokoeka.

$$H_0$$
: $mean_{GS} - mean_{Tokoeka} = 0$

$$H_A$$
: $mean_{GS} - mean_{Tokoeka} \neq 0$

CLT Conditions?

Number of male Great Spotted Kiwis is 82, Number of male Tokoekas is 119.

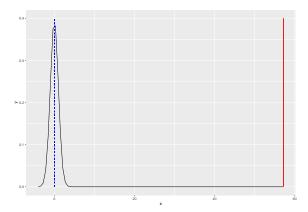


[1] 81

```
(point_estimate <- mean(kiwi_GS_M$`Height(cm)`) -
   mean(kiwi_T_M$`Height(cm)`) )
## [1] 8.295091
(SE<-sqrt((sd(kiwi GS M$`Height(cm)`)^2/nrow(kiwi GS M)) +
          (sd(kiwi_T_M$`Height(cm)`)^2/nrow(kiwi_T_M))))
## [1] 0.1452033
(df <- nrow(kiwi GS M) - 1)
```

```
(t_statistic <- (point_estimate - 0)/SE)</pre>
```

[1] 57.12745



```
(p_value < -2*(1-pt(t_statistic, df = df)))
```

[1] 0

We reject null hypothesis in favour of Alternative. Mean Male Great Spotted Kiwis are of different height than Tokoekas.

Other way of conducting the t test:

```
t.test(kiwi_GS_M$`Height(cm)`, kiwi_T_M$`Height(cm)`)
##
   Welch Two Sample t-test
##
##
## data: kiwi_GS_M$'Height(cm)' and kiwi_T_M$'Height(cm)'
## t = 57.127, df = 147.33, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not
## 95 percent confidence interval:
## 8.008141 8.582041
## sample estimates:
## mean of x mean of y
## 45.30854 37.01345
```

Note different df value.

$$df = \frac{\left(\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}\right)^2}{\frac{\left(\frac{s_1^2}{n_1}\right)^2}{n_1 - 1} + \frac{\left(\frac{s_2^2}{n_2}\right)^2}{n_2 - 1}}$$

```
M.v <- var(kiwi_GS_M$`Height(cm)`)
M.n <- nrow(kiwi_GS_M)
T.v <- var(kiwi_T_M$`Height(cm)`)
T.n <- nrow(kiwi_T_M)
(((M.v/M.n) + (T.v/T.n))^2)/
  ((((M.v/M.n)^2)/(M.n-1)) + (((T.v/T.n)^2)/(T.n-1)))</pre>
```

[1] 147.3307

##

Other way of conducting the t test:

```
hm <- kiwi %>% filter(Species_code == 'GS' |
                        Species code == 'Tok') %>%
  filter(Gender == 'M')
t.test(`Height(cm)`~Species code, data = hm)
##
##
   Welch Two Sample t-test
##
## data: Height(cm) by Species code
## t = 57.127, df = 147.33, p-value < 2.2e-16
## alternative hypothesis: true difference in means between
## 95 percent confidence interval:
## 8.008141 8.582041
## sample estimates:
## mean in group GS mean in group Tok
```

37.01345

45.30854

Used to compare two different groups.

Category or sample.

Both samples must fulfill conditions of CLT.

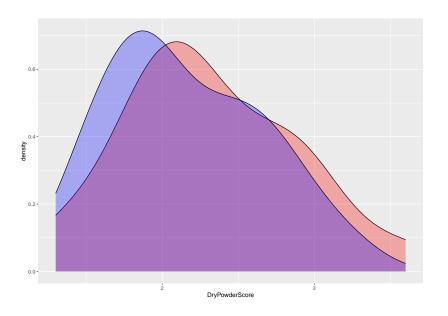
Used to compare two observations within same cases.

- Consumer grades of different product types,
- Prices of products in between shops,

H₀: mean DryPowderScore is equal to mean PowderInOilScore

 \mathcal{H}_A : mean DryPowderScore is not equal to mean PowderInOilScore

ID	DryPowderScore	PowderInOilScore
1	2.0	1.9
2	2.8	2.4
3	1.3	1.5
4	1.8	1.8
5	1.9	1.8



##

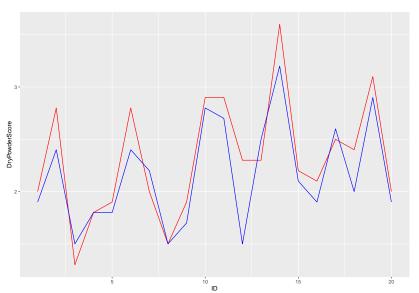
If we disregard paired observation and use simple difference of means scenario:

```
t.test(anti$DryPowderScore, anti$PowderInOilScore)
```

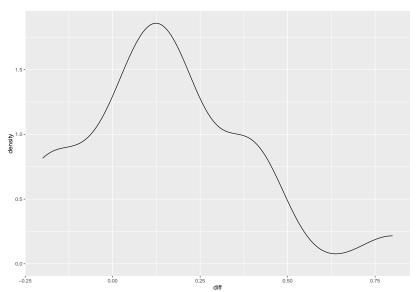
```
##
##
   Welch Two Sample t-test
##
## data: anti$DryPowderScore and anti$PowderInOilScore
## t = 0.88603, df = 37.417, p-value = 0.3813
## alternative hypothesis: true difference in means is not
## 95 percent confidence interval:
## -0.1928938 0.4928938
## sample estimates:
## mean of x mean of y
      2.315 2.165
```

we would keep null hypothesis.

But if we look at the data in a different way



So we compute difference variable:



$$H_0: \mu_{\mathit{diff}} = 0$$

$$H_A$$
 : $\mu_{diff} \neq 0$

$$SE = \frac{s_{diff}}{\sqrt{n}}$$

Degrees of freedom is again number of cases minus 1.

Back to antierspirant data

[1] 0.05548826

```
(point_estimate <- mean(anti$diff))
## [1] 0.15

(SE <- sd(anti$diff)/sqrt(nrow(anti)))</pre>
```

Back to antierspirant data

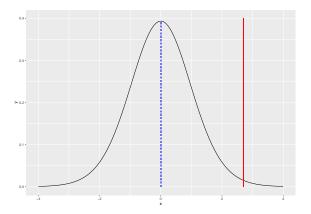
```
(df <- nrow(anti) - 1)
## [1] 19

(t_statistic <- (point_estimate - 0)/SE)</pre>
```

[1] 2.703274

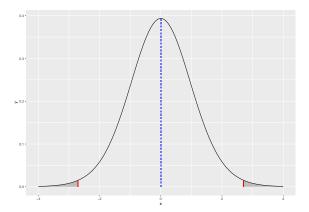
```
(p_value \leftarrow 2*(1- pt(t_statistic, df = df)))
```

[1] 0.01408939



```
(p_value \leftarrow 2*(1- pt(t_statistic, df = df)))
```

[1] 0.01408939



Lars was 6 years old, when on a fishing trip with his dad he caught 3 Atlantic herrings. All were almost 50 cm long, 1.5 kg heavy. Since then (and Lars is now 50) he always said, that the average Atlantic herring is 50 cm long and weights 1.5 kg. How would you test this hypothesis? How would you gather data, formulate hypothesis and which statistical test would you use?

Who loves licorice more: Danes or Swedes? An online survey was conducted throughout Sweden and Denmark to find licorice king. The participants were asked to grade, on the scale 1-10, how much do they love licorice. You are now in possession of the dataset. How do you decide who is the Scandinavian licorice king?

Gammeldags Ice-cream company wants to expand into the Irish market. They want to introduce all of the basic flavors of ice cream and one of the 'exotic' to the market flavors. Based on experience in Danish market they've decided to test two options: licorice and sea buckthorn. They recruited a sample of 100 true Irish inhabitants. Each of them was given sample of both flavors and graded each on the scale 1-10. How would you determine which of the flavors is more appropriate for the Irish market? Which test would you use? Are conditions fulfilled?

Nordisk Film Biografer is refurbishing their rooms. They have two chair types to choose from. The chairs have been tested by independent audience (12 hours LOTR extended marathon viewers). First chair received mean score of 7.8 (out of 10) with standard deviation of 1.3, the other 7.6 with standard deviation of 0.9. Both were tested by audiences of 50. Can you say with statistical certainty that one of them is better? Which test would you use to check this? How would you set up hypothesis? How would you calculate it?