

SWE3053

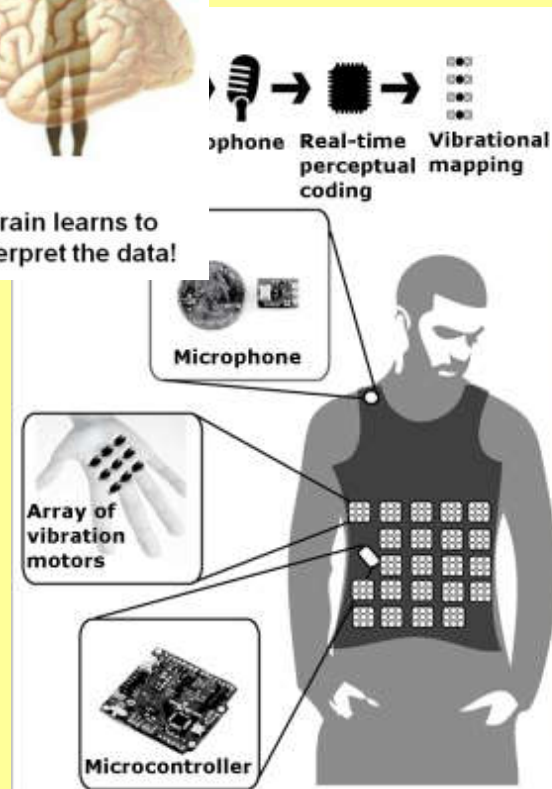
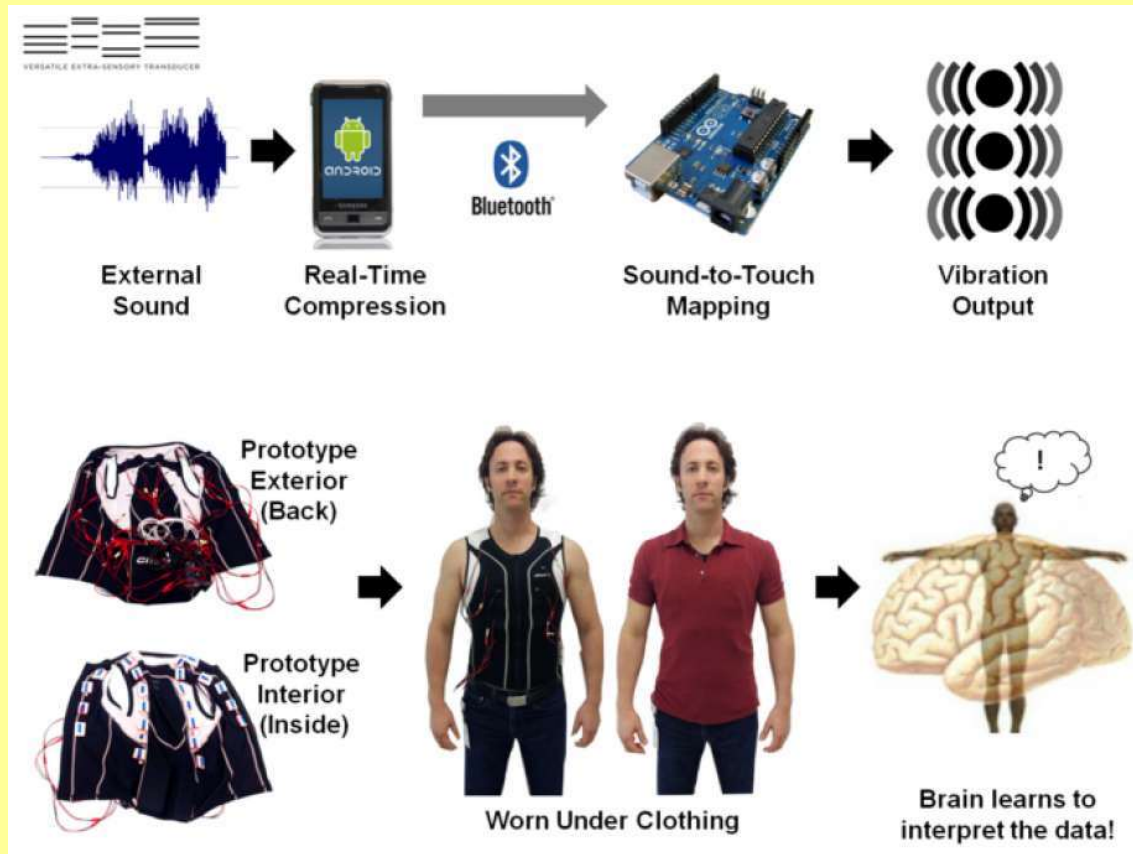
Human Computer Interaction

Lecture 21

Interpreting Data

Computer Aided Data Analysis

# Sensory Substitution – Extrasensation ...



# Agenda

- 
- Inferential Statistic
  - t-test
  - ANOVA

Refresh your memory .....

t-value

*difference between means*  
(SYSTEMATIC)

if "large enough" (e.g.,  
 $|t| > 1.96$ ) then you say  
difference is  
statistically significant

$$t = \frac{\bar{x} - \bar{y}}{S_{x\&y}}$$

*large values*  
*indicate statistical*  
*significance*

*variability of*  
*both distributions*  
(ERROR)

|    | Tail probability $p$ |       |       |       |       |       |       |       |       |       |       |       |
|----|----------------------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|-------|
| df | .25                  | .20   | .15   | .10   | .05   | .025  | .02   | .01   | .005  | .0025 | .001  | .0005 |
| 1  | 1.000                | 1.376 | 1.963 | 3.078 | 6.314 | 12.71 | 15.89 | 31.82 | 63.66 | 127.3 | 318.3 | 636.6 |
| 2  | .816                 | 1.061 | 1.386 | 1.886 | 2.920 | 4.303 | 4.849 | 6.965 | 9.925 | 14.09 | 22.33 | 31.60 |
| 3  | .765                 | .978  | 1.250 | 1.638 | 2.353 | 3.182 | 3.482 | 4.541 | 5.841 | 7.453 | 10.21 | 12.92 |
| 4  | .741                 | .941  | 1.190 | 1.533 | 2.132 | 2.776 | 2.999 | 3.747 | 4.604 | 5.598 | 7.173 | 8.610 |
| 5  | .727                 | .920  | 1.156 | 1.476 | 2.015 | 2.571 | 2.757 | 3.365 | 4.032 | 4.773 | 5.893 | 6.869 |
| 6  | .718                 | .906  | 1.134 | 1.440 | 1.943 | 2.447 | 2.612 | 3.143 | 3.707 | 4.317 | 5.208 | 5.959 |
| 7  | .711                 | .896  | 1.119 | 1.415 | 1.895 | 2.365 | 2.517 | 2.998 | 3.499 | 4.029 | 4.785 | 5.408 |
| 8  | .706                 | .889  | 1.108 | 1.397 | 1.860 | 2.306 | 2.449 | 2.896 | 3.355 | 3.833 | 4.501 | 5.041 |
| 9  | .703                 | .883  | 1.100 | 1.383 | 1.833 | 2.262 | 2.398 | 2.821 | 3.250 | 3.690 | 4.297 | 4.781 |
| 10 | .700                 | .879  | 1.093 | 1.372 | 1.812 | 2.228 | 2.359 | 2.764 | 3.169 | 3.581 | 4.144 | 4.587 |
| 11 | .697                 | .876  | 1.088 | 1.363 | 1.796 | 2.201 | 2.328 | 2.718 | 3.106 | 3.497 | 4.025 | 4.437 |
| 12 | .695                 | .873  | 1.083 | 1.356 | 1.782 | 2.179 | 2.303 | 2.681 | 3.055 | 3.428 | 3.930 | 4.318 |
| 13 | .694                 | .870  | 1.079 | 1.350 | 1.771 | 2.160 | 2.282 | 2.650 | 3.012 | 3.372 | 3.852 | 4.221 |
| 14 | .692                 | .868  | 1.076 | 1.345 | 1.761 | 2.145 | 2.264 | 2.624 | 2.977 | 3.326 | 3.787 | 4.140 |
| 15 | .691                 | .866  | 1.074 | 1.341 | 1.753 | 2.131 | 2.249 | 2.602 | 2.947 | 3.286 | 3.733 | 4.073 |
| 16 | .690                 | .865  | 1.071 | 1.337 | 1.746 | 2.120 | 2.235 | 2.583 | 2.921 | 3.252 | 3.686 | 4.015 |
| 17 | .689                 | .863  | 1.069 | 1.333 | 1.740 | 2.110 | 2.224 | 2.567 | 2.898 | 3.222 | 3.646 | 3.965 |
| 18 | .688                 | .862  | 1.067 | 1.330 | 1.734 | 2.101 | 2.214 | 2.552 | 2.878 | 3.197 | 3.611 | 3.922 |
| 19 | .688                 | .861  | 1.066 | 1.328 | 1.729 | 2.093 | 2.205 | 2.539 | 2.861 | 3.174 | 3.579 | 3.883 |
| 20 | .687                 | .860  | 1.064 | 1.325 | 1.725 | 2.086 | 2.197 | 2.528 | 2.845 | 3.153 | 3.552 | 3.850 |
| 21 | .686                 | .859  | 1.063 | 1.323 | 1.721 | 2.080 | 2.189 | 2.518 | 2.831 | 3.135 | 3.527 | 3.819 |
| 22 | .686                 | .858  | 1.061 | 1.321 | 1.717 | 2.074 | 2.183 | 2.508 | 2.819 | 3.119 | 3.505 | 3.792 |
| 23 | .685                 | .858  | 1.060 | 1.319 | 1.714 | 2.069 | 2.177 | 2.500 | 2.807 | 3.104 | 3.485 | 3.768 |
| 24 | .685                 | .857  | 1.059 | 1.318 | 1.711 | 2.064 | 2.172 | 2.492 | 2.797 | 3.091 | 3.467 | 3.745 |
| 25 | .684                 | .856  | 1.058 | 1.316 | 1.708 | 2.060 | 2.167 | 2.485 | 2.787 | 3.078 | 3.450 | 3.725 |
| 26 | .684                 | .856  | 1.058 | 1.315 | 1.706 | 2.056 | 2.162 | 2.479 | 2.779 | 3.067 | 3.435 | 3.707 |
| 27 | .684                 | .855  | 1.057 | 1.314 | 1.703 | 2.052 | 2.158 | 2.473 | 2.771 | 3.057 | 3.421 | 3.690 |
| 28 | .683                 | .855  | 1.056 | 1.313 | 1.701 | 2.048 | 2.154 | 2.467 | 2.763 | 3.047 | 3.408 | 3.674 |
| 29 | .683                 | .854  | 1.055 | 1.311 | 1.699 | 2.045 | 2.150 | 2.462 | 2.756 | 3.038 | 3.396 | 3.659 |
| 30 | .683                 | .854  | 1.055 | 1.310 | 1.697 | 2.042 | 2.147 | 2.457 | 2.750 | 3.030 | 3.385 | 3.646 |
| 40 | .681                 | .     |       |       |       |       |       |       |       |       |       |       |

# •The $t$ Distribution

We use  $t$  when

1. the population variance is unknown (the usual case), and
2. sample size is small ( $N < 100$ , the usual case).

The  $t$  distribution is a short, fat relative of the normal.

The shape of  $t$  depends on its  $df$  ( $df = N - 1$ ). As  $N$  becomes infinitely large,  $t$  becomes normal.

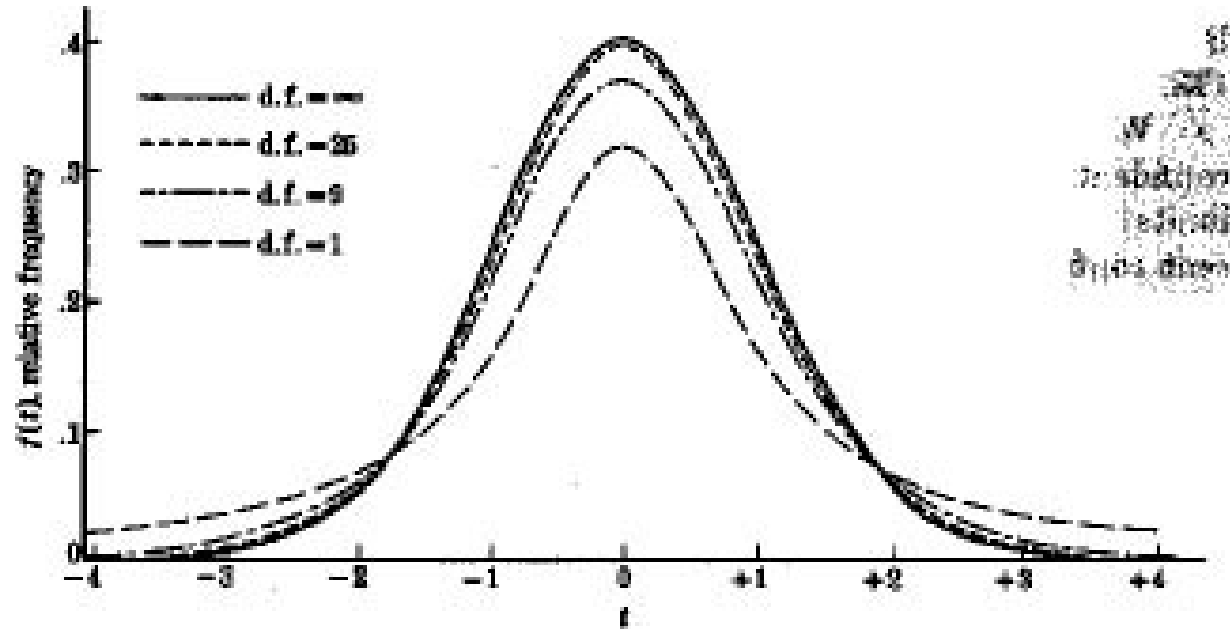
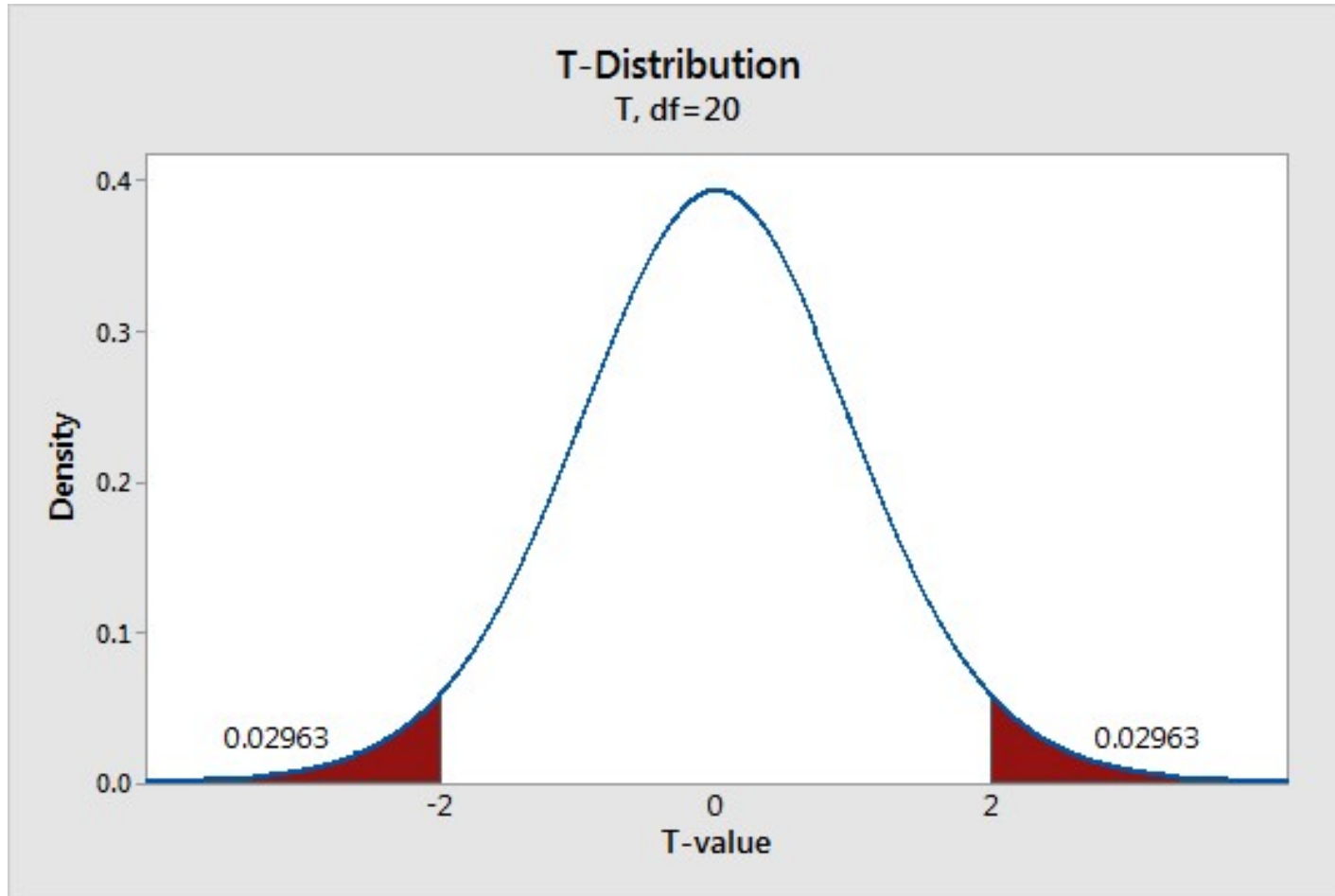


fig. 10.1 Distribution of  $t$  for various degrees of freedom. (From D. Lewis, *quantitative methods in psychology*, McGraw-Hill Book Company, New York, 1980.)

# The t-distribution

What is the meaning when you have a t-value of 2?



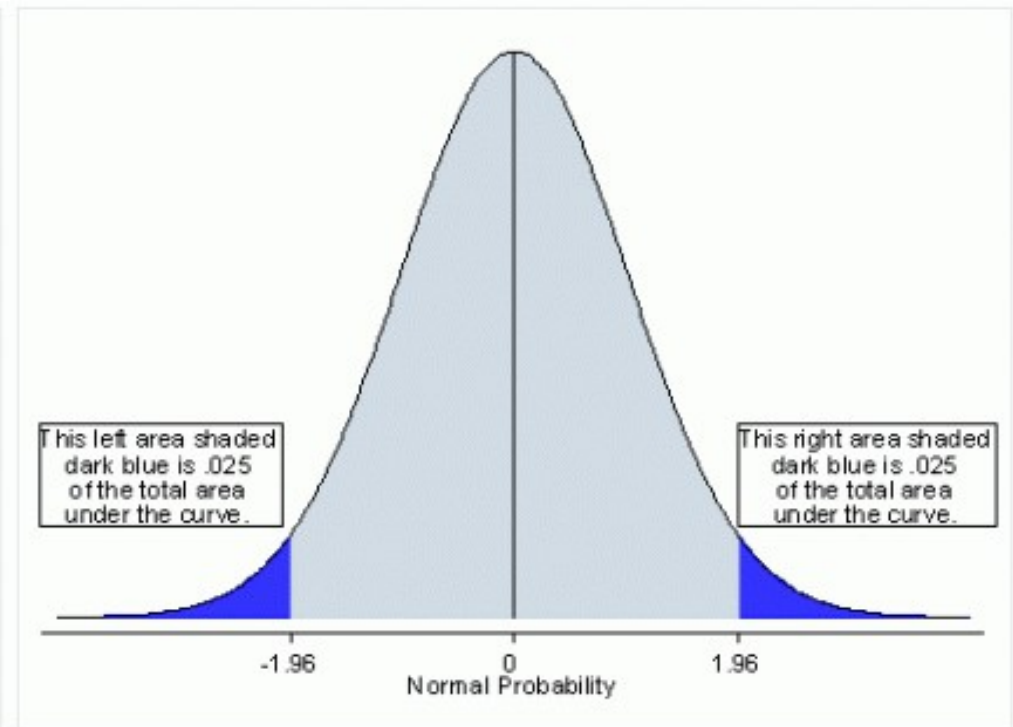
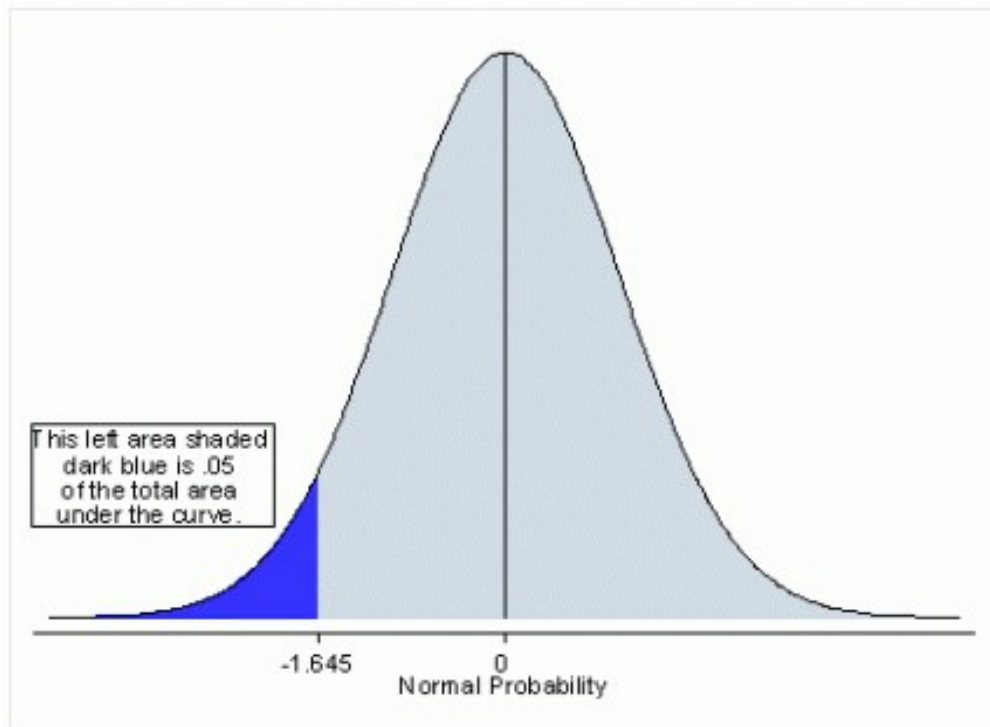
The probability for observing a difference from the null hypothesis that is at least as extreme as the difference present in our sample data while assuming that the null hypothesis is actually true – 5.962%.

This is called the p-value!



# 1-tailed vs. 2-tailed

- The t-value to achieve a significant level of 0.05 is:
  - 1-tailed – 1.645
  - 2-tailed – 1.96



## 2-tailed

- When you are comparing if the 2 mean values are the same.

## 1-tailed

- When you are comparing if one of the mean value significantly bigger/smaller than the other.



## •What kind of $t$ is it?

- 1)Single sample  $t$ -test – we have only 1 group; want to test against a hypothetical mean.
- 2)Independent samples  $t$ -test – we have 2 means, 2 groups; no relation between groups, e.g., people randomly assigned to a single group.
- 3)Dependent samples  $t$ -test – we have two means. Either same people in both groups, or people are related, e.g., husband-wife, left hand-right hand, hospital patient and visitor.

# 1. Single Sample t-test

- Used for comparing a sample mean to a population mean to determine if they are statistically significantly different
- Sample mean
  - The mean value of your sample
  - i.e. the mean of your data!
- Population mean
  - The real mean value

1. You know the population mean
2. You ran an experiment and obtained a sample mean from your samples
3. You want to know if the mean of your sample is significantly different from the known population mean

# 1. Single Sample t-test - Example

- You invented a drug to increase people's IQ!
- You know the population mean IQ score is 100.
- You recruited 20 people, give them the drug, and ask them to take the IQ test.
- And here's your sample data:

|     |     |
|-----|-----|
| 107 | 117 |
| 122 | 103 |
| 113 | 99  |
| 96  | 80  |
| 101 | 103 |
| 92  | 130 |
| 115 | 74  |
| 81  | 132 |
| 128 | 98  |
| 128 | 130 |

Mean = 107.45

STDEV = 17.83

df = N-1 = 19

# 1. Single Sample t-test - Example

- The t-value formula for single sample t-test:

$$t = \frac{\bar{x} - \mu_0}{s / \sqrt{n}}$$

|     |     |
|-----|-----|
| 107 | 117 |
| 122 | 103 |
| 113 | 99  |
| 96  | 80  |
| 101 | 103 |
| 92  | 130 |
| 115 | 74  |
| 81  | 132 |
| 128 | 98  |
| 128 | 130 |

Mean = 107.45

STDEV = 17.83

df = 19

$$\begin{aligned} t &= (107.45 - 100) / (17.83 / (20)^{1/2}) \\ &= 7.45 / 3.99 \\ &= 1.87 \end{aligned}$$

# 1. Single Sample t-test - Example

- Look up the t-table .....
- 1-tailed table

<http://www.statisticshowto.com/tables/t-distribution-table/>

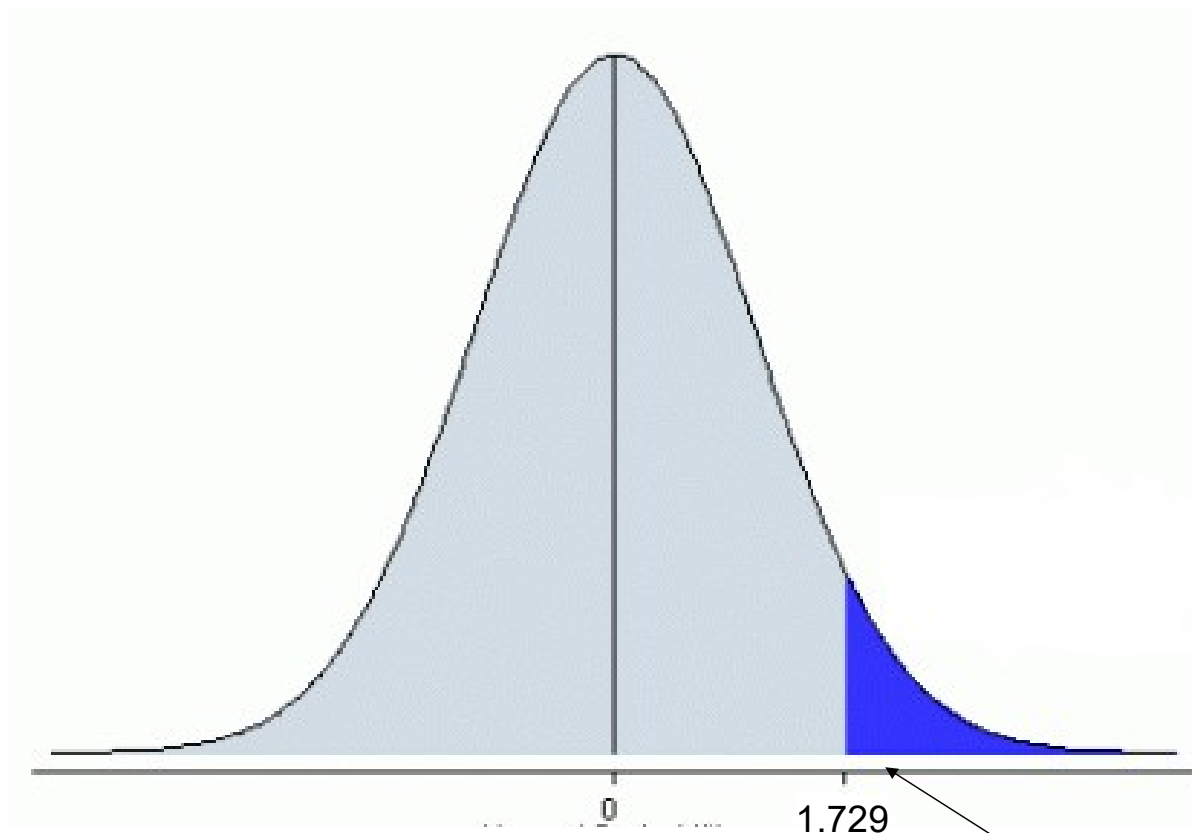
|     |     |
|-----|-----|
| 107 | 117 |
| 122 | 103 |
| 113 | 99  |
| 96  | 80  |
| 101 | 103 |
| 92  | 130 |
| 115 | 74  |
| 81  | 132 |
| 128 | 98  |
| 128 | 130 |

Mean = 108.45  
STDEV = 19.63  
t = 1.87

t value for achieving sig. level of 5% is 1.729

| DF | A = 0.1                | 0.05  | 0.025  | 0.01   | 0.005  | 0.001   | 0.0005  |
|----|------------------------|-------|--------|--------|--------|---------|---------|
| ∞  | t <sub>α</sub> = 1.282 | 1.645 | 1.960  | 2.326  | 2.576  | 3.091   | 3.291   |
| 1  | 3.078                  | 6.314 | 12.706 | 31.821 | 63.656 | 318.289 | 636.578 |
| 2  | 1.886                  | 2.920 | 4.303  | 6.965  | 9.925  | 22.328  | 31.600  |
| 3  | 1.638                  | 2.353 | 3.182  | 4.541  | 5.841  | 10.214  | 12.924  |
| 4  | 1.533                  | 2.132 | 2.776  | 3.747  | 4.604  | 7.173   | 8.610   |
| 5  | 1.476                  | 2.015 | 2.571  | 3.365  | 4.032  | 5.894   | 6.869   |
| 6  | 1.440                  | 1.943 | 2.447  | 3.143  | 3.707  | 5.208   | 5.959   |
| 7  | 1.415                  | 1.895 | 2.365  | 2.998  | 3.499  | 4.785   | 5.408   |
| 8  | 1.397                  | 1.860 | 2.306  | 2.896  | 3.355  | 4.501   | 5.041   |
| 9  | 1.383                  | 1.833 | 2.262  | 2.821  | 3.250  | 4.297   | 4.781   |
| 10 | 1.372                  | 1.812 | 2.228  | 2.764  | 3.169  | 4.144   | 4.587   |
| 11 | 1.363                  | 1.796 | 2.201  | 2.718  | 3.106  | 4.025   | 4.437   |
| 12 | 1.356                  | 1.782 | 2.179  | 2.681  | 3.055  | 3.930   | 4.318   |
| 13 | 1.350                  | 1.771 | 2.160  | 2.650  | 3.012  | 3.852   | 4.221   |
| 14 | 1.345                  | 1.761 | 2.145  | 2.624  | 2.977  | 3.787   | 4.140   |
| 15 | 1.341                  | 1.753 | 2.131  | 2.602  | 2.947  | 3.733   | 4.073   |
| 16 | 1.337                  | 1.746 | 2.120  | 2.583  | 2.921  | 3.686   | 4.015   |
| 17 | 1.333                  | 1.740 | 2.110  | 2.567  | 2.898  | 3.646   | 3.965   |
| 18 | 1.330                  | 1.734 | 2.101  | 2.552  | 2.878  | 3.610   | 3.922   |
| 19 | 1.328                  | 1.729 | 2.093  | 2.539  | 2.861  | 3.579   | 3.883   |
| 20 | 1.325                  | 1.725 | 2.086  | 2.528  | 2.845  | 3.552   | 3.850   |
| 21 | 1.323                  | 1.721 | 2.080  | 2.518  | 2.831  | 3.527   | 3.819   |
| 22 | 1.321                  | 1.717 | 2.074  | 2.508  | 2.819  | 3.505   | 3.792   |

df = 19; 1-tailed



t-value calculated from your sample is 1.87

$t(19) = 1.87, p < .05$

=> The effect is significant.

=> The Null Hypothesis is rejected.



# 5 steps for hypothesis proving

1. State the Null Hypothesis  $H_0$  and Alternative Hypothesis  $H_1$ .

$H_0$ : sample mean  $\leq 100$

$H_1$ : sample mean  $> 100$

2. Select the appropriate statistical method.

One sample t-test, 1-tailed

3. Select a level of significance.

.05;  $df = 19$ ; critical t-value needs to be larger than 1.729

4. Calculate the statistic.

$t(19) = 1.87$ ;  $p < .05$

5. Make the decision.

$H_0$  is reject;  $H_1$  is supported.

The effect of the drug is statistically significant



# How to do it in PSPP

\*[DataSet1] — PSPPIRE Data Editor

File Edit View Data Transform **Analyze** Graphs Utilities Windows Help

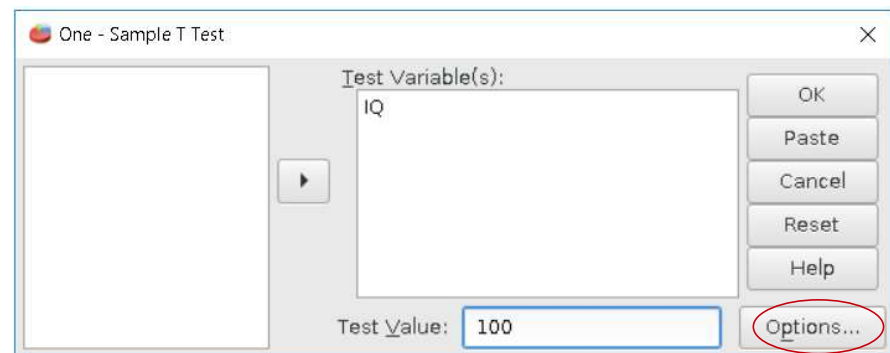
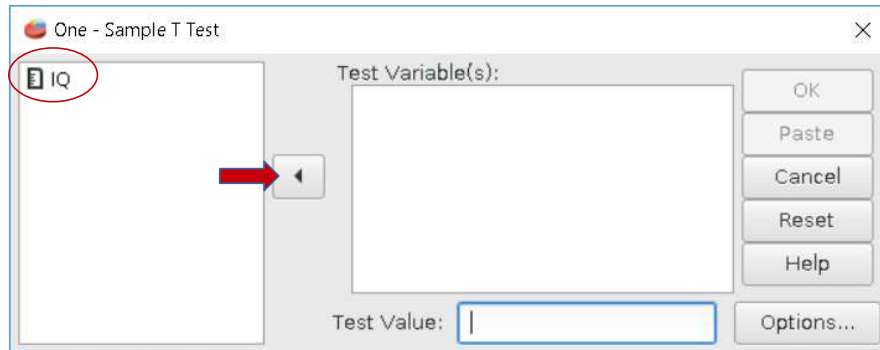
Descriptive Statistics  
Compare Means  
Univariate Analysis...  
Bivariate Correlation...  
K-Means Cluster...  
Factor Analysis...  
Reliability...  
Regression  
Non-Parametric Statistics  
ROC Curve...

Means...  
One Sample T Test...  
Independent Samples T Test...  
Paired Samples T Test...  
One Way ANOVA...

| Case | IQ     |  |
|------|--------|--|
| 1    | 107.00 |  |
| 2    | 122.00 |  |
| 3    | 113.00 |  |
| 4    | 96.00  |  |
| 5    | 101.00 |  |
| 6    | 92.00  |  |
| 7    | 115.00 |  |
| 8    | 81.00  |  |
| 9    | 128.00 |  |
| 10   | 128.00 |  |
| 11   | 117.00 |  |
| 12   | 103.00 |  |
| 13   | 99.00  |  |
| 14   | 80.00  |  |
| 15   | 103.00 |  |
| 16   | 130.00 |  |
| 17   | 74.00  |  |
| 18   | 132.00 |  |
| 19   | 98.00  |  |
| 20   | 130.00 |  |
| 21   | .      |  |
| 22   |        |  |

Data View Variable View

# How to do it in PSPP

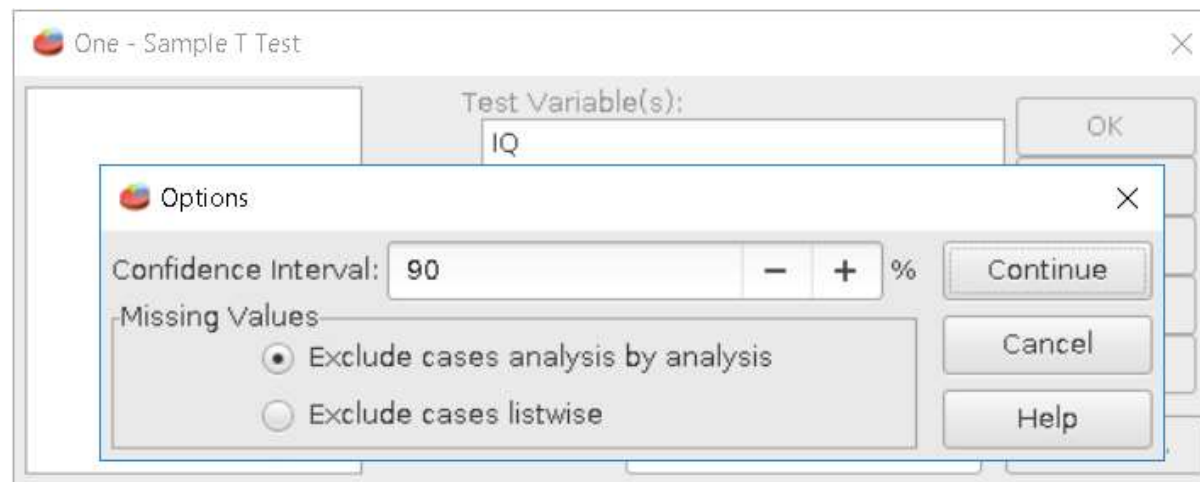


# How to do it in PSPP

Note: PSPP (and SPSS) does not have the option to do 1-tailed t-test; so you have to “hack” it...

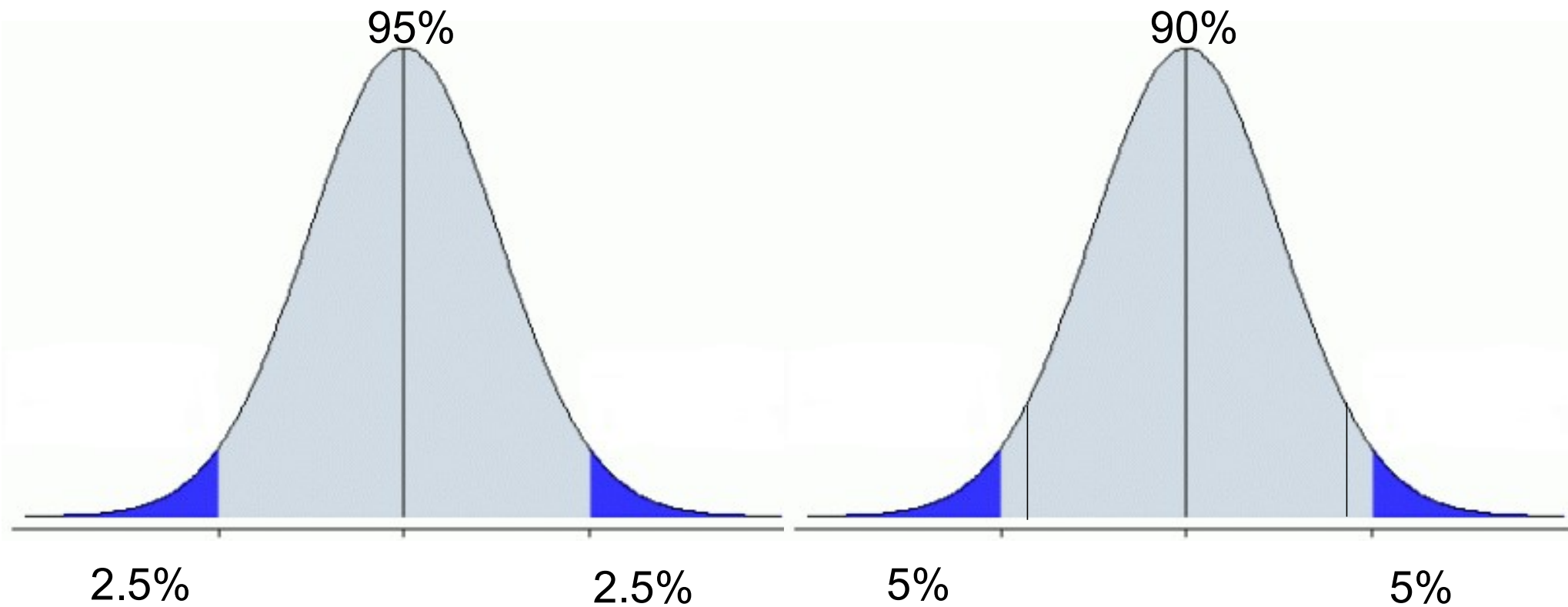
Your level of significant is 95%.

To do a 1-tailed test, change the significant level to 90%



# Explanation ...

- If you ignore the left hand side of the curve, 90% significant of 2-tailed is the same as 95% of 1-tailed



# Result

- Note the significant value reported from PSPP is for 2-tailed test
- To get the correct value for 1-tailed test, divide the value by 2
  - i.e. actual significant value should be **.0385**

```
T-TEST /TESTVAL=100
      /VARIABLES= IQ      /MISSING=ANALYSIS
      /CRITERIA=CI(0.9).
```

## One-Sample Statistics

|    | <i>N</i> | <i>Mean</i> | <i>Std. Deviation</i> | <i>S.E. Mean</i> |
|----|----------|-------------|-----------------------|------------------|
| IQ | 20       | 107.45      | 17.83                 | 3.99             |

Descriptive Statistic

## One-Sample Test

| Test Value = 100.000000 |          |           |                        |                        |   |              |
|-------------------------|----------|-----------|------------------------|------------------------|---|--------------|
|                         | <i>t</i> | <i>df</i> | <i>Sig. (2-tailed)</i> | <i>Mean Difference</i> | 90% Confidence Interval of the Difference |              |
|                         |          |           |                        |                        | <i>Lower</i>                              | <i>Upper</i> |
| IQ                      | 1.87     | 19        | .077                   | 7.45                   | .56                                       | 14.34        |

t-value



Significant value: p-value

# What is the meaning of the significant value $p$ ?

- A  $p$  value of .0385 means:
    - If you repeat the experiment 100 times, the Null Hypothesis  $H_0$  will be correct for 3.85 times.
    - i.e. the Alternative Hypothesis  $H_1$  will be correct 96.15 times.
- $H_0$ : sample mean  $\leq 100$   
 $H_1$ : sample mean  $> 100$
- If you recruit 2000 subjects, split them into 100 groups, with 20 subjects per group.
  - Then you give them drugs.
  - Then you ask them to do the IQ test.
  - Out of the 100 groups:
    - 96.15 groups will have mean score of  $>100$
    - 3.85 groups will have mean score of  $\leq 100$





# How do you compare means of 2 conditions?

- How do you analysis data in our cellphone driving experiment?

Independent samples *t-test* – we have 2 means, 2 groups; no relation between groups, e.g., people randomly assigned to a single group.

## 2. Independent Sample t-test

- Used for comparing the mean of one group to the mean of another independent group to determine if they are statistically significantly different
- Used for between-subject design comparison
- Requirement:
  - DV should be independence: Scores in one sample do not influence scores in other samples
  - Must be 2 groups (i.e. 2 conditions)
  - Measurement should be scaled
    - i.e. not categorical (e.g. male vs. female)
  - Homogenous assumption
    - Levene Test should not reach significance

## 2. Independent Sample t-test

$$t = \frac{\bar{X}_1 - \bar{X}_2}{\sqrt{\left[ \frac{(N_1 - 1)s_1^2 + (N_2 - 1)s_2^2}{N_1 + N_2 - 2} \right] \left[ \frac{1}{N_1} + \frac{1}{N_2} \right]}}$$


A blue arrow points from the label 'A' to the numerator  $\bar{X}_1 - \bar{X}_2$ . A green arrow points from the label 'B' to the square root symbol. A red arrow points from the label 'C' to the term  $\left[ \frac{1}{N_1} + \frac{1}{N_2} \right]$ .

- 
- $\bar{X}_1$  = mean of group 1;  $\bar{X}_2$  = mean of group 2;
- $N_1$  = number of sample of group 1
- $N_2$  = number of sample of group 2
- $S_1$  = Standard deviation of group 1
- $S_2$  = Standard deviation of group 2



A cartoon illustration of a man wearing a hat and a suit, driving a vintage car. He is smiling and holding the steering wheel. The car is moving to the right, indicated by motion lines behind it.

A cartoon illustration of a man and a woman driving a vintage car. The man is wearing a hat and a suit, and the woman is wearing a hat and a dress. They are both smiling. The car is a classic convertible with a dark body and light-colored wheels. There are motion lines behind the car, suggesting it is moving quickly.



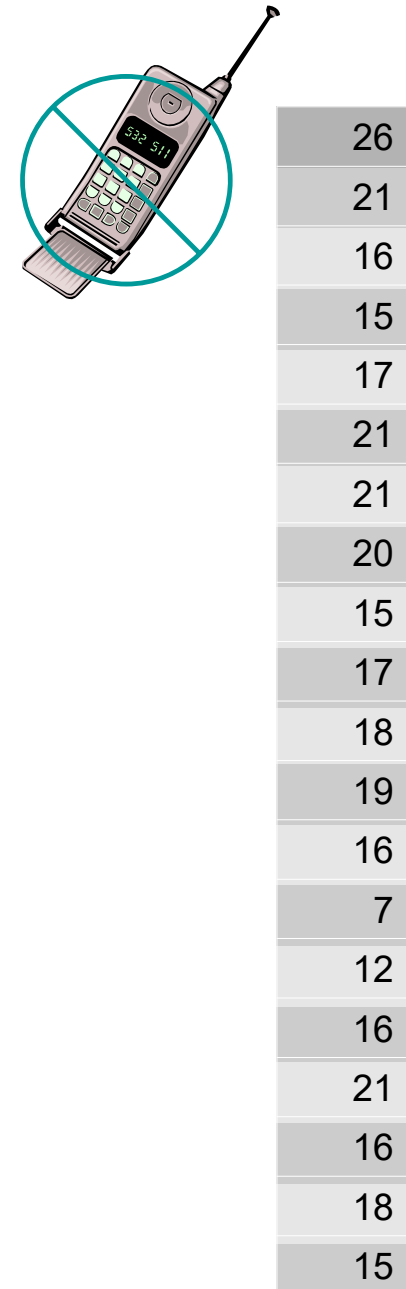
•

•

•

•

*Consider our cellphone driving example ...*



*And these are the data we collected (20 subjects in each condition)*

# 5 steps for hypothesis proving

1. State the Null Hypothesis  $H_0$  and Alternative Hypothesis  $H_1$ .

$H_0$ : sample mean a  $\leq$  sample mean b

$H_1$ : sample mean a  $>$  sample mean b

2. Select the appropriate statistical method.

Independent sample t-test, 1-tailed

3. Select a level of significance.

.05; df = 38; critical t-value needs to be larger than 1.684

4. Calculate the statistic.

5. Make the decision.



|    |
|----|
| 17 |
| 25 |
| 33 |
| 17 |
| 18 |
| 18 |
| 22 |
| 25 |
| 18 |
| 28 |
| 23 |
| 27 |
| 26 |
| 22 |
| 14 |
| 20 |
| 24 |
| 23 |
| 25 |
| 20 |

*Mean = 22.5*  
*STDEV = 4.59*



|    |
|----|
| 26 |
| 21 |
| 16 |
| 15 |
| 17 |
| 21 |
| 21 |
| 20 |
| 15 |
| 17 |
| 18 |
| 19 |
| 16 |
| 7  |
| 12 |
| 16 |
| 21 |
| 16 |
| 18 |
| 15 |

*Mean = 17.35*  
*STDEV = 3.96*



## 2. Independent Sample t-test

$$t = \frac{\bar{X}_1 - \bar{X}_2}{\sqrt{\left[ \frac{(N_1 - 1)s_1^2 + (N_2 - 1)s_2^2}{N_1 + N_2 - 2} \right] \left[ \frac{1}{N_1} + \frac{1}{N_2} \right]}}$$

The diagram includes three labels with arrows pointing to parts of the formula:   
 - A blue arrow points from 'A' to the numerator  $\bar{X}_1 - \bar{X}_2$ .   
 - A green arrow points from 'B' to the square root symbol.   
 - A red arrow points from 'C' to the term  $\left[ \frac{1}{N_1} + \frac{1}{N_2} \right]$ .

- $X_1 = 22.25$ ;  $X_2 = 17.35$ ;
- $N_1 = 20$
- $N_2 = 20$
- $S_1 = 4.59$
- $S_2 = 3.96$
- $A = 4.9$
- $B = 18.38$
- $C = 0.1$
- $t = 3.61$

# 5 steps for hypothesis proving

1. State the Null Hypothesis  $H_0$  and Alternative Hypothesis  $H_1$ .

$H_0$ : sample mean a  $\leq$  sample mean b

$H_1$ : sample mean a  $>$  sample mean b

2. Select the appropriate statistical method.

Independent group t-test, 1-tailed

3. Select a level of significance.

.05; df = 38; critical t-value needs to be larger than 1.684

4. Calculate the statistic.

$t(38) = 3.61$ ;  $p < .05$

5. Make the decision.

$H_0$  is reject;  $H_1$  is supported.

The phone condition has a significantly higher score than the no phone condition.



# How to do it in PSPP

\*phone.sav [DataSet1] — PSPPIRE Data Editor

File Edit View Data Transform Analyze Graphs Utilities Windows Help

Icons: Save, Open, Undo, Redo, Print, Find, etc.

| Variable | Name      | Type    | Width | Decimal | Label | Value Labels | Missing Values | Column | Align | Measure | Role  |
|----------|-----------|---------|-------|---------|-------|--------------|----------------|--------|-------|---------|-------|
| 1        | Score     | Numeric | 8     | 2       |       | None         | None           | 8      | Right | Scale   | Input |
| 2        | Condition | Numeric | 8     | 2       |       | None         | None           | 8      | Right | Scale   | Input |
| 3        |           |         |       |         |       |              |                |        |       |         |       |

Data View Variable View

Filter off Weights off No Split

# How to do it in PSPP

\*phone.sav [DataSet1] — PSPPIRE Data Editor

File Edit View Data Transform Analyze Graphs Utilities Windows Help

Icons: Save, Open, Undo, Redo, Find, Sort, Filter, Split, Tag

| Case | Score | Condition |  |
|------|-------|-----------|--|
| 1    | 17.00 | 1.00      |  |
| 2    | 25.00 | 1.00      |  |
| 3    | 33.00 | 1.00      |  |
| 4    | 17.00 | 1.00      |  |
| 5    | 18.00 | 1.00      |  |
| 6    | 18.00 | 1.00      |  |
| 7    | 22.00 | 1.00      |  |
| 8    | 25.00 | 1.00      |  |
| 9    | 18.00 | 1.00      |  |
| 10   | 28.00 | 1.00      |  |
| 11   | 23.00 | 1.00      |  |
| 12   | 27.00 | 1.00      |  |
| 13   | 26.00 | 1.00      |  |
| 14   | 22.00 | 1.00      |  |
| 15   | 14.00 | 1.00      |  |
| 16   | 20.00 | 1.00      |  |
| 17   | 24.00 | 1.00      |  |
| 18   | 23.00 | 1.00      |  |
| 19   | 25.00 | 1.00      |  |
| 20   | 20.00 | 1.00      |  |
| 21   | 26.00 | 2.00      |  |
| 22   | 21.00 | 2.00      |  |
| 23   | 16.00 | 2.00      |  |

Data View Variable View

Filter off Weights off No Split

# How to do it in PSPP

\*phone.sav [DataSet1] — PSPP Data Editor

File Edit View Data Transform Analyze Graphs Utilities Windows Help

Descriptive Statistics  
Compare Means  
Univariate Analysis...  
Bivariate Correlation...  
K-Means Cluster...  
Factor Analysis...  
Reliability...  
Regression  
Non-Parametric Statistics  
ROC Curve...

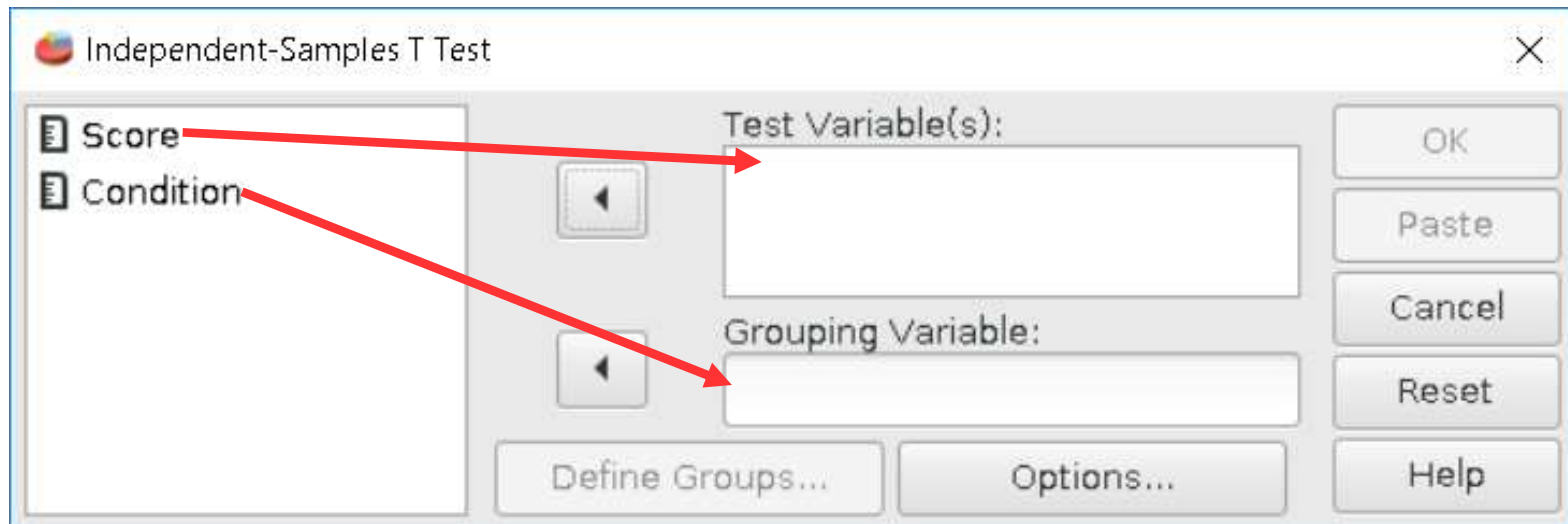
Means...  
One Sample T Test...  
Independent Samples T Test...  
Paired Samples T Test...  
One Way ANOVA...

| Case | Score | Condition |
|------|-------|-----------|
| 1    | 17.00 | 1.00      |
| 2    | 25.00 | 1.00      |
| 3    | 33.00 | 1.00      |
| 4    | 17.00 | 1.00      |
| 5    | 18.00 | 1.00      |
| 6    | 18.00 | 1.00      |
| 7    | 22.00 | 1.00      |
| 8    | 25.00 | 1.00      |
| 9    | 18.00 | 1.00      |
| 10   | 28.00 | 1.00      |
| 11   | 23.00 | 1.00      |
| 12   | 27.00 | 1.00      |
| 13   | 26.00 | 1.00      |
| 14   | 22.00 | 1.00      |
| 15   | 14.00 | 1.00      |
| 16   | 20.00 | 1.00      |
| 17   | 24.00 | 1.00      |
| 18   | 23.00 | 1.00      |
| 19   | 25.00 | 1.00      |
| 20   | 20.00 | 1.00      |
| 21   | 26.00 | 2.00      |
| 22   | 21.00 | 2.00      |
| 23   | 16.00 | 2.00      |

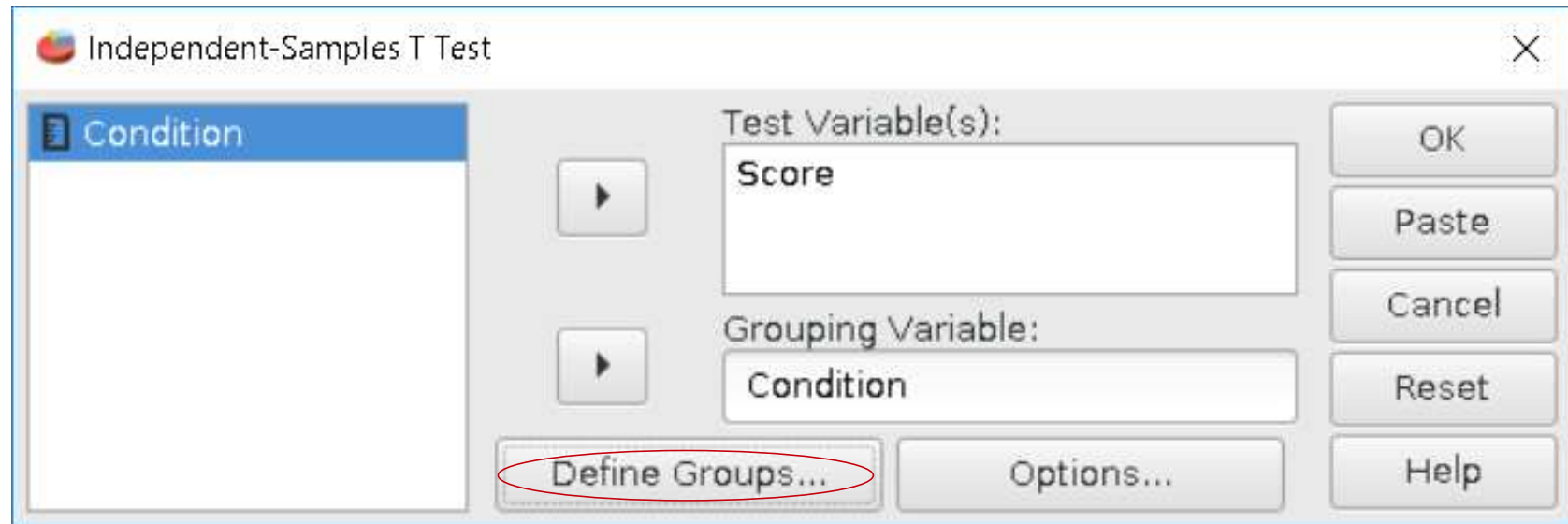
Data View Variable View

Filter off Weights off No Split

# How to do it in PSPP

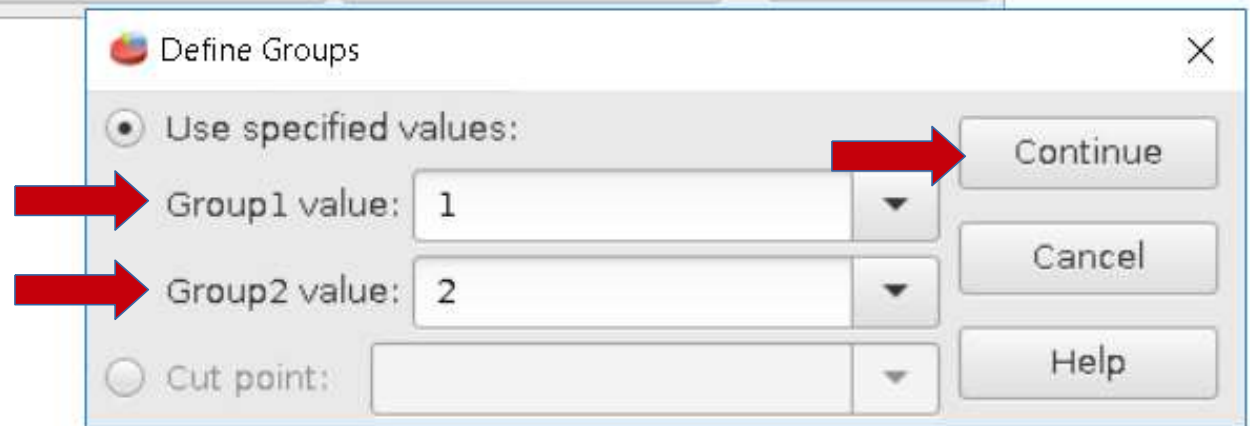
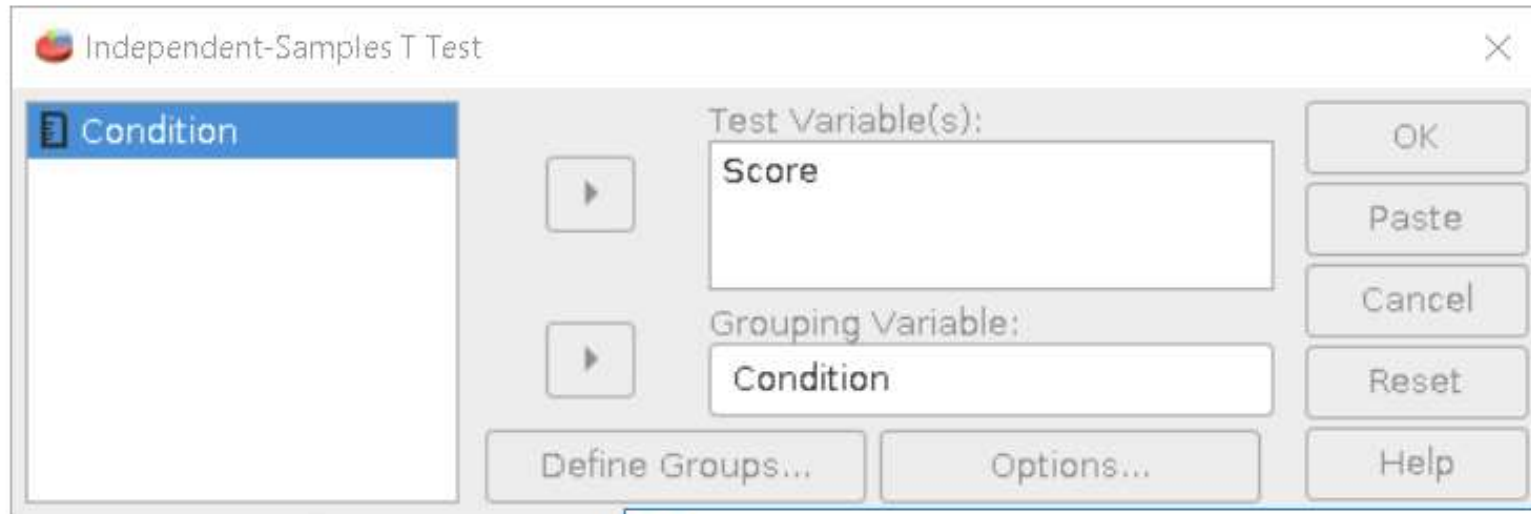


# How to do it in PSPP

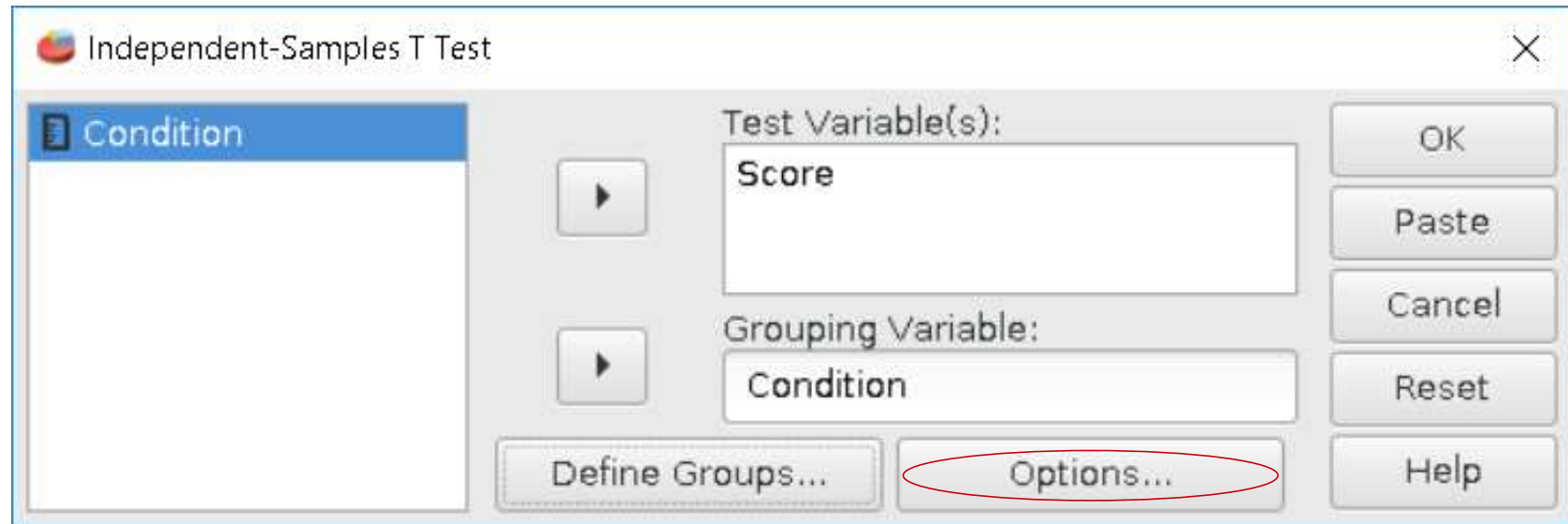




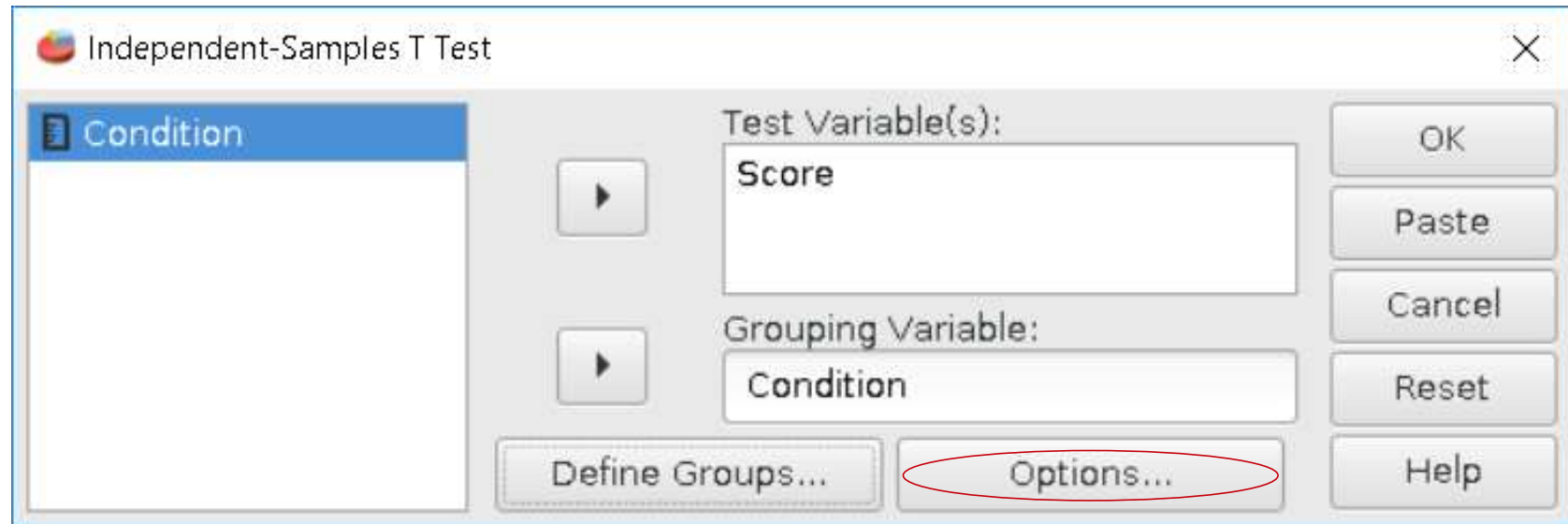
# How to do it in PSPP



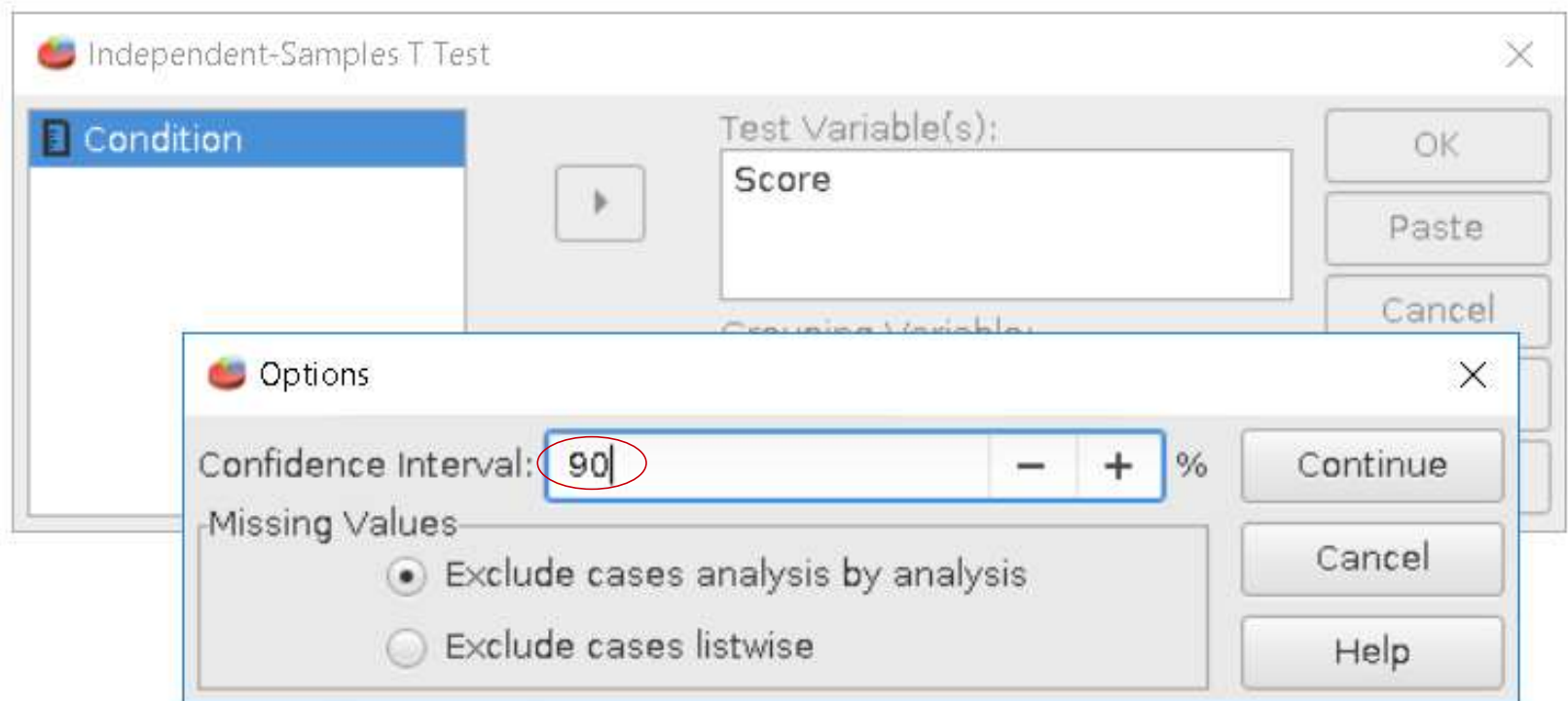
# How to do it in PSPP



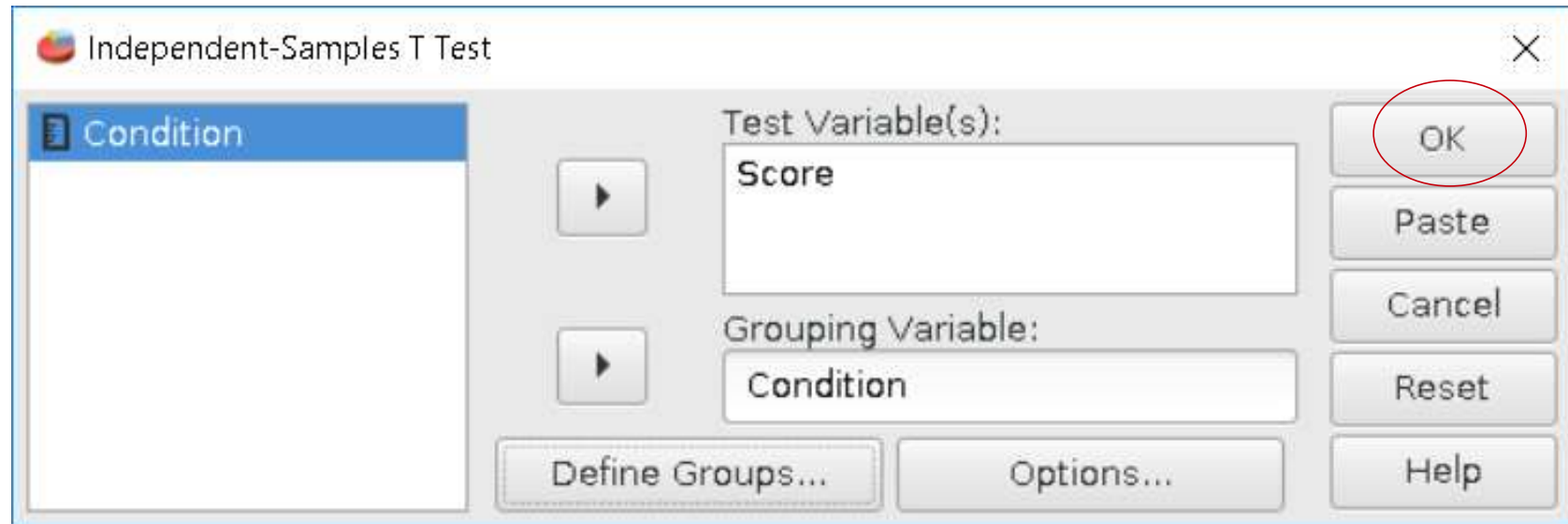
# How to do it in PSPP



# Set 90% confidence interval ... (For 1-Tailed 95% confidence)



# How to do it in PSPP



# Result

## T-TEST

T-TEST /VARIABLES= Score  
/GROUPS=Condition(1,2) /MISSING=ANALYSIS  
/CRITERIA=CI(0.9).

## Descriptive Statistic

## Group Statistics

|       | Condition | N  | Mean  | Std. Deviation | S.E. Mean |
|-------|-----------|----|-------|----------------|-----------|
| Score | 1.00      | 20 | 22.25 | 4.59           | 1.03      |
|       | 2.00      | 20 | 17.35 | 3.96           | .89       |

## Independent Samples Test

|       |                             | Levene's Test for Equality of Variances |      | t-test for Equality of Means |       |                 |                 |                       |   |       |
|-------|-----------------------------|---|------|------------------------------|-------|-----------------|-----------------|-----------------------|---|-------|
|       |                             | F                                       | Sig. | t                            | df    | Sig. (2-tailed) | Mean Difference | Std. Error Difference | 90% Confidence Interval of the Difference |       |
|       |                             |   |      |                              |       |                 |                 |                       | Lower                                     | Upper |
| Score | Equal variances assumed     | .84                                     | .366 | 3.61                         | 38.00 | .001            | 4.90            | 1.36                  | 2.61                                      | 7.19  |
|       | Equal variances not assumed |   |      | 3.61                         | 37.22 | .001            | 4.90            | 1.36                  | 2.61                                      | 7.19  |

## Levene's Test Result

- If Sig. is below .05, then Levene's Test shows significance
- i.e. The variance between groups is not homogenous (not equal)
- In that case, we should read the second row in the t-test result
- In our case, Levene's Test does not show significance
- Read the first row (equal variances)

# Result

## T-TEST

T-TEST /VARIABLES= Score  
/GROUPS=Condition(1,2) /MISSING=ANALYSIS  
/CRITERIA=CI(0.9).

### Group Statistics

|       | Condition | N  | Mean  | Std. Deviation | S.E. Mean |
|-------|-----------|----|-------|----------------|-----------|
| Score | 1.00      | 20 | 22.25 | 4.59           | 1.03      |
|       | 2.00      | 20 | 17.35 | 3.96           | .89       |

Descriptive Statistic

### Independent Samples Test

|       |                             | Levene's Test for Equality of Variances |      | t-test for Equality of Means |       |                 |                 |                       |   |      |
|-------|-----------------------------|---|------|------------------------------|-------|-----------------|-----------------|-----------------------|---|------|
|       |                             | F                                       | Sig. | t                            | df    | Sig. (2-tailed) | Mean Difference | Std. Error Difference | 90% Confidence Interval of the Difference |      |
| Score | Equal variances assumed     | .84                                     | .366 | 3.61                         | 38.00 | .001            | 4.90            | 1.36                  | 2.61                                      | 7.19 |
|       | Equal variances not assumed |   |      | 3.61                         | 37.22 | .001            | 4.90            | 1.36                  | 2.61                                      | 7.19 |

t-value

Significance value (p value) for 2-tailed  
p value for 1-tailed should be divided by 2  
i.e. .0005






### 3. Paired Sample t-test

- Used for comparing the mean of one group to the mean of two dependent conditions
- Test if the average difference of two measurements is different from zero
- Used for within-subject design comparison
- Often used for “before-after effect” analysis
- Requirement:
  - Must be 2 conditions
  - Measurement should be scaled
    - i.e. not categorical (e.g. male vs. female)
- No requirement for variances homogeneity
  - Levene’s Test is not required
  -

# NHST Settings

- Null Hypothesis  $H_0$ :
  - The mean difference between condition 1 and condition 2 is zero
  - $\text{Mean}_1 - \text{Mean}_2 = 0$
- Alternative Hypothesis  $H_1$ :
  - The mean difference between condition 1 and condition 2 is not zero
  - $\text{Mean}_1 - \text{Mean}_2 \neq 0$
- $df = N - 1$
- 2-tailed (since you're comparing if the means are the same or different)

# Scenario

- You created 2 interfaces for the same task
  - Performance of the task was measured
  - $N = 20$  (i.e. 20 subjects)
  - Using within subject design
    - The same subject uses Interface #1 and Interface #2 to complete the task
    - Performances were compared within subject
    - Performance was measured using a 10-point scale
    - Sequence of the Interfaces was counterbalanced
- 

# The Data

| Subject | Interface1   | Interface2   |
|---------|--------------|--------------|
| 1       | 9            | 1            |
| 2       | 7            | 5            |
| 3       | 8            | 4            |
| 4       | 5            | 10           |
| 5       | 7            | 5            |
| 6       | 4            | 4            |
| 7       | 10           | 10           |
| 8       | 3            | 5            |
| 9       | 10           | 5            |
| 10      | 10           | 5            |
| 11      | 7            | 6            |
| 12      | 6            | 4            |
| 13      | 5            | 3            |
| 14      | 4            | 6            |
| 15      | 10           | 3            |
| 16      | 6            | 3            |
| 17      | 5            | 5            |
| 18      | 7            | 5            |
| 19      | 5            | 2            |
| 20      | 7            | 10           |
| Mean    | 6.75         | 5.05         |
| STDEV   | 2.1974866026 | 2.4809802816 |

# 5 steps for hypothesis proving

1. State the Null Hypothesis  $H_0$  and Alternative Hypothesis  $H_1$ .

$H_0$ : sample mean 1 = sample mean 2

$H_1$ : sample mean 1  $\neq$  sample mean 2

2. Select the appropriate statistical method.

Paired sample t-test, 2-tailed

3. Select a level of significance.

.05; df = 19; critical t-value needs to be larger than 2.093

4. Calculate the statistic.

5. Make the decision.

# Formula for paired-sample t-test

$$t = \frac{\frac{\sum d}{N}}{\sqrt{\frac{\sum d^2 - \frac{(\sum d)^2}{N}}{N(N-1)}}$$

# Formula for paired-sample t-test

$$t = \frac{\frac{\sum d}{N}}{\sqrt{\frac{\sum d^2 - \frac{(\sum d)^2}{N}}{N(N-1)}}$$

$$t = (34/20) / \sqrt{(260 - (34^2/20)) / (20 \times 19)}$$

= 2.33

| Subject | Interface1 | Interface2 | d  | d square |
|---------|------------|------------|----|----------|
| 1       | 9          | 1          | 8  | 64       |
| 2       | 7          | 5          | 2  | 4        |
| 3       | 8          | 4          | 4  | 16       |
| 4       | 5          | 10         | -5 | 25       |
| 5       | 7          | 5          | 2  | 4        |
| 6       | 4          | 4          | 0  | 0        |
| 7       | 10         | 10         | 0  | 0        |
| 8       | 3          | 5          | -2 | 4        |
| 9       | 10         | 5          | 5  | 25       |
| 10      | 10         | 5          | 5  | 25       |
| 11      | 7          | 6          | 1  | 1        |
| 12      | 6          | 4          | 2  | 4        |
| 13      | 5          | 3          | 2  | 4        |
| 14      | 4          | 6          | -2 | 4        |
| 15      | 10         | 3          | 7  | 49       |
| 16      | 6          | 3          | 3  | 9        |
| 17      | 5          | 5          | 0  | 0        |
| 18      | 7          | 5          | 2  | 4        |
| 19      | 5          | 2          | 3  | 9        |
| 20      | 7          | 10         | -3 | 9        |
| Sum     | 135        | 101        | 34 | 260      |

# 5 steps for hypothesis proving

1. State the Null Hypothesis  $H_0$  and Alternative Hypothesis  $H_1$ .

$H_0$ : sample mean 1 = sample mean 2

$H_1$ : sample mean 1  $\neq$  sample mean 2

2. Select the appropriate statistical method.

Paired sample t-test, 2-tailed

3. Select a level of significance.

.05; df = 19; critical t-value needs to be larger than 2.093

4. Calculate the statistic.

$t(19) = 2.33$ ;  $p < .05$

5. Make the decision.

$H_0$  is reject;  $H_1$  is supported.











Mean for Interface 1 is significantly different from Mean for Interface 2





# How to do it in PSPP

\*[DataSet2] — PSPPIRE Data Editor

| File  | Edit  | View  | Data  | Transform   | Analyze   | Graphs  | Utilities   | Windows   | Help  |
|---|---|---|---|---|---|---|---|---|---|
|  |  |  |  |  |  |  |  |  |  |

| Case | Interface1 | Interface2 |  |
|------|------------|------------|--|
| 1    | 9.00       | 1.00       |  |
| 2    | 7.00       | 5.00       |  |
| 3    | 8.00       | 4.00       |  |
| 4    | 5.00       | 10.00      |  |
| 5    | 7.00       | 5.00       |  |
| 6    | 4.00       | 4.00       |  |
| 7    | 10.00      | 10.00      |  |
| 8    | 3.00       | 5.00       |  |
| 9    | 10.00      | 5.00       |  |
| 10   | 10.00      | 5.00       |  |
| 11   | 7.00       | 6.00       |  |
| 12   | 6.00       | 4.00       |  |
| 13   | 5.00       | 3.00       |  |
| 14   | 4.00       | 6.00       |  |
| 15   | 10.00      | 3.00       |  |
| 16   | 6.00       | 3.00       |  |
| 17   | 5.00       | 5.00       |  |
| 18   | 7.00       | 5.00       |  |
| 19   | 5.00       | 2.00       |  |
| 20   | 7.00       | 10.00      |  |
| 21   | .          | .          |  |
| 22   |            |            |  |

|                  |                      |
|------------------|----------------------|
| <b>Data View</b> | <b>Variable View</b> |
|                  | Filter off           |
|                  | Weights off          |
|                  | No Split             |

# How to do it in PSPP

\*[DataSet2] — PSPPIRE Data Editor

File Edit View Data Transform **Analyze** Graphs Utilities Windows Help

Descriptive Statistics  
**Compare Means**  
Univariate Analysis...  
Bivariate Correlation...  
K-Means Cluster...  
Factor Analysis...  
Reliability...  
Regression  
Non-Parametric Statistics  
ROC Curve...

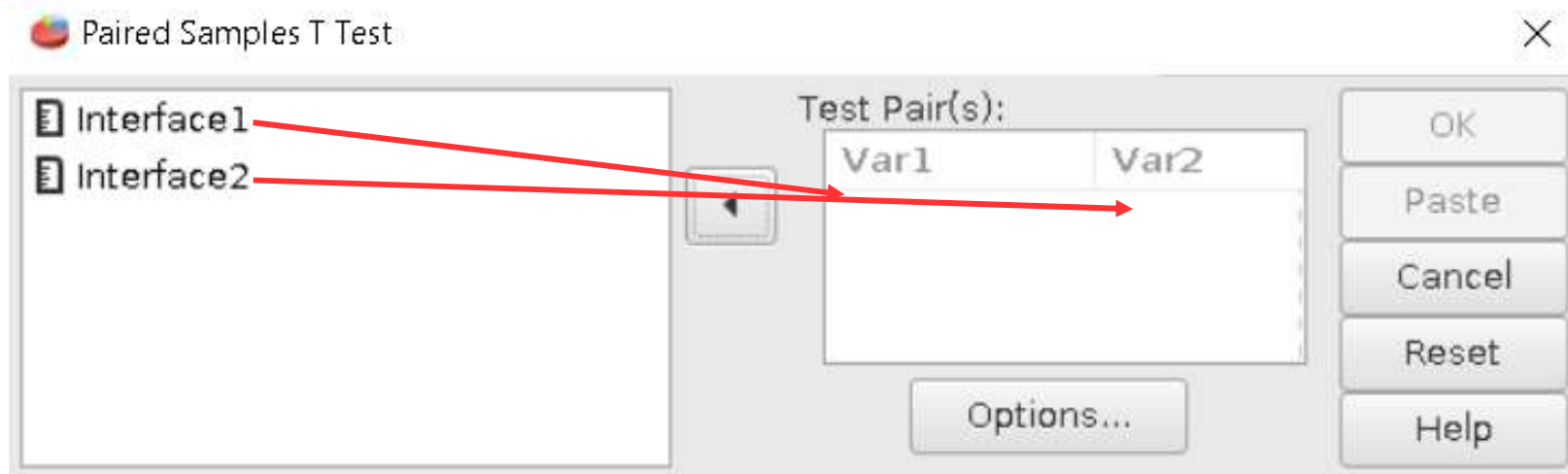
Means...  
One Sample T Test...  
Independent Samples T Test...  
**Paired Samples T Test...**  
One Way ANOVA...

| Case | Interface1 | Interface2 |
|------|------------|------------|
| 1    | 9.00       | 1.00       |
| 2    | 7.00       | 5.00       |
| 3    | 8.00       | 4.00       |
| 4    | 5.00       | 10.00      |
| 5    | 7.00       | 5.00       |
| 6    | 4.00       | 4.00       |
| 7    | 10.00      | 10.00      |
| 8    | 3.00       | 5.00       |
| 9    | 10.00      | 5.00       |
| 10   | 10.00      | 5.00       |
| 11   | 7.00       | 6.00       |
| 12   | 6.00       | 4.00       |
| 13   | 5.00       | 3.00       |
| 14   | 4.00       | 6.00       |
| 15   | 10.00      | 3.00       |
| 16   | 6.00       | 3.00       |
| 17   | 5.00       | 5.00       |
| 18   | 7.00       | 5.00       |
| 19   | 5.00       | 2.00       |
| 20   | 7.00       | 10.00      |
| 21   | .          | .          |
| 22   |            |            |

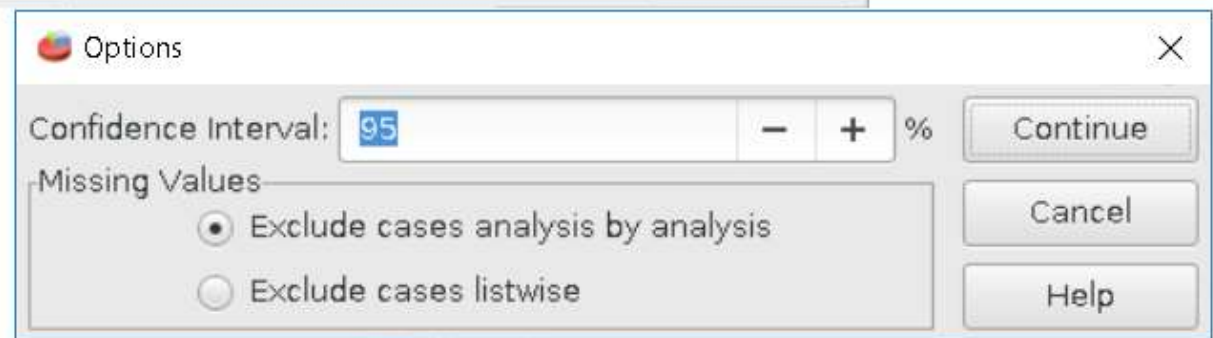
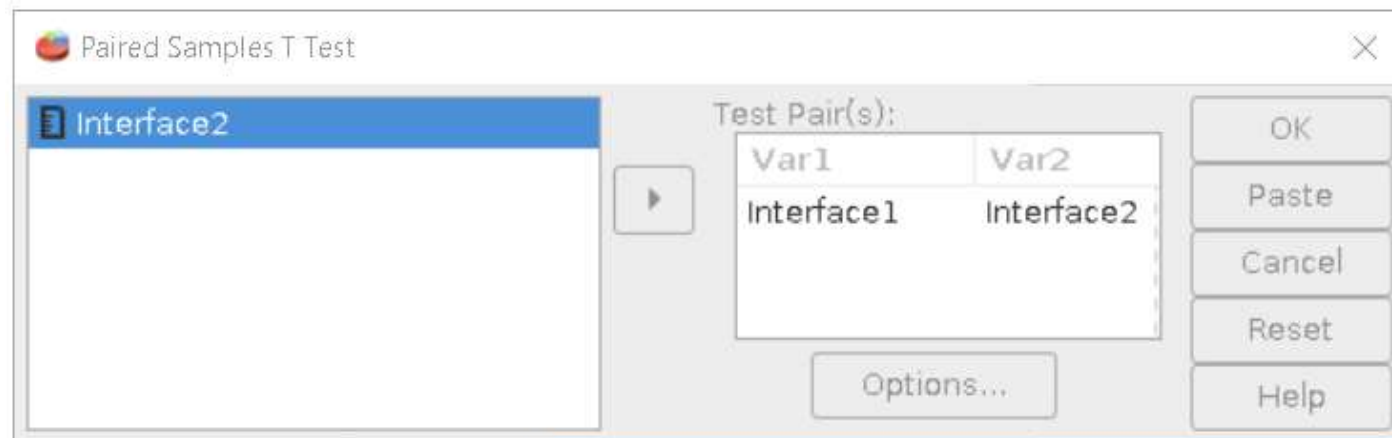
**Data View** Variable View

Filter off Weights off No Split

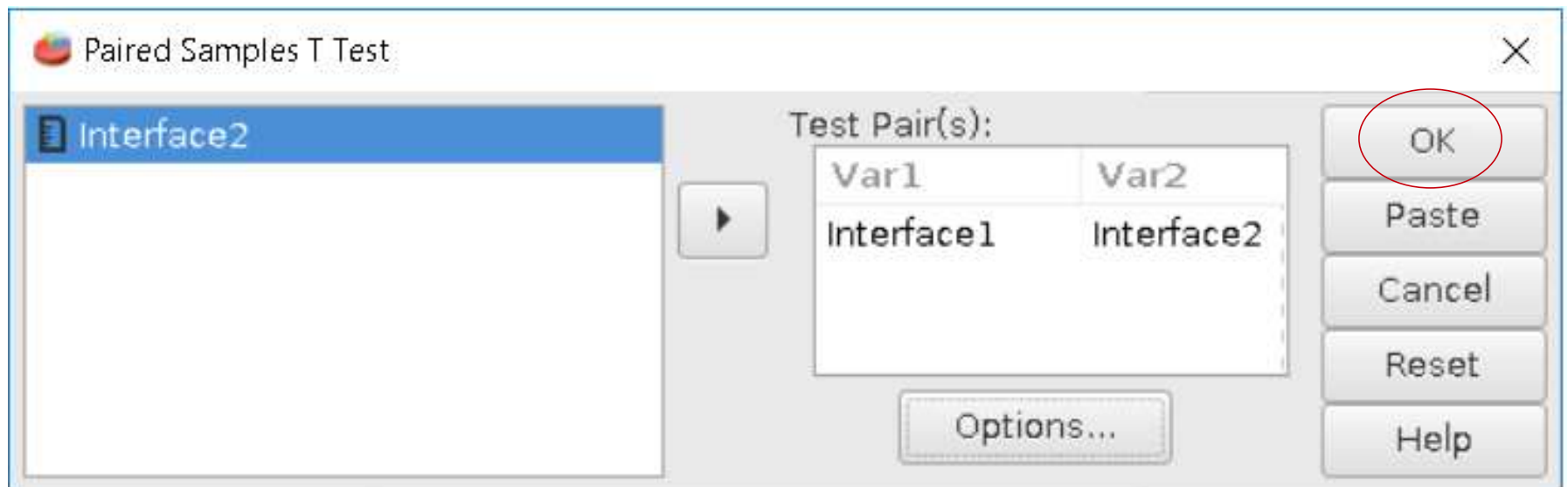
# How to do it in PSPP



# Set 95% Confidence Interval for 2-tailed test



# How to do it in PSPP



# Result

T-TEST

T-TEST

PAIRS = Interface1 WITH Interface2 (PAIRED)  
/MISSING=ANALYSIS  
/CRITERIA=CI(0.95).

## Paired Sample Statistics

|        |            | Mean | N  | Std. Deviation | S.E. Mean |
|--------|------------|------|----|----------------|-----------|
| Pair 1 | Interface1 | 6.75 | 20 | 2.20           | .49       |
|        | Interface2 | 5.05 | 20 | 2.48           | .55       |

Descriptive Statistic

## Paired Samples Correlations

|        |                         | N  | Correlation | Sig. |
|--------|-------------------------|----|-------------|------|
| Pair 1 | Interface1 & Interface2 | 20 | .03         | .896 |

## Paired Samples Test

|        |                         | Paired Differences |                |                 |   |       | t    | df | Sig. (2-tailed) |
|--------|-------------------------|--------------------|----------------|-----------------|---|-------|------|----|-----------------|
|        |                         | Mean               | Std. Deviation | Std. Error Mean | 95% Confidence Interval of the Difference |       |      |    |                 |
|        |                         |                    |                |                 | Lower                                     | Upper |      |    |                 |
| Pair 1 | Interface1 - Interface2 | 1.70               | 3.26           | .73             | .17                                       | 3.23  | 2.33 | 19 | .031            |

# Summary: t-test

## T-tests

For comparing 2 mean values

### 1. Single sample t-test

Comparing a sample mean to a population mean

### 2. Independent sample t-test

Comparing two sample mean values

- The two sample means has to be independent
- Levene's Test must not pass

### 3. Paired sample t-test

Comparing two sample mean values

- The two sample means does not need to be independent
- Levene's Test not required





# What if you have more than 2 conditions?

For example, you may have one Independent Variable (IV) with three levels:

Interface #1

Interface #2

Interface #3

When your IV has more than 2 levels, you can't use the t-test!!!

You need to use ANOVA!




# ANOVA

ANOVA – Analysis of Variances

Similar to t-test, but for more than 2 conditions

Different variants of ANOVA

- One way ANOVA
  - MANOVA
  - ANCOVA
  - MANCOVA
- 

# Requirement of ANOVA

## Independent Variable (IV)

- must be independent
- Sample size of each condition should be roughly equal

## Dependent Variable (DV)

- Measurement must be quantitative
- Normality: DV must be normally distributed
- Homogeneity of Variance:  
Equal variance (tested by Levene's Test)

# NHST using ANOVA

1. State the Null Hypothesis  $H_0$  and Alternative Hypothesis  $H_1$ .

Null Hypothesis  $N_0$ : Mean1 = Mean2 = Mean3

Alternative Hypothesis  $N_1$ : Mean1  $\neq$  Mean2  $\neq$  Mean3

2. Select the appropriate statistical method.

One way ANOVA

3. Select a level of significance.

.05

4. Calculate the statistic.

5. Make the decision.

6. Calculate the post-hoc test

# A note about the hypotheses

**In the case of 3 conditions:**

Null Hypothesis  $N_0$ : **Mean1 = Mean2 = Mean3**

Alternative Hypothesis  $N_1$ : **Mean1 != Mean2 != Mean3**

**Mean1 = Mean2 = Mean3**

Mean1=Mean2 AND Mean2=Mean3 AND Mean1=Mean3

=> (Mean1=Mean2 AND Mean1=Mean3 AND Mean2=Mean3)

**Mean1 != Mean2 != Mean3**

Mean1 != Mean2 OR Mean2 != Mean3 OR Mean1 != Mean3

=> (Mean1!=Mean2 AND Mean1=Mean3 AND Mean2=Mean3) OR  
(Mean1=Mean2 AND Mean1!=Mean3 AND Mean2=Mean3) OR  
(Mean1=Mean2 AND Mean1=Mean3 AND Mean2!=Mean3) OR  
(Mean1!=Mean2 AND Mean1!=Mean3 AND Mean2=Mean3) OR  
(Mean1=Mean2 AND Mean1!=Mean3 AND Mean2!=Mean3) OR  
(Mean1!=Mean2 AND Mean1!=Mean3 AND Mean2!=Mean3)

# A note about the hypotheses

When an ANOVA analysis is significant,

- (1) the Null Hypothesis is rejected;
- (2) the Alternative Hypothesis is supported.

That means at least one of the following cases are significant:

(Mean1!=Mean2 AND Mean1=Mean3 AND Mean2=Mean3) OR  
(Mean1=Mean2 AND Mean1!=Mean3 AND Mean2=Mean3) OR  
(Mean1=Mean2 AND Mean1=Mean3 AND Mean2!=Mean3) OR  
(Mean1!=Mean2 AND Mean1!=Mean3 AND Mean2=Mean3) OR  
(Mean1=Mean2 AND Mean1!=Mean3 AND Mean2!=Mean3) OR  
(Mean1!=Mean2 AND Mean1!=Mean3 AND Mean2!=Mean3)

But we don't know which cases are significant!!!

A further **post-hoc test** is required in order to find this out!

# NHST using ANOVA

1. State the Null Hypothesis  $H_0$  and Alternative Hypothesis  $H_1$ .

Null Hypothesis  $N_0$ : Mean1 = Mean2 = Mean3

Alternative Hypothesis  $N_1$ : Mean1  $\neq$  Mean2  $\neq$  Mean3

2. Select the appropriate statistical method.

One way ANOVA

3. Select a level of significance.

.05

4. Calculate the statistic.

I am skipping the maths and stats for this course.....

Let's go directly into SPSS and let the computer do the calculations!

5. Make the decision.

6. Calculate the post-hoc test







|    |
|----|
| 17 |
| 25 |
| 33 |
| 17 |
| 18 |
| 18 |
| 22 |
| 25 |
| 18 |
| 28 |
| 23 |
| 27 |
| 26 |
| 22 |
| 14 |
| 20 |
| 24 |
| 23 |
| 25 |
| 20 |

*Mean = 22.5*  
*STDEV = 4.59*



|    |
|----|
| 26 |
| 21 |
| 16 |
| 15 |
| 17 |
| 21 |
| 21 |
| 20 |
| 15 |
| 17 |
| 18 |
| 19 |
| 16 |
| 7  |
| 12 |
| 16 |
| 21 |
| 16 |
| 18 |
| 15 |

*Mean = 17.35*  
*STDEV = 3.96*



|    |
|----|
| 16 |
| 26 |
| 34 |
| 16 |
| 19 |
| 15 |
| 24 |
| 22 |
| 19 |
| 29 |
| 24 |
| 19 |
| 24 |
| 26 |
| 19 |
| 21 |
| 26 |
| 22 |
| 22 |
| 22 |

*Mean = 22.5*  
*STDEV = 4.64*

# How to do it in PSPP

\*anova.sav [DataSet1] — PSPPIRE Data Editor

| File | Edit      | View  | Data | Transform | Analyze | Graphs | Utilities | Windows | Help |
|------|-----------|-------|------|-----------|---------|--------|-----------|---------|------|
|      |           |       |      |           |         |        |           |         |      |
| Case | condition | score |      |           |         |        |           |         |      |
| 1    | 1.00      | 17.00 |      |           |         |        |           |         |      |
| 2    | 1.00      | 25.00 |      |           |         |        |           |         |      |
| 3    | 1.00      | 33.00 |      |           |         |        |           |         |      |
| 4    | 1.00      | 17.00 |      |           |         |        |           |         |      |
| 5    | 1.00      | 18.00 |      |           |         |        |           |         |      |
| 6    | 1.00      | 18.00 |      |           |         |        |           |         |      |
| 7    | 1.00      | 22.00 |      |           |         |        |           |         |      |
| 8    | 1.00      | 25.00 |      |           |         |        |           |         |      |
| 9    | 1.00      | 18.00 |      |           |         |        |           |         |      |
| 10   | 1.00      | 28.00 |      |           |         |        |           |         |      |
| 11   | 1.00      | 23.00 |      |           |         |        |           |         |      |
| 12   | 1.00      | 27.00 |      |           |         |        |           |         |      |
| 13   | 1.00      | 26.00 |      |           |         |        |           |         |      |
| 14   | 1.00      | 22.00 |      |           |         |        |           |         |      |
| 15   | 1.00      | 14.00 |      |           |         |        |           |         |      |
| 16   | 1.00      | 20.00 |      |           |         |        |           |         |      |
| 17   | 1.00      | 24.00 |      |           |         |        |           |         |      |
| 18   | 1.00      | 23.00 |      |           |         |        |           |         |      |
| 19   | 1.00      | 25.00 |      |           |         |        |           |         |      |
| 20   | 1.00      | 20.00 |      |           |         |        |           |         |      |
| 21   | 2.00      | 26.00 |      |           |         |        |           |         |      |
| 22   | 2.00      | 21.00 |      |           |         |        |           |         |      |
| 23   | 2.00      | 16.00 |      |           |         |        |           |         |      |

Data View

Variable View











Filter off

Weights off

No Split

\*anova.sav [DataSet1] — PSPPIRE Data Editor

File Edit View Data Transform Analyze Graphs Utilities Windows Help



| Case | condition | score |  |
|------|-----------|-------|--|
| 39   | 2.00      | 18.00 |  |
| 40   | 2.00      | 15.00 |  |
| 41   | 3.00      | 16.00 |  |
| 42   | 3.00      | 26.00 |  |
| 43   | 3.00      | 34.00 |  |
| 44   | 3.00      | 16.00 |  |
| 45   | 3.00      | 19.00 |  |
| 46   | 3.00      | 15.00 |  |
| 47   | 3.00      | 24.00 |  |
| 48   | 3.00      | 22.00 |  |
| 49   | 3.00      | 19.00 |  |
| 50   | 3.00      | 29.00 |  |
| 51   | 3.00      | 24.00 |  |
| 52   | 3.00      | 19.00 |  |
| 53   | 3.00      | 24.00 |  |
| 54   | 3.00      | 26.00 |  |
| 55   | 3.00      | 19.00 |  |
| 56   | 3.00      | 21.00 |  |
| 57   | 3.00      | 26.00 |  |
| 58   | 3.00      | 22.00 |  |
| 59   | 3.00      | 22.00 |  |
| 60   | 3.00      | 22.00 |  |
| 61   | .         | .     |  |

Data View

Variable View

Filter off

Weights off

No Split

# How to do it in PSPP

\*anova.sav [DataSet1] — PSPP Data Editor

File Edit View Data Transform **Analyze** Graphs Utilities Windows Help

Descriptive Statistics  
**Compare Means**  
Univariate Analysis...  
Bivariate Correlation...  
K-Means Cluster...  
Factor Analysis...  
Reliability...  
Regression  
Non-Parametric Statistics  
ROC Curve...

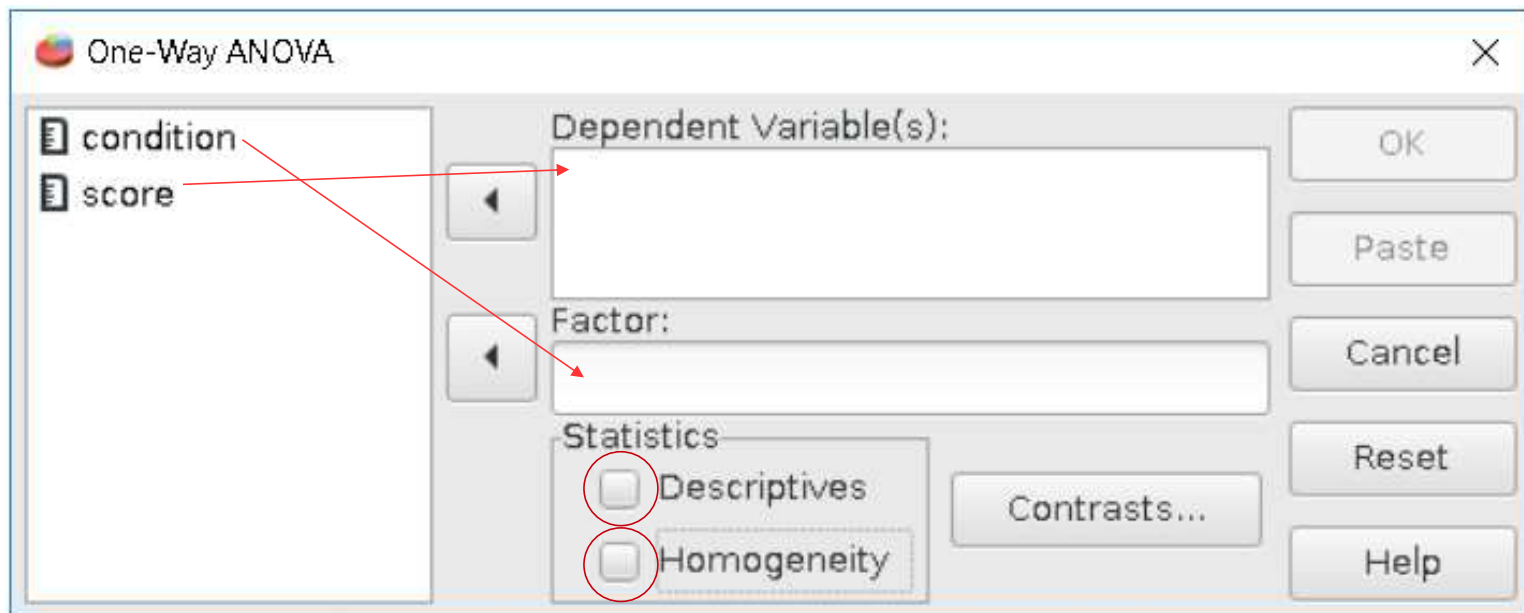
Means...  
One Sample T Test...  
Independent Samples T Test...  
Paired Samples T Test...  
**One Way ANOVA...**

| Case | condition | score |
|------|-----------|-------|
| 39   | 2.00      | 16.00 |
| 40   | 2.00      | 15.00 |
| 41   | 3.00      | 16.00 |
| 42   | 3.00      | 26.00 |
| 43   | 3.00      | 34.00 |
| 44   | 3.00      | 16.00 |
| 45   | 3.00      | 19.00 |
| 46   | 3.00      | 15.00 |
| 47   | 3.00      | 24.00 |
| 48   | 3.00      | 22.00 |
| 49   | 3.00      | 19.00 |
| 50   | 3.00      | 29.00 |
| 51   | 3.00      | 24.00 |
| 52   | 3.00      | 19.00 |
| 53   | 3.00      | 24.00 |
| 54   | 3.00      | 26.00 |
| 55   | 3.00      | 19.00 |
| 56   | 3.00      | 21.00 |
| 57   | 3.00      | 26.00 |
| 58   | 3.00      | 22.00 |
| 59   | 3.00      | 22.00 |
| 60   | 3.00      | 22.00 |
| 61   | .         | .     |

**Data View** Variable View

Filter off Weights off No Split

# How to do it in PSPP



# Results

ONEWAY

ONEWAY /VARIABLES= score BY condition  
/STATISTICS=DESCRIPTIVES HOMOGENEITY .

## Descriptives

|       |       | N  | Mean  | Std. Deviation | Std. Error | 95% Confidence Interval for Mean |             | Minimum | Maximum |
|-------|-------|----|-------|----------------|------------|----------------------------------|-------------|---------|---------|
|       |       |    |       |                |            | Lower Bound                      | Upper Bound |         |         |
| score | 1.00  | 20 | 22.25 | 4.59           | 1.03       | 20.10                            | 24.40       | 14.00   | 33.00   |
|       | 2.00  | 20 | 17.35 | 3.96           | .89        | 15.49                            | 19.21       | 7.00    | 26.00   |
|       | 3.00  | 20 | 22.25 | 4.64           | 1.04       | 20.08                            | 24.42       | 15.00   | 34.00   |
|       | Total | 60 | 20.62 | 4.92           | .64        | 19.35                            | 21.89       | 7.00    | 34.00   |

## Test of Homogeneity of Variances

|       | Levene Statistic | df1 | df2 | Sig. |
|-------|------------------|-----|-----|------|
| score | .44              | 2   | 57  | .649 |

## ANOVA

|       |                | Sum of Squares | df | Mean Square | F    | Sig. |
|-------|----------------|----------------|----|-------------|------|------|
| score | Between Groups | 320.13         | 2  | 160.07      | 8.23 | .001 |
|       | Within Groups  | 1108.05        | 57 | 19.44       |      |      |
|       | Total          | 1428.18        | 59 |             |      |      |

Descriptive  
Statistics

Levene's  
Test

# NHST using ANOVA

1. State the Null Hypothesis  $H_0$  and Alternative Hypothesis  $H_1$ .

Null Hypothesis  $N_0$ : Mean1 = Mean2 = Mean3

Alternative Hypothesis  $N_1$ : Mean1  $\neq$  Mean2  $\neq$  Mean3

2. Select the appropriate statistical method.

One way ANOVA

3. Select a level of significance.

.05

4. Calculate the statistic.

I am skipping the maths and stats for this course.....

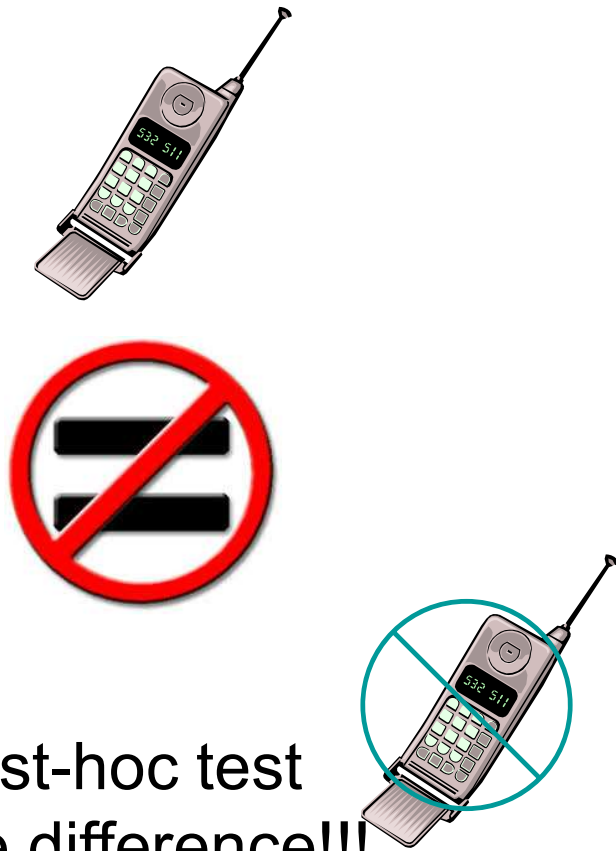
Let's go directly into SPSS and let the computer do the calculations!

5. Make the decision.

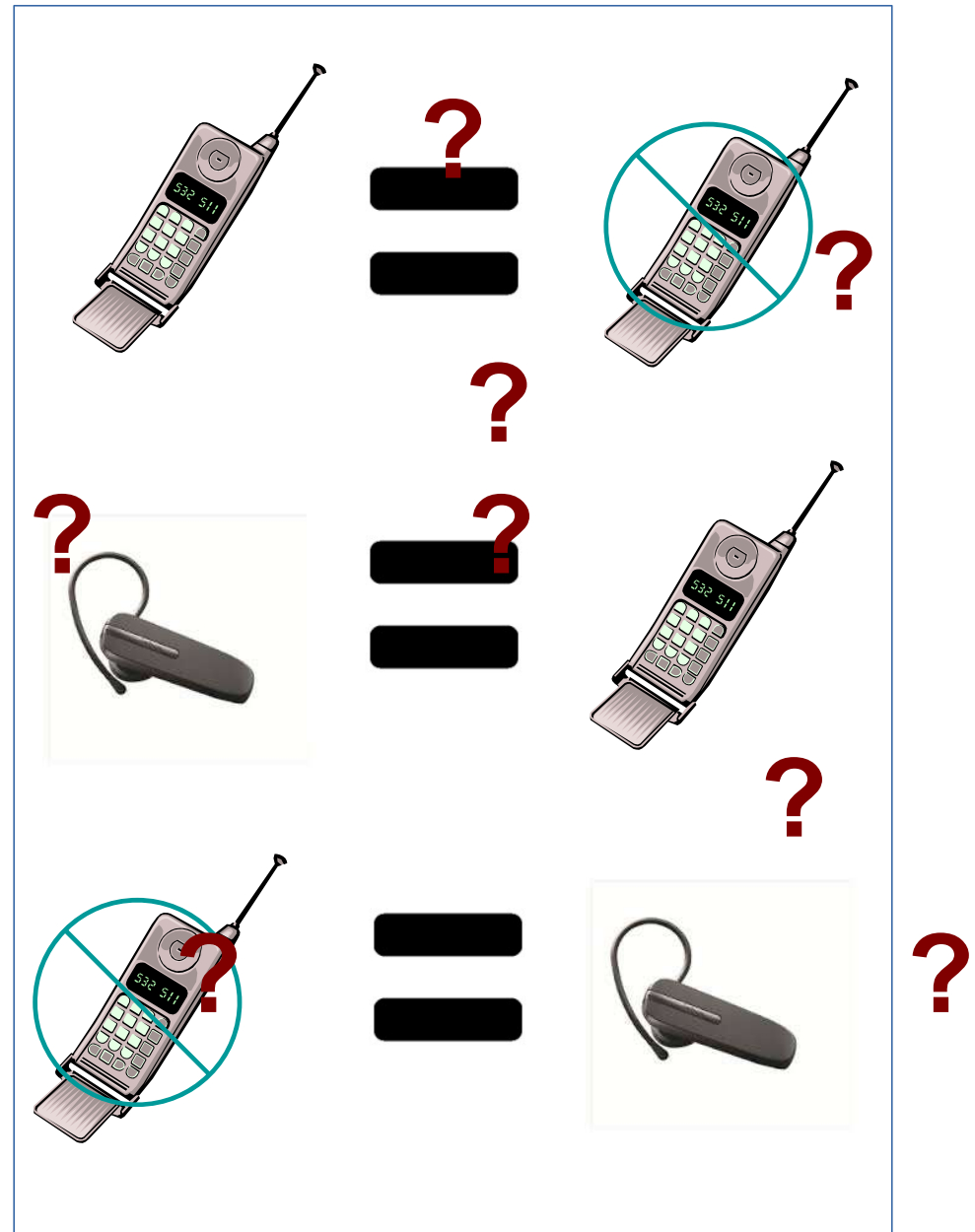
$H_0$  is rejected.

- We know [Mean1  $\neq$  Mean2  $\neq$  Mean3]

6. Calculate the post-hoc test



Need Post-hoc test  
to tell the difference!!!





# How to do posthoc test in PSPP

The screenshot displays the PSPP Data Editor interface. The 'Analyze' menu is open, showing the path: Analyze > Compare Means > One Way ANOVA... The 'One Way ANOVA...' option is highlighted. The data table below shows 14 cases with columns for 'Case', 'condition', and 'score'.

| Case | condition | score |
|------|-----------|-------|
| 1    | 1.00      | 17.00 |
| 2    | 1.00      | 25.00 |
| 3    | 1.00      | 33.00 |
| 4    | 1.00      | 17.00 |
| 5    | 1.00      | 18.00 |
| 6    | 1.00      | 18.00 |
| 7    | 1.00      | 22.00 |
| 8    | 1.00      | 25.00 |
| 9    | 1.00      | 18.00 |
| 10   | 1.00      | 28.00 |
| 11   | 1.00      | 23.00 |
| 12   | 1.00      | 27.00 |
| 13   | 1.00      | 26.00 |
| 14   | 1.00      | 22.00 |

At the bottom of the window, the 'Data View' tab is active, and the status bar shows 'Filter off', 'Weights off', and 'No Split'.

# How to do posthoc test in PSPP

The screenshot displays the PSPP (Predictive Software for Psychology) interface. On the left is the 'PSPPIRE Syntax Editor' window, which is currently empty. On the right is the '\*anova.sav [DataSet1] — PSPPIRE Data Editor' window, showing a data table with columns 'Case', 'condition', and 'score'. The data table contains 14 rows of data. A 'One-Way ANOVA' dialog box is open in the center, with 'condition' selected as the factor and 'score' as the dependent variable. The 'Statistics' section is checked for 'Descriptives' and 'Homogeneity'. The 'Contrasts...' button is visible. The 'Paste' button is highlighted with a red circle. The bottom of the window shows the 'Data View' tab selected, with 'Filter off', 'Weights off', and 'No Split' options.

| Case | condition | score |
|------|-----------|-------|
| 1    | 1.00      | 17.00 |
| 2    |           |       |
| 3    |           |       |
| 4    |           |       |
| 5    |           |       |
| 6    |           |       |
| 7    |           |       |
| 8    |           |       |
| 9    |           |       |
| 10   | 1.00      | 28.00 |
| 11   | 1.00      | 23.00 |
| 12   | 1.00      | 27.00 |
| 13   | 1.00      | 26.00 |
| 14   | 1.00      | 22.00 |

One-Way ANOVA

Dependent Variable(s):  
score

Factor:  
condition

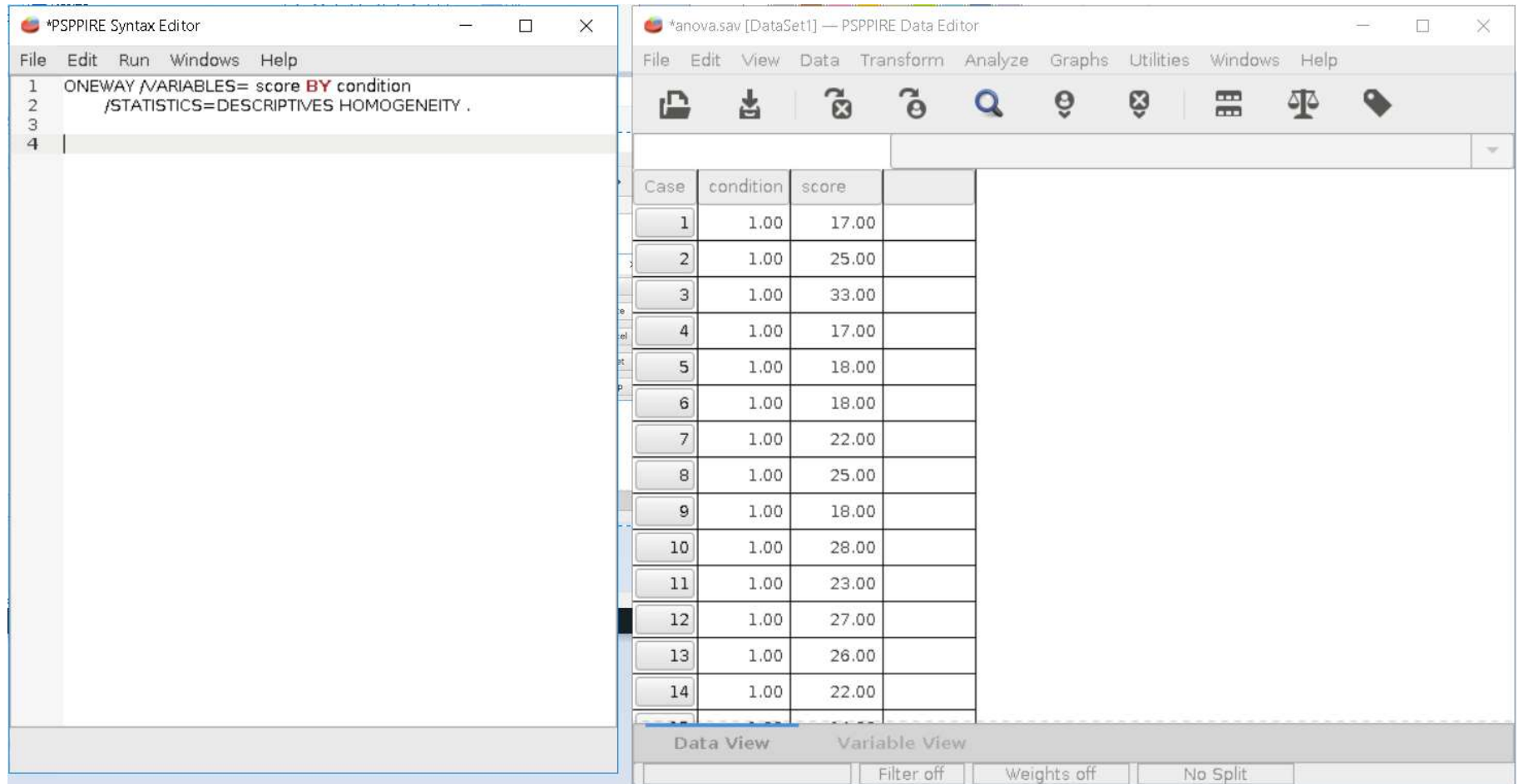
Statistics  
☒ Descriptives  
☒ Homogeneity

Contrasts...

OK  
Paste  
Cancel  
Reset  
Help

Data View Variable View  
Filter off Weights off No Split

# How to do posthoc test in PSPP



The image shows two windows from the PSPP (Predictive Statistical Projector) software. The left window is the 'Syntax Editor' and the right window is the 'Data Editor'.

**PSPP Syntax Editor:**

```
1 ONEWAY /VARIABLES= score BY condition
2 /STATISTICS=DESCRIPTIVES HOMOGENEITY .
3
4
```

**PSPP Data Editor:**

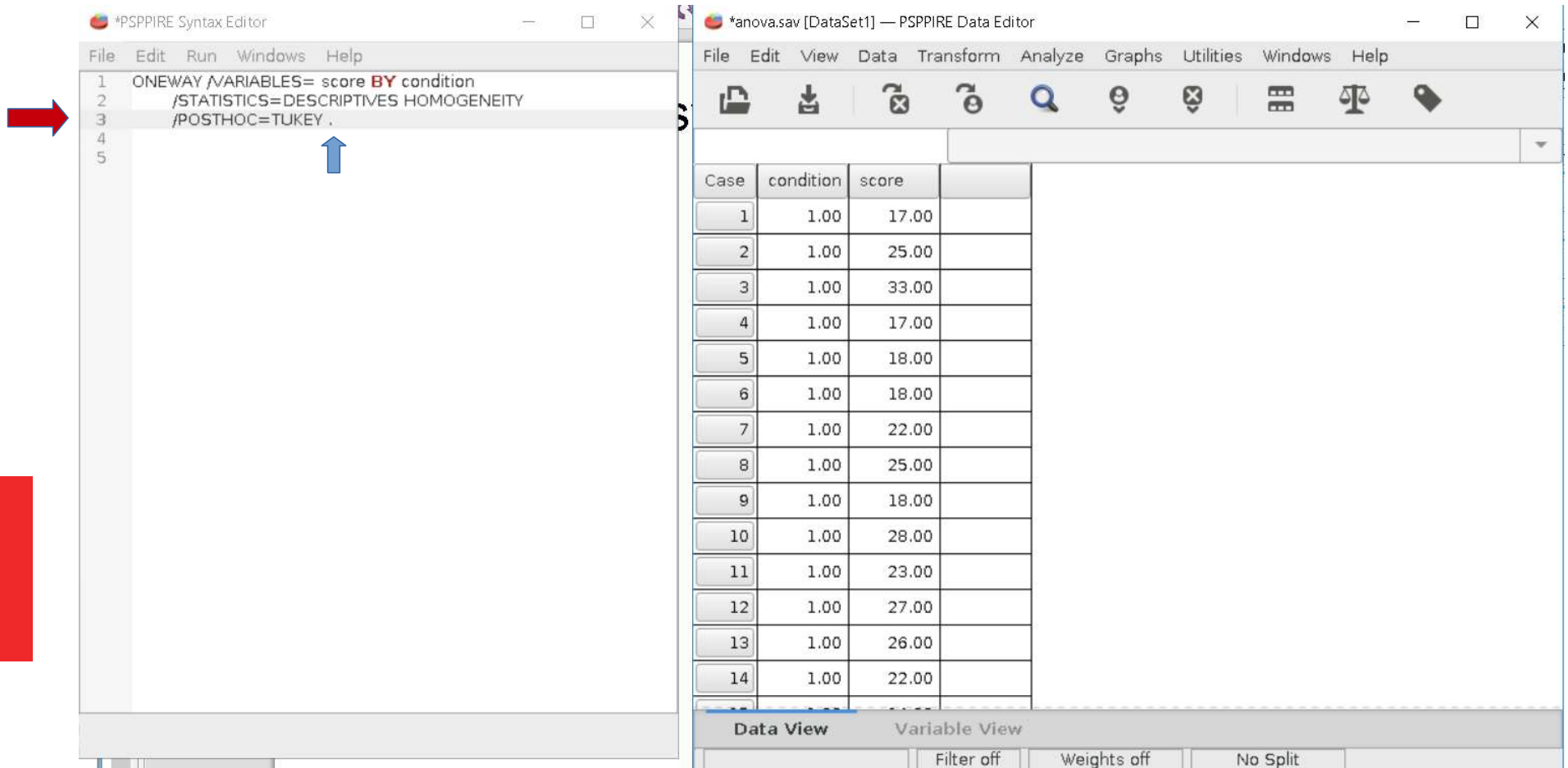
The data editor shows a table with 14 cases. The columns are 'Case', 'condition', and 'score'.

| Case | condition | score |
|------|-----------|-------|
| 1    | 1.00      | 17.00 |
| 2    | 1.00      | 25.00 |
| 3    | 1.00      | 33.00 |
| 4    | 1.00      | 17.00 |
| 5    | 1.00      | 18.00 |
| 6    | 1.00      | 18.00 |
| 7    | 1.00      | 22.00 |
| 8    | 1.00      | 25.00 |
| 9    | 1.00      | 18.00 |
| 10   | 1.00      | 28.00 |
| 11   | 1.00      | 23.00 |
| 12   | 1.00      | 27.00 |
| 13   | 1.00      | 26.00 |
| 14   | 1.00      | 22.00 |

The bottom of the Data Editor window shows tabs for 'Data View' and 'Variable View', with 'Data View' selected. Below the tabs are buttons for 'Filter off', 'Weights off', and 'No Split'.

# Now, edit the syntax to add posthoc test

Note: the full stop indicate the end of the script



The image shows two windows from the PSPPIRE software. The left window, titled '\*PSPPIRE Syntax Editor', contains the following syntax script:

```
1 ONEWAY /VARIABLES= score BY condition
2 /STATISTICS=DESCRIPTIVES HOMOGENEITY
3 /POSTHOC=DUKEY .
4
5
```

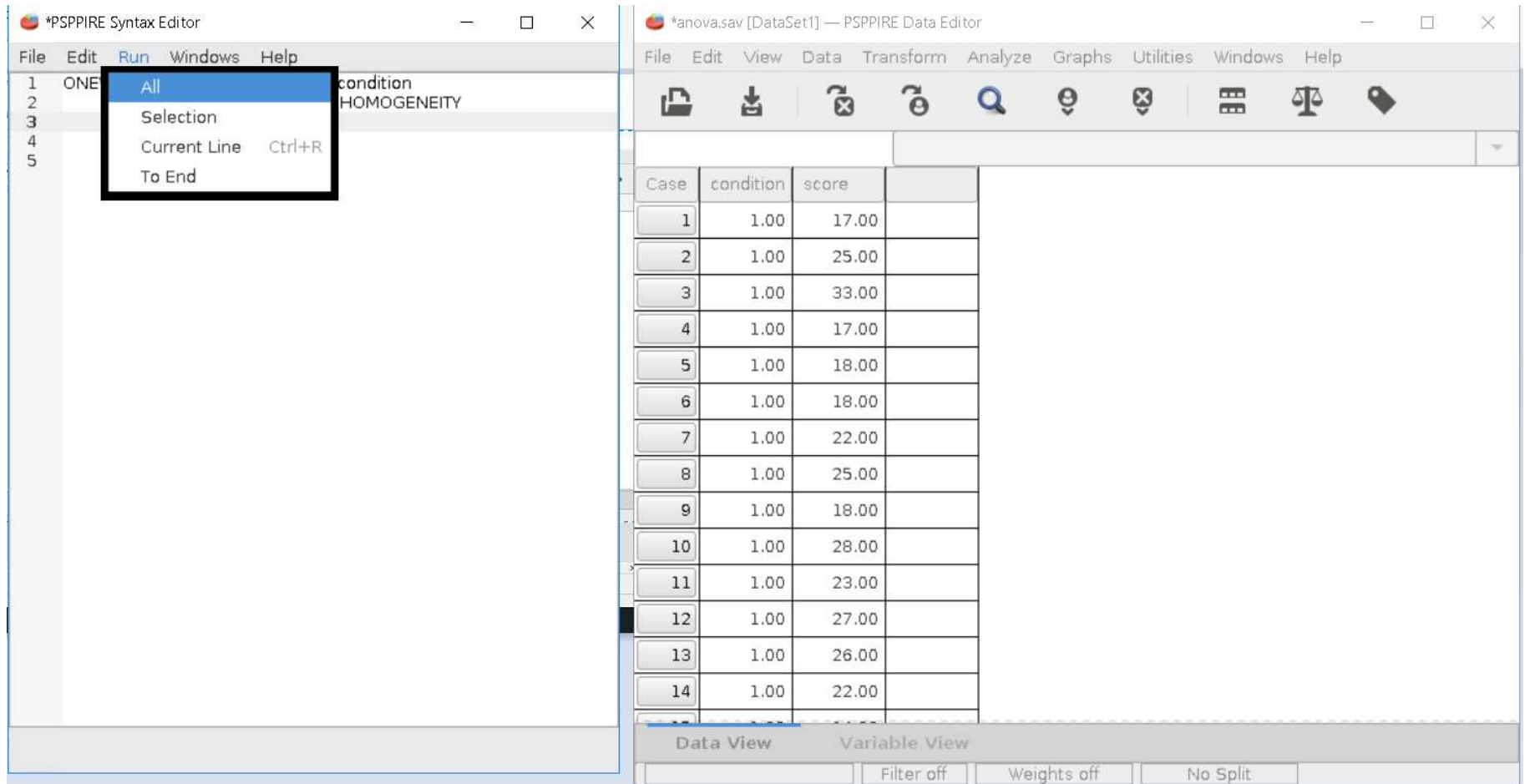
A red arrow points to the first line of the script, and a blue arrow points to the period at the end of the third line. The right window, titled '\*anova.sav [DataSet1] — PSPPIRE Data Editor', displays a data table with 14 cases. The table has three columns: 'Case', 'condition', and 'score'.

| Case | condition | score |
|------|-----------|-------|
| 1    | 1.00      | 17.00 |
| 2    | 1.00      | 25.00 |
| 3    | 1.00      | 33.00 |
| 4    | 1.00      | 17.00 |
| 5    | 1.00      | 18.00 |
| 6    | 1.00      | 18.00 |
| 7    | 1.00      | 22.00 |
| 8    | 1.00      | 25.00 |
| 9    | 1.00      | 18.00 |
| 10   | 1.00      | 28.00 |
| 11   | 1.00      | 23.00 |
| 12   | 1.00      | 27.00 |
| 13   | 1.00      | 26.00 |
| 14   | 1.00      | 22.00 |

The bottom of the right window shows the 'Data View' tab selected, with buttons for 'Filter off', 'Weights off', and 'No Split'.

# Now, edit the syntax to add posthoc test

Note: the full stop indicate the end of the script



The image shows two windows from the PSPPIRE software. The left window, titled '\*PSPPIRE Syntax Editor', displays a script with line numbers 1 through 5. A context menu is open over the script, showing options: 'All', 'Selection', 'Current Line Ctrl+R', and 'To End'. The right window, titled '\*anova.sav [DataSet1] — PSPPIRE Data Editor', shows a data table with columns 'Case', 'condition', and 'score'. The table contains 14 rows of data. Below the table, there are tabs for 'Data View' and 'Variable View', and buttons for 'Filter off', 'Weights off', and 'No Split'.

| Case | condition | score |
|------|-----------|-------|
| 1    | 1.00      | 17.00 |
| 2    | 1.00      | 25.00 |
| 3    | 1.00      | 33.00 |
| 4    | 1.00      | 17.00 |
| 5    | 1.00      | 18.00 |
| 6    | 1.00      | 18.00 |
| 7    | 1.00      | 22.00 |
| 8    | 1.00      | 25.00 |
| 9    | 1.00      | 18.00 |
| 10   | 1.00      | 28.00 |
| 11   | 1.00      | 23.00 |
| 12   | 1.00      | 27.00 |
| 13   | 1.00      | 26.00 |
| 14   | 1.00      | 22.00 |

# Result

ONEWAY

ONEWAY /VARIABLES= score BY condition  
/STATISTICS=DESCRIPTIVES HOMOGENEITY  
/POSTHOC=TUKEY .

Descriptives

|       |       | N  | Mean  | Std. Deviation | Std. Error | 95% Confidence Interval for Mean |             | Minimum | Maximum |
|-------|-------|----|-------|----------------|------------|----------------------------------|-------------|---------|---------|
|       |       |    |       |                |            | Lower Bound                      | Upper Bound |         |         |
| score | 1.00  | 20 | 22.25 | 4.59           | 1.03       | 20.10                            | 24.40       | 14.00   | 33.00   |
|       | 2.00  | 20 | 17.35 | 3.96           | .89        | 15.49                            | 19.21       | 7.00    | 26.00   |
|       | 3.00  | 20 | 22.25 | 4.64           | 1.04       | 20.08                            | 24.42       | 15.00   | 34.00   |
|       | Total | 60 | 20.62 | 4.92           | .64        | 19.35                            | 21.89       | 7.00    | 34.00   |

Test of Homogeneity of Variances

|       | Levene Statistic | df1 | df2 | Sig. |
|-------|------------------|-----|-----|------|
| score | .44              | 2   | 57  | .649 |

ANOVA

|       |                | Sum of Squares | df | Mean Square | F    | Sig. |
|-------|----------------|----------------|----|-------------|------|------|
| score | Between Groups | 320.13         | 2  | 160.07      | 8.23 | .001 |
|       | Within Groups  | 1108.05        | 57 | 19.44       |      |      |
|       | Total          | 1428.18        | 59 |             |      |      |

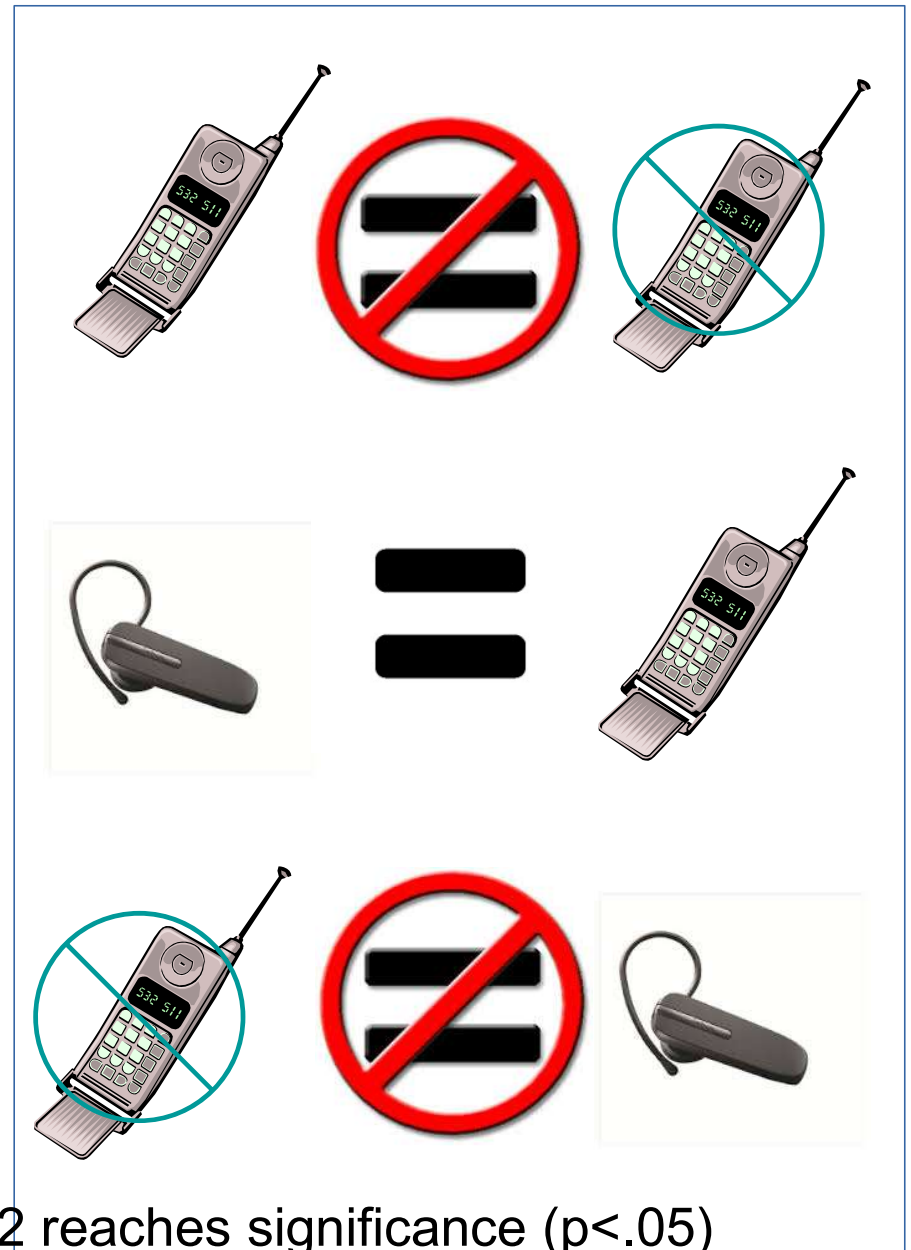
Multiple Comparisons (score)

|               |               | Mean Difference<br>(i - j) |       | Std. Error | Sig.  | 95% Confidence Interval |             |
|---------------|---------------|----------------------------|-------|------------|-------|-------------------------|-------------|
| (i) condition | (j) condition |                            |       |            |       | Lower Bound             | Upper Bound |
| Tukey HSD     | 1.00          | 2.00                       | 4.90  | 1.39       | .002  | 1.54                    | 8.26        |
|               |               | 3.00                       | .00   | 1.39       | 1.000 | -3.36                   | 3.36        |
|               | 2.00          | 1.00                       | -4.90 | 1.39       | .002  | -8.26                   | -1.54       |
|               |               | 3.00                       | -4.90 | 1.39       | .002  | -8.26                   | -1.54       |
|               | 3.00          | 1.00                       | .00   | 1.39       | 1.000 | -3.36                   | 3.36        |
|               |               | 2.00                       | 4.90  | 1.39       | .002  | 1.54                    | 8.26        |

Effect between Condition 1 and Condition 2 reaches significance ( $p < .05$ )

Effect between Condition 1 and Condition 3 does not reach significance ( $p > .05$ )

Effect between Condition 2 and Condition 3 reaches significance ( $p < .05$ )



Effect between Condition 1 and Condition 2 reaches significance ( $p < .05$ )  
Effect between Condition 1 and Condition 3 does not reach significance ( $p > .05$ )  
Effect between Condition 2 and Condition 3 reaches significance ( $p < .05$ )

