# Parametric Significance Tests

#### Parametric Signifiance Tests

Experiment design	Two Conditions	Three <sup>(*)</sup> or more conditions
Between-group	t Test (independent samples)	One-way ANOVA
Within-group	<i>t</i> Test (paired samples)	Repeated Measures ANOVA

- Require normal distribution of the scores
- Variances should be nearly equal
- (\*) Can actually be applied also on two conditions

#### t Test

- Between-group design (independent samples)
  - Means are contributed from different groups
- Within-group design (paired samples)
  - Means are contributed from the same group

Can be two-tailed or one-tailed

 Simplified version of ANalysis Of VAriance (ANOVA) with only two groups or conditions

 H<sub>0</sub>: There is no significant difference in the task completion time between individuals who use the word-prediction software (WP group) and those who do not use the software (NP group).

Commonly accepted!

- Select reasonable significance level:  $\alpha = 5\%$
- If t-test shows significance at p < .05, we can conclude that under  $\alpha$  = 5%, i.e., in 95% of the time, the test result correctly applies to the entire population.

Sample data from between-group experiment

Group	Participants	Task completion time	Coding
No prediction	Participant 1	245	0
No prediction	Participant 2	236	O
No prediction	Participant 3	321	O
No prediction	Participant 4	212	0
No prediction	Participant 5	267	0
No prediction	Participant 6	334	0
No prediction	Participant 7	287	0
No prediction	Participant 8	259	O
With prediction	Participant 1	246	1
With prediction	Participant 2	213	1
With prediction	Participant 3	265	1
With prediction	Participant 4	189	1
With prediction	Participant 5	201	1
With prediction	Participant 6	197	1
With prediction	Participant 7	289	1
With prediction	Participant 8	224	1

#### **Testing for Normal Distribution**

- Kolmogorov-Smirnov Test
- Shapiro-Wilk Test (n ≤ 50)
  - -D'Agostino for n > 50
  - –Similar test Shapiro-Francia for n > 50

#### Shapiro-Wilk Test

(Conover, pp. 450)

- Assumption: sample is a random sample
- Compute test denominator D for 'No Prediction'

$$D = \sum_{i=1}^{n} (x_i - \overline{x})^2 = \sum_{i=1}^{8} (x_i - 270.125)^2 = 12260.875$$

Order 'No Prediction' samples

$$X = \{245, 236, 321, 212, 267, 334, 287, 259\}$$

from smallest to largest

•  $x^{(i)}$  denotes the *i*th order statistic, e.g.,  $x^{(4)} = 259$ 

#### Shapiro-Wilk Test

(Conover, pp. 450)

- Obtain coefficients  $a_1$ , ...,  $a_k$  with k = n/2 = 4  $a_1 = 0.6052$ ,  $a_2 = 0.3164$ ,  $a_3 = 0.1743$ ,  $a_4 = 0.0561$  (represent what the order statistics should look like if the population is normal)
- Compute test statistics

$$T = \frac{1}{D} \left[ \sum_{i=1}^{k} a_i \left( x^{(n-i+1)} - x^{(i)} \right) \right]^2$$

$$= \frac{1}{D} \left[ a_1 \left( x^{(8)} - x^{(1)} \right) + a_2 \left( x^{(7)} - x^{(2)} \right) + a_3 \left( x^{(6)} - x^{(3)} \right) + a_4 \left( x^{(5)} - x^{(4)} \right) \right]^2$$

$$= \dots$$

## Shapiro-Wilk Tables

n	2	- 6 3 5 6	6 4 6 6	8 55 8	6	8 7 8 1	8	9	10
	0.7071	0.7071	0.6872	0.6646	0.6431	0.6233	0.6052	0.0588	0.5739
2		0.0000	0.1667	0.2413	0.2806	0.3031	0.3164	0.3244	0.329
3	- 1			0.0000	0.0875	0.1401	0.1743	0.1976	0.214
4	· . —	_	onenos <u></u> -ones		- K	0.0000	0.0561	0.0947	0.1224
5			U. III.	<u> </u>		<del></del>		0.0000	0.0399

#### Shapiro-Wilk Test

(Conover, pp. 450)

$$T = \frac{1}{D} \left[ a_1 \left( x^{(8)} - x^{(1)} \right) + a_2 \left( x^{(7)} - x^{(2)} \right) + a_3 \left( x^{(6)} - x^{(3)} \right) + a_4 \left( x^{(5)} - x^{(4)} \right) \right]^2$$

$$= \frac{1}{D} \left[ a_1 \left( 334 - 212 \right) + a_2 \left( 321 - 236 \right) + \right]^2$$

$$= \frac{1}{D} \left[ 122 a_1 + 85 a_2 + 42 a_3 + 8a_4 \right]^2$$

$$= \frac{108.4978^2}{12260.875} \approx 0.96$$

- Sample: 212 < 236 < 245 < 259 < 267 < 287 < 321 < 334</li>
- Coefficients:  $a_1 = 0.6052$ ,  $a_2 = 0.3164$ ,  $a_3 = 0.1743$ ,  $a_4 = 0.0561$

#### Shapiro-Wilk Test

(Conover, pp. 450)

- Look up quantile of Shapiro-Wilk test for n = 8 and at α = .05, which is 0.818
- As  $T \approx 0.96 > 0.818$ , we accept  $H_0$  saying that the sample is normally distributed

(T close to 1.0, the sample behaves like normal sample, otherwise sample looks nonnormal)

Repeat for 'With Prediction' sample: T ≈ 0.92

## Shapiro-Wilk Tables

<b>n</b> :0.0	0.01	0.02	0.05	0.10	0.50	0.90	0.95	0.98	0.99
3	0.753	0.756	0.767	0.789	0.959	0.998	0.999	1.000	1.000
4	0.687	0.707	0.748	0.792	0.935	0.987	0.992	0.996	0.997
5	0.686	0.715	0.762	0.806	0.927	0.979	0.986	0.991	0.993
6	0.713	0.743	0.788	0.826	0.927	0.974	0.981	0.986	0.989
7	0.730	0.760	0.803	0.838	0.928	0.972	0.979	0.985	0.988
8	0.749	0.778	0.818	0.851	0.932	0.972	0.978	0.984	0.987
9	0.764	0.791	0.829	0.859	0.935	0.972	0.978	0.984	0.986
10	0.781	0.806	0.842	0.869	0.938	0.972	0.978	0.983	0.986
	0.700	0017	0.050			- · · · · · ·	0.770	0.703	0.700

#### Testing for Equality of Variances

Levene's test:

$$F_{Levene} = \frac{(N-t) \cdot \sum_{i=1}^{t} n_i (\overline{D}_i - \overline{D})^2}{(t-1) \cdot \sum_{i=1}^{t} \sum_{j=1}^{n_i} (D_{ij} - \overline{D}_i)^2}$$

t:number of treatments

 $n_i$ : number of observations from treatment i

 $N = n_1 + n_2 + ... + n_t$  (overall size of combined samples)

 $y_{ii}$ : observation j from treatment i  $(j = 1,...,n_i \text{ and } i = 1,...t)$ 

 $\overline{y}_i$ : mean of sample data from treatment i

 $D_{ij} = \left| \mathbf{y}_{ij} - \overline{\mathbf{y}}_{i} \right|$  (absolute deviation of observation j from treatment i mean)

 $\overline{D}_i$ : average of the n<sub>i</sub> absolute deviations from treatment i

 $\overline{D}$ : average of all N absolute deviations

#### Testing for Equality of Variances

Computing Levene's test score:

$$t = 2$$
;  $n_1 = n_2 = 8$ ;  $N = 16$ 

$$\begin{split} F_{Levene} &= \frac{(N-t) \cdot \sum_{i=1}^{t} n_{i} (\overline{D_{i}} - \overline{D})^{2}}{(t-1) \cdot \sum_{i=1}^{t} \sum_{j=1}^{n_{i}} (D_{ij} - \overline{D_{i}})^{2}} \\ &= \frac{14 \cdot 8 \cdot ((\overline{D_{1}} - \overline{D})^{2} + (\overline{D_{2}} - \overline{D})^{2})}{1 \cdot (\sum_{j=1}^{n_{1}} (D_{1j} - \overline{D_{1}})^{2} + \sum_{j=1}^{n_{2}} (D_{2j} - \overline{D_{2}})^{2})} \end{split}$$

$$\overline{D}_1 = 32.90625$$

$$\overline{D}_2 = 29$$

$$\overline{D} = 30.953125$$

#### Testing for Equality of Variances

Computing Levene's test score:

... = 
$$\frac{854.4921875}{\sum_{j=1}^{n_1} (D_{1j} - \overline{D}_1)^2 + \sum_{j=1}^{n_2} (D_{2j} - \overline{D}_2)^2}$$
$$= \frac{854.4921875}{3598.3046875 + 2138} \approx 0.149$$

- Compare with critical value of F-distribution<sup>1)</sup> for  $\alpha = .05$ :  $F_{Levene} \approx 0.149 < 4.600$
- We retain H<sub>0</sub> and conclude that the variances are equal

<sup>1)</sup> http://www.itl.nist.gov/div898/handbook/eda/section3/eda3673.htm

#### Critical Values of F-Distribution

				5% si	gnificance	level
				Ĭ	$F_{.05}(\nu_1, \nu_2)$	
	\ \ <sup>v</sup> 1 1	2	3	4	5	6
	<b>v</b> <sub>2</sub>					
1	161.448	199.500	215.707	224.583	230.162	233.986
2	18.513	19.000	19.164	19.247	19.296	19.330
3	10.128	9.552	9.277	9.117	9.013	8.941
4	7.709	6.944	6.591	6.388	6.256	6.163
5	6.608	5.786	5.409	5.192	5.050	4.950
6	5.987	5.143	4.757	4.534	4.387	4.284
7	5.591	4.737	4.347	4.120	3.972	3.866
8	5.318	4.459	4.066	3.838	3.687	3.581
9	5.117	4.256	3.863	3.633	3.482	3.374
16	4.965	4.103	3.708	3.478	3.326	3.217
11	4.844	3.982	3.587	3.357	3.204	3.095
12	4.747	3.885	3.490	3.259	3.106	2.996
13		3.806	3.411	3.179	3.025	2.915
14	4.600	3.739	3.344	3.112	2.958	2.848
15	4.543	3.682	3.287	3.056	2.901	2.790
16	4.494	3.634	3.239	3.007	2.852	2.741
17	4.451	3.592	3.197	2.965	2.810	2.699

- Central question: does the IV affect the DV?
- Recall
  - IV: Typing support (No prediction vs. With prediction)
  - DV: Task completion time

Group	Participants	Task completion time	Coding
No prediction	Participant 1	245	0
No prediction	Participant 2	236	0
No prediction	Participant 3	321	O
No prediction	Participant 4	212	0
No prediction	Participant 5	267	O
No prediction	Participant 6	334	0
No prediction	Participant 7	287	0
No prediction	Participant 8	259	O
With prediction	Participant 1	246	1
With prediction	Participant 2	213	1
With prediction	Participant 3	265	1
With prediction	Participant 4	189	1
With prediction	Participant 5	201	1
With prediction	Participant 6	197	1
With prediction	Participant 7	289	1
With prediction	Participant 8	224	1

- No prediction:
  - -M=270.125

- With prediction:
  - -M=228

- If the IV did not affect the DV, then the population means for the two groups are equal
- However, because of sampling error, the two means are not exactly equal even if the H<sub>∩</sub> is true
- For the example: not known whether the difference of 270.125 and 228 can be explained by sampling error
- Thus, question is: how large is the sampling error in  $\bar{X}_1 - \bar{X}_2$  given that the population means are equal?
- Aim: precisely estimate how unusual the difference between means is relative to its standard error:

$$\frac{\bar{X}_1 - \bar{X}_2}{SE(\bar{X}_1 - \bar{X}_2)}$$



- Standard error of single distribution:  $s \cdot \sqrt{\frac{1}{n}}$  with s is SD
- Standard error of the difference between two means randomly and independently sampled from the same population is:  $\sigma \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$

with  $\sigma$  is population SD and group sizes  $n_1$  and  $n_2$ 

- How to estimate the SD  $\sigma$ ?
- Pooled variance  $s_p^2$ : compute variance for each group and combine it by weighting based on their df

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

 Answer: precisely estimate how unusual the difference between means is relative to its standard error:

$$(*) \quad \frac{\bar{X}_1 - \bar{X}_2}{SE(\bar{X}_1 - \bar{X}_2)} = \frac{\bar{X}_1 - \bar{X}_2}{s_p \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}} = \frac{\bar{X}_1 - \bar{X}_2}{\sqrt{\frac{(n_1 - 1)s_1^2 + (n_2 - 1)s_2^2}{n_1 + n_2 - 2} \cdot \left[\frac{1}{n_1} + \frac{1}{n_2}\right]}}$$

- It happens that (\*) has a t distribution with  $n_1+n_2-2$  degrees of freedom, under the assumptions that
  - Residual term has normal distribution
  - Variances of  $s_1^2$  and  $s_1^2$  are equal
- Basically, this is the independent samples t-test

Determine degree of freedom:

$$d_f = df_1 + df_2$$

$$= (n_1 - 1) + (n_2 - 1)$$

$$= (8 - 1) + (8 - 1) = 14$$

 Look up critical t-value (α=.05, two-tailed test, df=14): t-critical = 2.1448

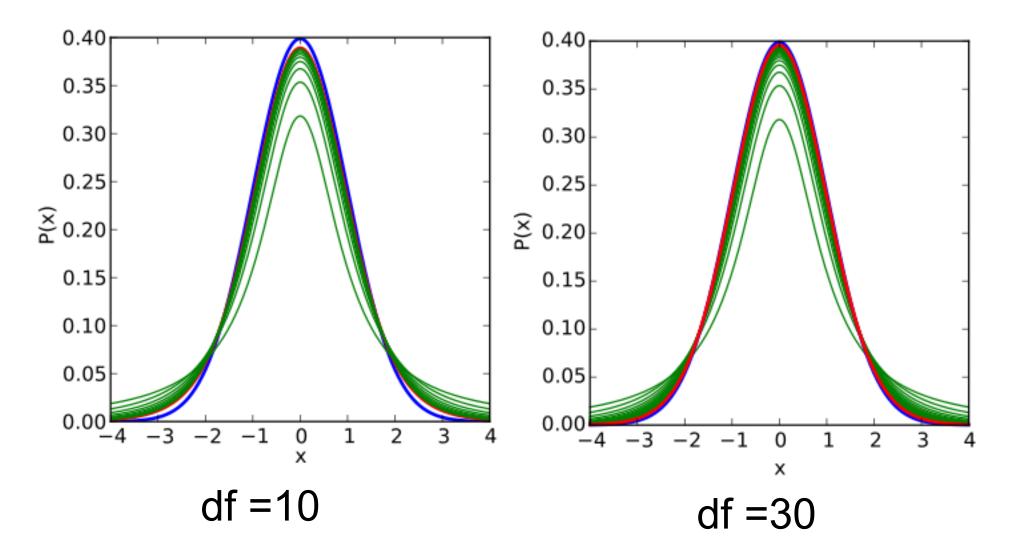
#### t Table

df

1-tailed	0.1	0.05	0.025
2-tailed	0.2	0.1	0.05
1	3.0777	6.3138	12.7062
2	1.8856	2.9200	4.3027
13	1.3502	1.7709	2.1604
14	1.3450	1.7613	2.1448

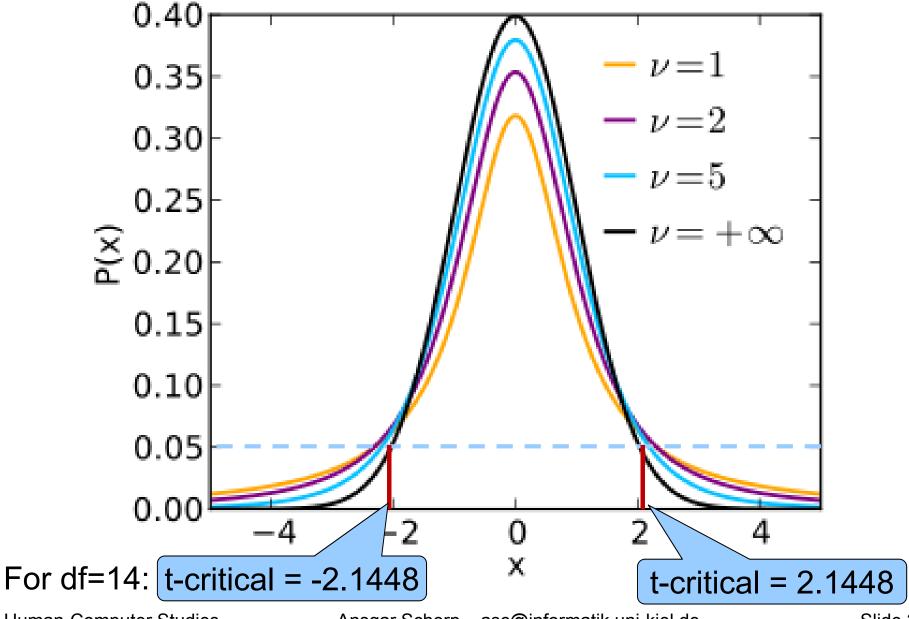
• Source: <a href="http://statisticslectures.com/tables/ttable/">http://statisticslectures.com/tables/ttable/</a>

#### Let's look at the t distribution



Source: Wikipedia

#### Let's look at the t distribution



Calculate pooled variance

Estimate the population variance

$$s_p^2 = \frac{s_1^2 df_1 + s_2^2 df_2}{df_1 + df_2}$$

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

$$= \frac{\sum_{i=1}^{n} (x_{1_i} - \overline{x}_1)^2 + \sum_{i=1}^{n} (x_{2_i} - \overline{x}_2)^2}{(n_1 - 1) + (n_2 - 1)} = \dots$$

Calculate pooled variance (continued)

$$s_p^2 = \dots = \frac{\sum_{i=1}^{n} (x_{1_i} - \overline{x}_1)^2 + \sum_{i=1}^{n} (x_{2_i} - \overline{x}_2)^2}{(n_1 - 1) + (n_2 - 1)}$$
$$= \frac{12260.875 + 8866}{(8 - 1) + (8 - 1)} = 1509.0625$$

$$\bar{x}_1 = 270.125, \bar{x}_2 = 228$$

Apply test statistics

$$t = \frac{\overline{x}_1 - \overline{x}_2}{S_p \sqrt{\frac{n_1 + n_2}{n_1 n_2}}} = \frac{\overline{x}_1 - \overline{x}_2}{\sqrt{\frac{s_p^2 n_1 + s_p^2 n_2}{n_1 n_2}}}$$

In textbooks, you may see various versions

$$= \frac{\overline{x}_{1} - \overline{x}_{2}}{\sqrt{\frac{s_{p}^{2}}{n_{1}} + \frac{s_{p}^{2}}{n_{2}}}} = \frac{\overline{x}_{1} - \overline{x}_{2}}{\sqrt{\frac{1}{n_{1}} + \frac{1}{n_{2}}}}$$

Normalization factor for pooled variance

$$= \frac{270.125 - 228}{\sqrt{\frac{1509.0625}{8} + \frac{1509.0625}{8}}} \approx \frac{42.125}{19.423} \approx 2.169$$

- Returned t-value t ≈ 2.169 > 2.1448
- Higher than t-value for specific df=14 at 95% confidence interval, which is 2.1448
  - → Reject H<sub>0</sub>
- Given the significance level  $\alpha = 5\%$ . There is a significant difference in the task completion time between 'No prediction' group and 'With prediction' group, t (14)=2.169, p < .05.

#### Please Note: p-value vs. a

• Given the significance level  $\alpha = 5\%$ . There is a significant difference in the task completion time between 'No prediction' group and 'With prediction' group, t (14)=2.169, p < .05.

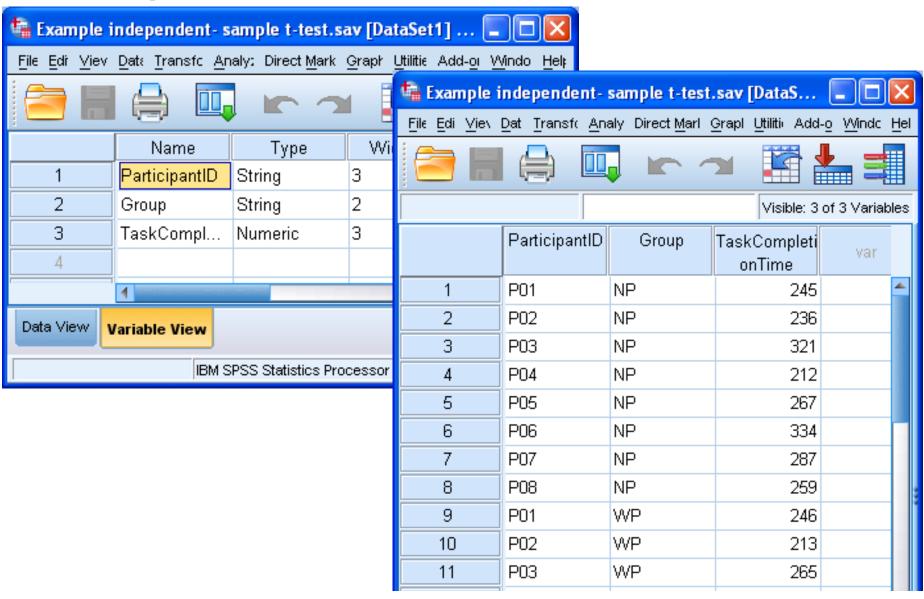
- p-value: probability of observing an effect given that H<sub>0</sub> is true (i.e., we actually should not observe that effect)
- Significance level α: probability of rejecting
   H<sub>0</sub> given that it is actually true (Type I error)

 Example computations with Gnumeric and SPSS 20 in Dropbox

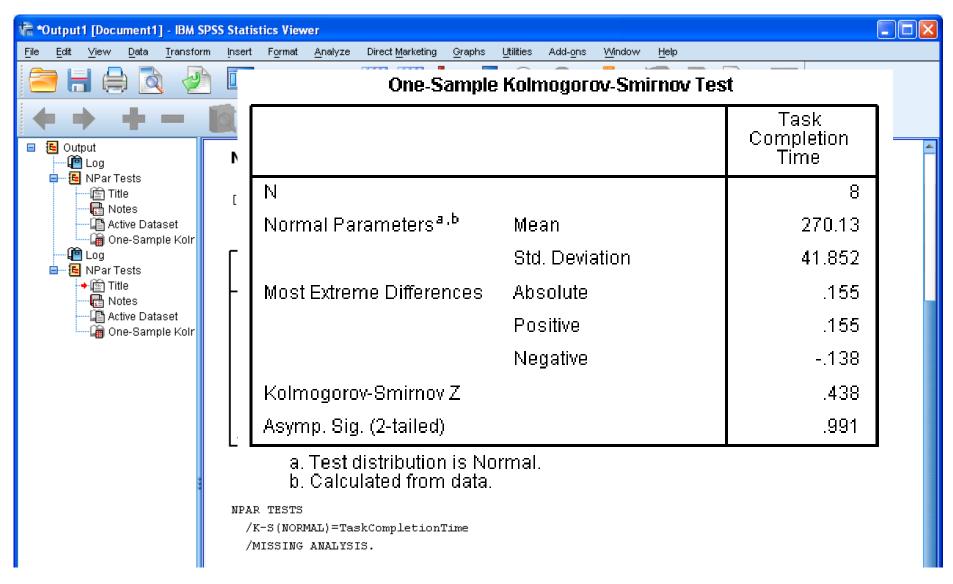
http://dl.dropbox.com/u/8830452/RMinHCI/DataAnalysis.zip

Also: computations with sample data from within-group design

#### Example in SPSS



## Example in SPSS (cont.)



## Example in SPSS (cont.)

#### 😬 🏓 NPar Tests

[DataSet1] X:\Dropbox\Public\RMinHCI\DataAnalysis\Example indepe

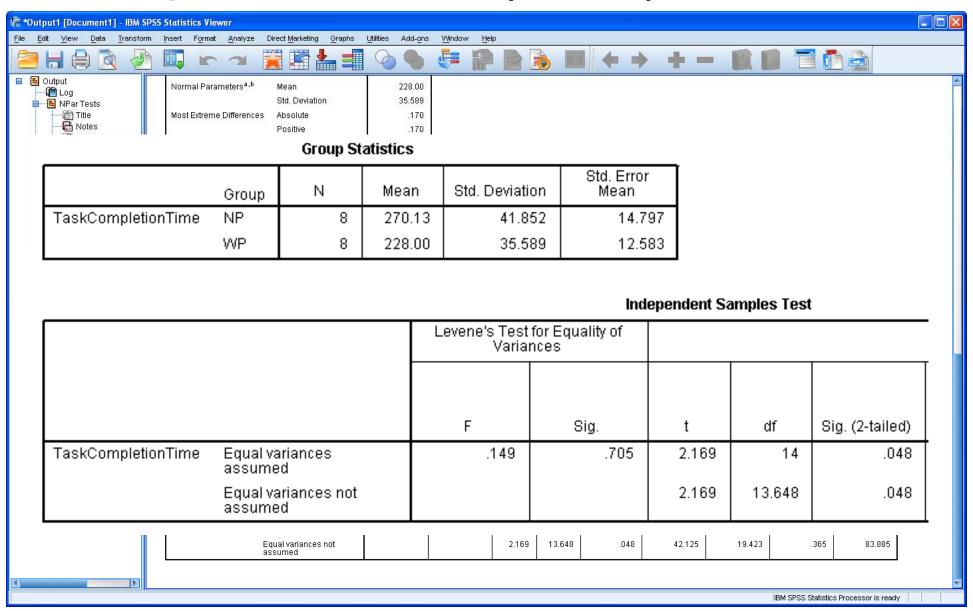
One-Sample Kolmogorov-Smirnov Test

		Task Completion Time
N		8
Normal Parameters <sup>a,b</sup>	Mean	228.00
	Std. Deviation	35.589
Most Extreme Differences	Absolute	.170
	Positive	.170
	Negative	137
Kolmogorov-Smirnov Z		.480
Asymp. Sig. (2-tailed)		.975

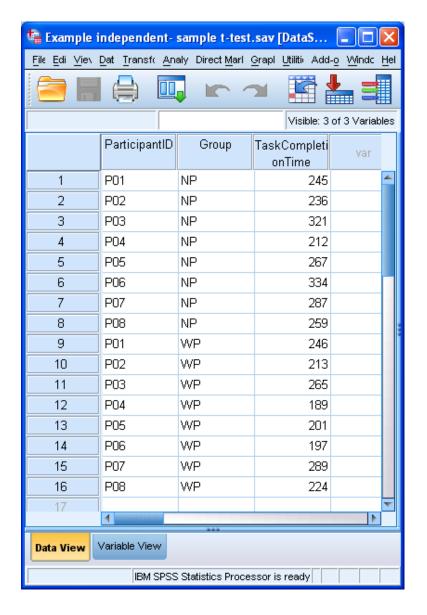
a. Test distribution is Normal.

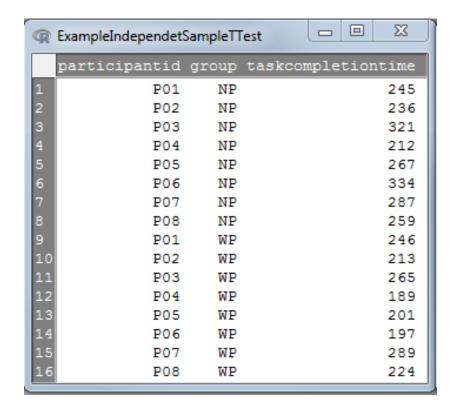
b. Calculated from data.

## Example in SPSS (cont.)

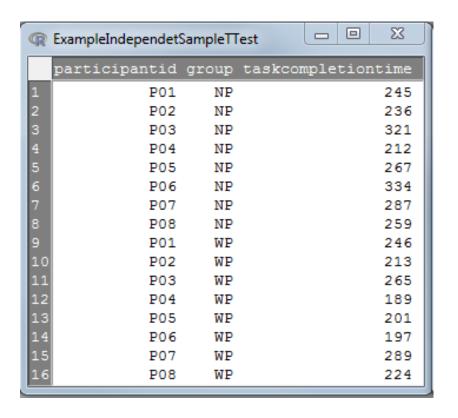


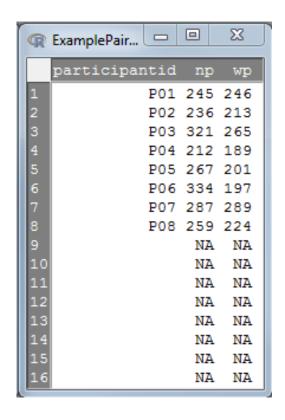
#### SPSS vs. R Commander (Rcmdr)





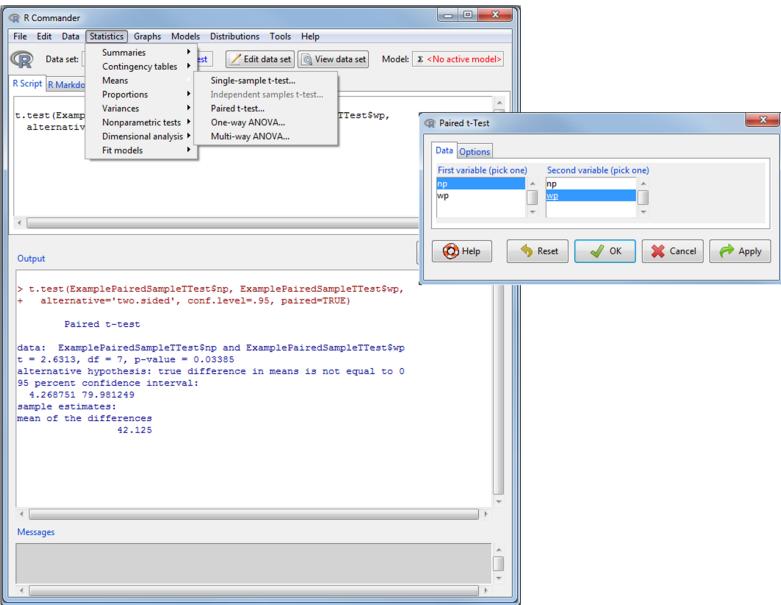
## Organizing Data in Rcmdr





Independent sample vs. paired sample

## Example in R Commander (Rcmdr)



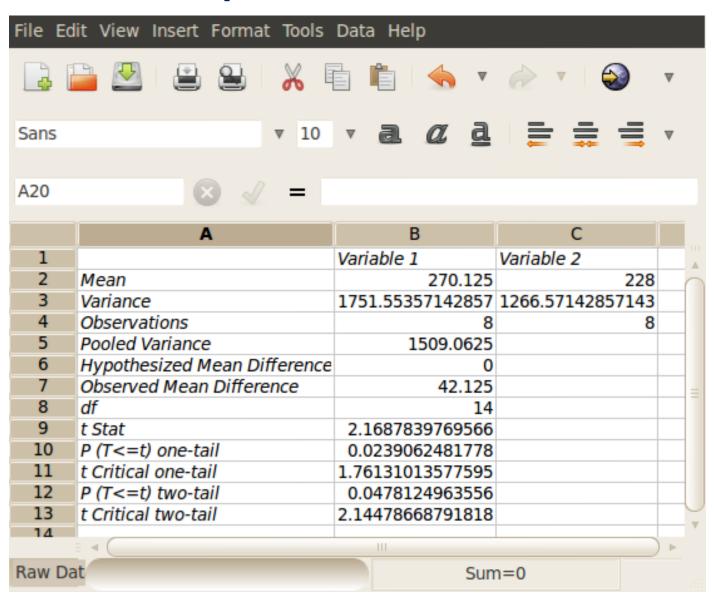
## t Test: Example of One-tailed Test

 H<sub>0</sub>: Individuals who use word-prediction software can type faster than those who do not use word-prediction software.

#### • Difference?

- Direction is specified in hypothesis
- Implying that word-prediction may improve typing speed or have no impact at all
- One-tailed t test is appropriate

### t Test: Example of One-tailed Test



## ANalysis Of VARiance (ANOVA)

- One-way ANOVA
  - One IV with two or more conditions
  - Returns a value F (thus, also called F-test)
- Factorial ANOVA
  - Like one-way ANOVA, but two or more IVs involved
- Both are for between-group designs

One-way/Factorial Repeated Measures ANOVA

#### One-way ANOVA: Example

Group	Participants	Task completion time	Coding
Standard	Participant 1	245	0
Standard	Participant 2	236	0
Standard	Participant 3	321	0
Standard	Participant 4	212	0
Standard	Participant 5	267	0
Standard	Participant 6	334	0
Standard	Participant 7	287	0
Standard	Participant 8	259	0
Prediction	Participant 1	246	1
Prediction	Participant 2	213	1
Prediction	Participant 3	265	1
Prediction	Participant 4	189	1
Prediction	Participant 5	201	1
Prediction	Participant 6	197	1
Prediction	Participant 7	289	1
Prediction	Participant 8	224	1
Speech-based dictation	Participant 1	178	2
Speech-based dictation	Participant 2	289	2
Speech-based dictation	Participant 3	222	2

- Three conditions
- No prediction (standard)
- With prediction
- Speech-based dictations

Source	Sum of squares	df	Mean square	$\boldsymbol{F}$	Significance
Between-group	7842.250	2	3921.125	2.174	0.139
Within-group	37880.375	21	1803.827		

#### One-way Repeated Measures ANOVA

	Standard	Prediction	Speech
Participant 1	245	246	178
Participant 2	236	213	289
Participant 3	321	265	222
Participant 4	212	189	189
Participant 5	267	201	245
Participant 6	334	197	311
Participant 7	287	289	267
Participant 8	259	224	197

Source	Sum of square	Df	Mean square	$\boldsymbol{F}$	Significance
Entry method	7842.25	2	3921.125	2.925	0.087
Error	18767.083	14	1340.506		

### One-way Repeated Measures ANOVA

- Significance does not mean that all means are actually different
- Conduct a pairwise t-tests
- Relative high risk of error due to multiple tests

- Reduce risk of error by using post-hoc test like Student-Newman-Keuls-Test
- Or apply Bonferroni correction, i.e., compare  $\alpha_{new} = \alpha/m$ , m being the number of conditions

### Split-plot ANOVA

- Involves between-group & within-group factors
- Example experiment design
  - Group 1 for task "Transcription"
  - Group 2 for task "Composition"
  - Both groups experience 3 conditions (K, P, S)

	Keyboard	Prediction	Speech
Transcription	Group 1	Group 1	Group 1
Composition	Group 2	Group 2	Group 2

 Could be considered two separate experiments with group 1 and group 2, respectively?

## Split-plot ANOVA data layout

Task type	Participant number	Task type coding	Standard	Prediction	Speech
Transcription	Participant 1	0	245	246	178
Transcription	Participant 2	0	236	213	289
Transcription	Participant 3	0	321	265	222
Transcription	Participant 4	0	212	189	189
Transcription	Participant 5	0	267	201	245
Transcription	Participant 6	0	334	197	311
Transcription	Participant 7	0	287	289	267
Transcription	Participant 8	0	259	224	197
Composition	Participant 9	1	256	265	189
Composition	Participant 10	1	269	232	321
Composition	Participant 11	1	333	254	202
Composition	Participant 12	1	246	199	198
Composition	Participant 13	1	259	194	278
Composition	Participant 14	1	357	221	341
Composition	Participant 15	1	301	302	279
Composition	Participant 16	1	278	243	229

#### Split-plot ANOVA summary report

Source	Sum of square	df	Mean square	F	Significance
Task type	2745.187	1	2745.187	0.995	0.335
Error	38625.125	14	2758.937		

Table 4.18 Results of the split-plot test for the between-group variable.

Source	Sum of square	df	Mean square	F	Significance
Entry method	17564.625	2	8782.313	5.702	0.008
Entry method * task type	114.875	2	57.437	0.037	0.963
Error (entry method)	43126.5	28	1540.232		

Table 4.19 Results of the split-plot test for the within-group variable.

### Summary: Parametric Methods

## Commonly used significance tests for comparing means and their application context

Experiment design	Independent variables (IV)	Conditions for each IV	Types of test
	1	2	Independent-samples t test
Between-group	1	3 or more	One-way ANOVA
	2 or more	2 or more	Factorial ANOVA
	1	2	Paired-samples t test
Within-group	1	3 or more	Repeated measures ANOVA
	2 or more	2 or more	Repeated measures ANOVA
Between- and within-group	2 or more	2 or more	Split-plot ANOVA

# Correlation, Effect Sizes, and Confidence Intervals

#### Statistical Power: Lessons Learned?

- One-tailed tests are more powerful in rejecting H<sub>0</sub> than two-tailed tests as they do not require such strong differences in the conditions
- Increasing the number of participants allows to find the smallest effect having statistical significance (=rejecting H<sub>0</sub> at "any costs")
- Thus, it is extremely important to report the effect size of the experiment result

Reality check!!!

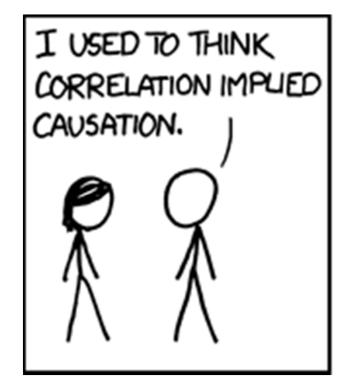
## Types of Effect Sizes

- Correlation family
  - Effect size based on "variance explained"
  - Example: Pearson's r
- Difference family
  - Effect size based on differences between means
  - -Example: Cohen's d
- And others, e.g., for categorical variables

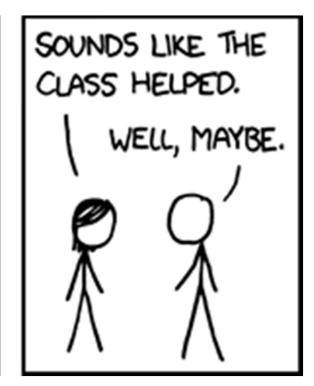
## **Correlation Analysis**

- Identify relationships between two factors
- Pearson's correlation coefficient (Pearson's r)
  - Linear relationships
  - -Perfect positive prediction when r = 1, no linear relationship when r = 0, perfect negative relationship when r = -1.
- Negative relationship equally good as positive
- Pearson's r as the most common effect size
- Caution: correlation does not imply causation

#### Got it?







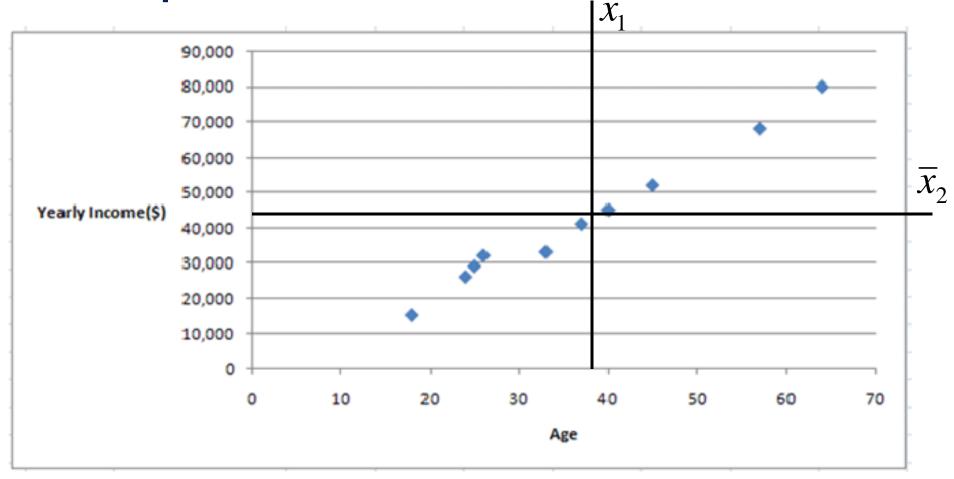
## Computing Pearson's r

Deviation score formula

$$r = \frac{\sum_{i=1}^{n} (x_{1_i} - \overline{x}_1)(x_{2_i} - \overline{x}_2)}{\sqrt{\sum_{i=1}^{n} (x_{1_i} - \overline{x}_1)^2} \sqrt{\sum_{i=1}^{n} (x_{2_i} - \overline{x}_2)^2}}$$

• The *i*-th score on the *j*-th treatment:  $x_{j_i}$ 

Example of Pearson's r



• Age and yearly income have a strong positive relationship (r(8) = .99, p < .05).

Source: http://www.statisticslectures.com/topics/pearsonr/

#### Effect Sizes in Pearson's r

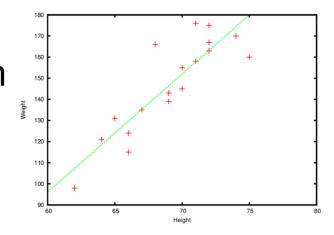
- Allow to compare different experiments in an objective way
  - -r = .10 (small effect): in this case the effect explains 1% of the total variance
  - -r = .30 (medium effect): explains 9% of total variance
  - -r = .50 (large effect): explains 25% of the variances

Spearman's r for non-linear relationships

#### **Excursion DMML: Goodness of Fit**

 Compare the regression sum of squares (SSM) with the total sum of squares (SST):

$$R^{2} = \frac{SSM}{SST} = \frac{\sum_{i=1...m} (f(x_{i}) - \bar{y})^{2}}{\sum_{i=1...m} (y_{i} - \bar{y})^{2}}$$



- Denotes how much variability can be explained with the regression model
- Note:  $R^2$  increases when more explanatory variables are added to the model  $\rightarrow$  use adjusted  $R^2$  measure

$$AdjustedR^2 = 1 - (\frac{m-1}{m-d})(1 - R^2)$$
 with

- m number of objects and
- d+1 number of parameters in the model

#### Example: Entity Resolution in mobEx

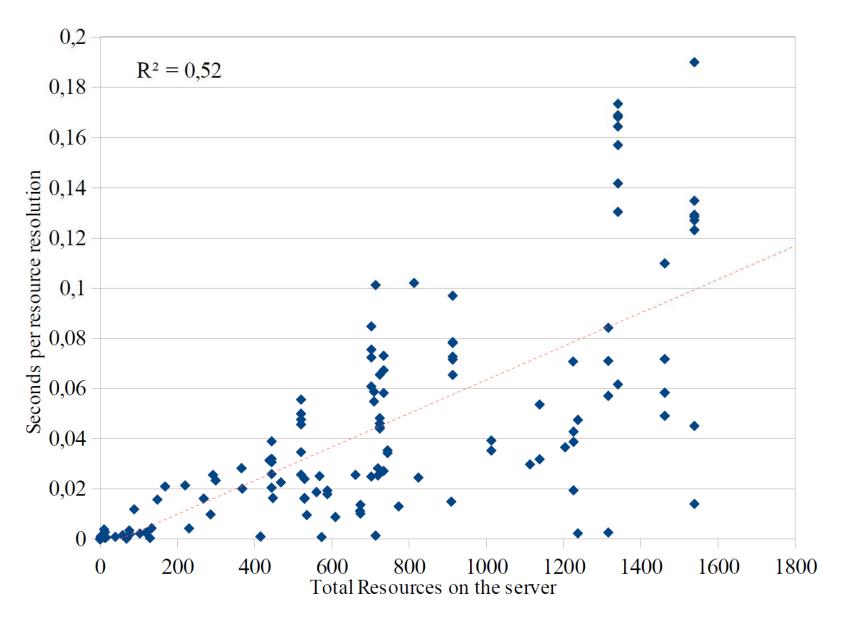






- Integration of nine data providers
- Query-time entity resolution
- Delivery of results once first merges are available

## Example: Runtime Performance



#### Difference between Means: Cohen's d

 Most common measure of how much the treatment affects the dependent variable

$$\delta = \frac{\mu_1 - \mu_2}{\sigma}$$
 (population standardized mean difference)

 Can be estimated from the means' estimators and the pooled variance

$$d = \frac{\bar{X}_1 - \bar{X}_2}{s_p} \qquad \text{with} \qquad t_{independent} = \frac{\bar{X}_1 - \bar{X}_2}{s_p \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}$$
 Note: the pooled variance disappears 
$$d = t_{independent} \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$$
 Results in: 
$$d = t_{independent} \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$$

#### Difference between Means: Cohen's d

Cohen's d for independent samples t test

$$d = t_{independent} \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$$
 - Small effect:  $d \ge .2$  - Medium effect:  $d \ge .5$ 

- Large effect:  $d \ge .8$

Cohen's d for dependent samples t test

$$d = t_{dependent} \cdot \sqrt{\frac{2(1-r)}{n}}$$

$$t_{dependent} = \frac{\bar{X}_D}{s_D \cdot \sqrt{1/n}}$$

where n is the number of pairs, r is correlation between paired scores

#### Difference between Means: Cohen's d

- Cohen's d can range from 0 to positive infinity
- Most values of d vary from 0 to 1
- Cohen's d is like a Z score in that its denominator is a standard deviation

## Confidence Intervals (CI)

- Compute an upper and lower limit of an interval [l, u]
  - Such that the probability that the *fixed* parameter (e.g., population mean) is contained in [l, u] is  $1 \alpha$
  - $-1-\alpha$  is the confidence level coverage
  - $\alpha$  is the Type I error rate
- Typically  $\alpha = 0.05$ , i.e., we compute 95% CIs (also written as:  $\text{CI}_{.95}$ )
- Cls can be computed for single means and differences between means (two distributions)

## CI on Mean (Single Distribution)

• Compute the 95% CI [l, u] given mean  $\bar{x}$ , standard deviation s and a constant c then

$$l = \bar{x} - c \cdot \frac{s}{\sqrt{n}}$$
 and  $u = \bar{x} + c \cdot \frac{s}{\sqrt{n}}$ 

- For large samples with  $n \ge 30$  (Central Limit Theorem)
  - Use standard normal distribution:  $c = z_{(1-\alpha/2)}$
  - Example for  $\alpha = 0.05$ :  $c = z_{0.975} = 1.96$
- For smaller samples with n < 30
  - Use approximation provided by Student's t-distribution at degree of freedom  $\nu = n-1$ :

$$c = z_{(1-\alpha/2;\nu)}$$

## CI on Mean (Single Distribution)

• More formally, a confidence interval is given as  $p[\theta_L(X) \le \theta \le \theta_U(X)] = 1 - \alpha$ 

#### where

- $\theta$  is some parameter of interest (mean, difference in mean, variance, etc.)
- $-\alpha = \alpha_L + \alpha_U$  (i.e., lower and upper can be different)
- $\theta_L(X)$  and  $\theta_U(X)$  the lower and upper random confidence limits of the observed data X (we call them random limits, since they are based on random data)

## CI on Mean (Single Distribution)

• Example for  $\mu$  over normally distributed data ( $n \ge 30$ ):

$$\begin{split} p\left[z_{(\alpha/_2)} \leq \frac{\bar{X} - \mu}{s_{\bar{X}}} \leq z_{(1 - \alpha/_2)}\right] &= 1 - \alpha \\ \Leftrightarrow p\left[z_{(\alpha/_2)} \cdot s_{\bar{X}} - \bar{X} \leq -\mu \leq z_{(1 - \alpha/_2)} \cdot s_{\bar{X}} - \bar{X}\right] &= 1 - \alpha \\ \Leftrightarrow p\left[\bar{X} - z_{(1 - \alpha/_2)} \cdot s_{\bar{X}} \leq \mu \leq \bar{X} - z_{(\alpha/_2)} \cdot s_{\bar{X}}\right] &= 1 - \alpha \\ \Leftrightarrow p\left[\bar{X} - z_{(1 - \alpha/_2)} \cdot s_{\bar{X}} \leq \mu \leq \bar{X} + z_{(1 - \alpha/_2)} \cdot s_{\bar{X}}\right] &= 1 - \alpha \\ \Leftrightarrow p\left[\bar{X} - z_{(1 - \alpha/_2)} \cdot s_{\bar{X}} \leq \mu \leq \bar{X} + z_{(1 - \alpha/_2)} \cdot s_{\bar{X}}\right] &= 1 - \alpha \\ -z_{(\alpha/_2)} = z_{(1 - \alpha/_2)} \text{ since the z-distribution is symmetric} \end{split}$$

 $\bar{X}$ : sample mean

 $\mu$ : population mean (parameter we seek to estimate for)

 $s_{\bar{X}}$ : population standard deviation of the sampling distribution of the mean (standard error)

 $\alpha/2 = \alpha_U = \alpha_L$ : significance level of computing the CI

#### CI on the Difference between Means

• Analog to the single distribution case, the equation in the two group situation with  $n_1, n_2 < 30$  is defined as:

$$p\left[t_{(\alpha_L;\,\nu)} \leq \frac{(\bar{X}_1 - \bar{X}_2) - (\mu_1 - \mu_2)}{s_p \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}} \leq t_{(1 - \alpha_U;\,\nu)}\right] = 1 - \alpha$$

$$\Leftrightarrow p \left| (\bar{X}_1 - \bar{X}_2) - t_{(1-\alpha_L; \nu)} \cdot s_p \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}} \le \mu_1 - \mu_2 \right|$$

$$\leq (\bar{X}_1 - \bar{X}_2) + t_{(1-\alpha_U; \nu)} \cdot s_p \cdot \sqrt{\frac{1}{n_1} + \frac{1}{n_2}} = 1 - \alpha$$

with  $v = n_1 + n_2 - 2$  degrees of freedom

 Further transformation needed to provide CI over standardized mean difference

#### Confidence Intervals

- A 95% CI can detect with a probability of 95% an interval that contains the true value of the parameter  $\theta$  (e.g., the mean). Thus, after computing the CI, it is a binary decision. Either the parameter value is included in the CI or not. It is not longer a matter of probability!
- Thus, common misunderstandings of CIs are:
  - A 95% CI does contain 95% of the sample data.
  - A 95% CI is a range of plausible values for the sample mean. → but plausible values for the CI parameters
  - A 95% CI contains with 95% probability the population mean / population parameter.
  - A particular 95% CI has a 95% probability that a sample mean is falling into the interval when one does repeat the experiment.

#### Continued on Part 3 ...