

The work of medical entomologist Bart Knols revealed that a species of mosquito known for its penchant for sucking blood from people's feet and ankles also quite likes the stinky German cheese, Limberger. Bemused by this discovery upon watching Dr Knols' TED talk¹, a researcher decides to try to extend this research by assessing mosquito preferences for a range of stinky cheeses in the hopes of finding the ultimate cheese mosquito magnet. The researcher captures 10 mosquitoes and releases them one by one into four sealed plastic boxes, each containing four different cheese varieties for 30 seconds. Sensors built into the boxes record the amount of time each mosquito spends within a 1cm radius of the cheeses. The researcher is presented with data that reveals how long each mosquito spent hovering over each cheese variety and decides to conduct a one-way repeated measures ANOVA to determine which cheese is most attractive to mosquitoes.

Step 1 – Taking a look at the data.

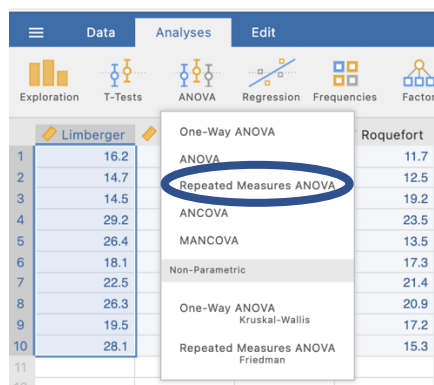
The screenshot shows the Jamovi 'DATA VARIABLE' settings for the variable 'Limberger'. The 'Measure type' is set to 'Continuous', the 'Data type' is 'Decimal', and 'Missing values' is empty. Below the settings is a data table with 10 rows of mosquito data across four cheese varieties: Limberger, Epoisses, Blue Stilton, and Roquefort.

	Limberger	Epoisses	Blue Stilton	Roquefort
1	16.2	21.2	5.6	11.7
2	14.7	16.7	3.2	12.5
3	14.5	15.1	7.2	19.2
4	29.2	12.5	1.4	23.5
5	26.4	17.3	8.3	13.5
6	18.1	11.5	11.1	17.3
7	22.5	22.1	17.3	21.4
8	26.3	24.3	15.2	20.9
9	19.5	18.6	8.1	17.2
10	28.1	20.6	9.4	15.3

Our data takes the form of four continuous variables that represent the number of seconds spent by each mosquito hovering within 1cm of each of four stinky cheese varieties, namely German Limberger, French Epoisses and Roquefort and British Blue Stilton. The variables are designated continuous.

¹ You can view Dr Knols' TED talk here <https://ed.ted.com/lessons/cheese-dogs-and-a-pill-to-kill-mosquitoes-and-end-malaria-bart-knols>.

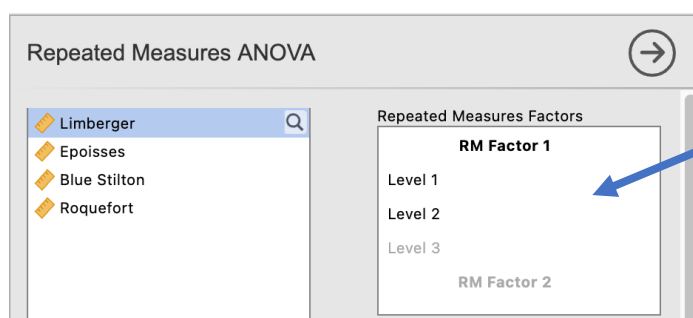
Step 3 – Navigating to the Repeated Measures ANOVA analysis menu.



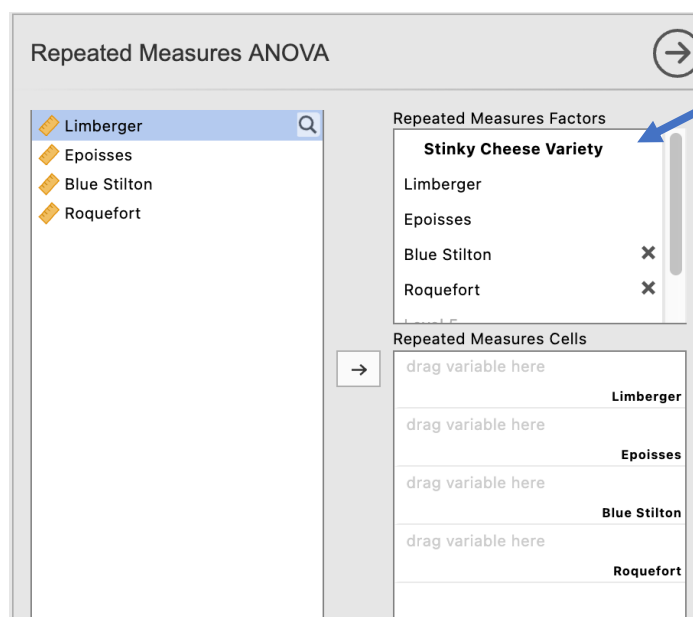
On the Analyses tab select the ANOVA menu, then select Repeated Measures ANOVA.

Step 4a – Selecting analysis options to get the output we need for our omnibus results

The first thing we will do is to let *jamovi* know that these four variables represent levels of our repeated measures factor. This set up is somewhat more complex than a between groups ANOVA.



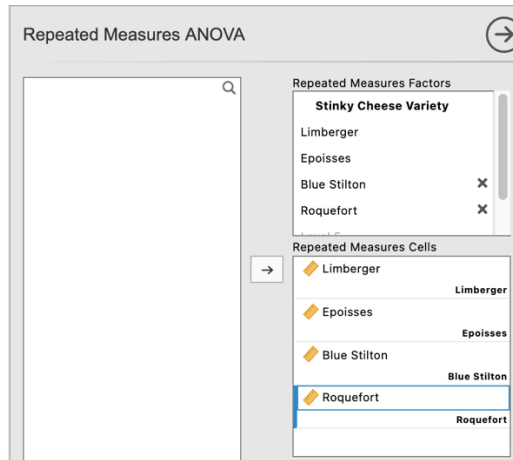
Within the Repeated Measures Factors box we need to type in the labels of our factor and its four levels.



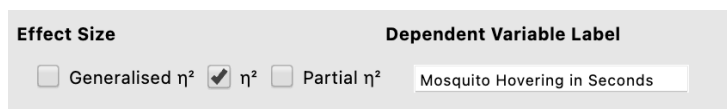
By clicking in each line you can edit and add in the variable name and its levels. Label the RM Factor 1 as Stinky Cheese Variety. Then click in each Level and type the four cheeses in. Note that once you hit the default last level (Level 3), hitting return will create an additional slot to type in our fourth cheese.

You'll notice that the four cheese varieties now appear down the right hand side of the Repeated Measures Cells box. The next stage is to drag the variables across to these slots.

When you have completed the variable set up the Repeated Measures ANOVA variable specification should look like this:



At the bottom of this menu we can also give our Dependent Variable a label to neaten our output, and ask for effect sizes while we are at it.



We now have some initial ANOVA output to interpret.

Repeated Measures ANOVA

Within Subjects Effects						
	Sum of Squares	df	Mean Square	F	p	η^2
Stinky Cheese Variety	893.67275	3	297.89092	17.84432	<.00001	0.52464
Residual	450.73475	27	16.69388			

Note. Type 3 Sums of Squares

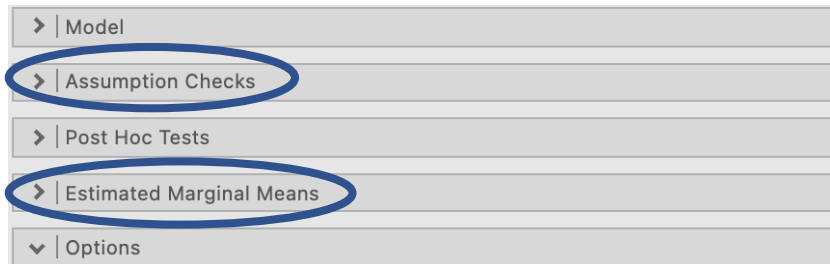
Our p value here is <.00001. This value is less than .05 so we will reject the null hypothesis that there is no difference in the number of seconds mosquitoes hover within 1cm of each stinky cheese variety.

In our ANOVA table we have the components of the calculations that help us arrive at our F statistic (namely the SS_{treat} and SS_{error} and their associated dfs , which lead us to our MS_{treat} which we divide by the MS_{error} to get our F statistic.

And hidden on the end is our η^2 effect size.

There are a couple of extra things we need to ask for from some of the drop down menus in our one-way repeated ANOVA. These are found under the Assumption Checks menu and the Estimated Marginal Means menu.

NB – We are not going to ask for our post hoc pairwise comparisons here as a pooled error term is used for all comparisons. It is preferable to use error terms that relate only to the two levels being compared. We will get to this soon.



Under Assumption Checks we are going to ask for “Sphericity tests” as well as a “Greenhouse-Geisser correction” under Sphericity corrections. This will enable us to check whether the sphericity assumption has been breached and to report corrected/adjusted results if a breach is present.

Assumptions

Tests of Sphericity				
	Mauchly's W	p	Greenhouse-Geisser ϵ	Huynh-Feldt ϵ
Stinky Cheese Variety	0.65375	0.65890	0.79939	1.00000

Repeated Measures ANOVA

Within Subjects Effects							
	Sphericity Correction	Sum of Squares	df	Mean Square	F	p	η^2
Stinky Cheese Variety	None	893.67275	3	297.89092	17.84432	<.00001	0.52464
	Greenhouse-Geisser	893.67275	2.39817	372.64847	17.84432	0.00001	0.52464
Residual	None	450.73475	27	16.69388			
	Greenhouse-Geisser	450.73475	21.58349	20.88331			

Note. Type 3 Sums of Squares

Our output now contains Mauchly's test of sphericity, and our ANOVA table now includes results with a Greenhouse-Geisser correction if needed

In our case the p value for Mauchly's test of sphericity is greater than .05 so we can say the sphericity assumption has not been breached. This means we can move forward with uncorrected results.

N.B., Some argue that Greenhouse-Geisser corrected results should always be reported to be on the safe side.

Under the Estimated Marginal Means drop down menu we can ask for our plot to help illustrate our findings.

Estimated Marginal Means

Stinky Cheese Variety

Marginal Means

Term 1

Stinky Cheese Variety

+ Add New Term

Output

☒ Marginal means plots

☐ Marginal means tables

General Options

☒ Equal cell weights

Confidence interval 95 %

Plot

Error bars Confidence interval

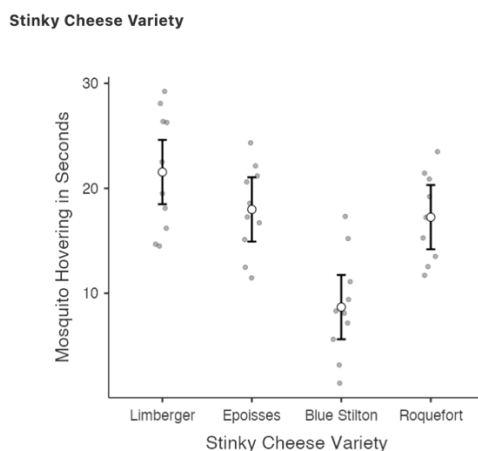
☒ Observed scores

We need to move our repeated measures factor Stinky Cheese Variety across to the Term 1 slot under the Marginal Means heading.

Marginal means plot with 95% confidence interval error bars is selected as a default.

We'll also tick observed scores so that we can see where the individual data points fall in relation to the means and error bars.

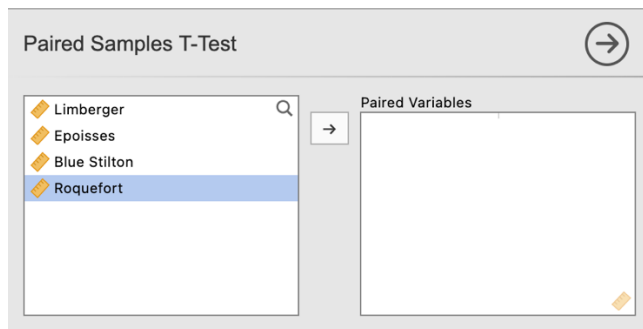
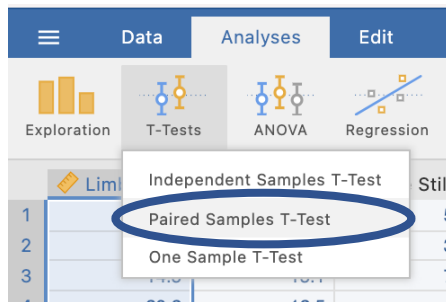
Estimated Marginal Means



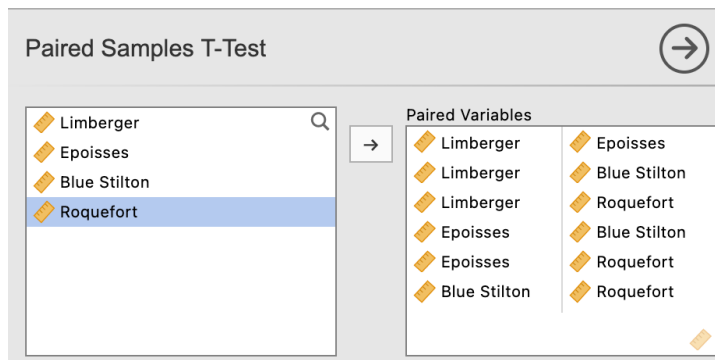
We can see visually here that it seems the mosquitoes most likely are hovering around the Blue Stilton for significantly fewer seconds than the other stinky cheeses. However we need to run our pairwise comparisons to obtain these significance tests.

Step 4a – Selecting analysis options to get the output we need for our post hoc pairwise comparisons following our significant omnibus result

While it is possible to ask for these pairwise comparisons within the Repeated Measures ANOVA menu, the comparisons that are generated use a pooled error term (across all four of our stinky cheese conditions). However, it is preferable to conduct these pairwise comparisons with error terms that are generated from the two levels being compared. In order to obtain these we will need to run a series of pairwise *t*-tests.



We can specify multiple paired *t*-tests at once by moving pairs over to the Paired Variables box. We want to create each combination of cheese comparisons to give us the set of comparisons we need.



Here we have specified each possible combination of comparisons. Depending on your specific research question or hypothesis you may not need to consider every combination but they have been asked for for completeness here.

In the analysis options for the paired *t*-tests output we'll select Mean difference (to illustrate a learning point only – you don't need to select this otherwise), effect sizes and associated confidence intervals and descriptives under the "Additional Statistics" heading.

Additional Statistics

☒ Mean difference

☐ Confidence interval 95 %

☒ Effect size

☒ Confidence interval 95 %

☒ Descriptives

☐ Descriptives plots

These selections will return the following output:

Paired Samples T-Test

Paired Samples T-Test

paired samples t-test										95% Confidence Interval	
			statistic	df	p	Mean difference	SE difference	Effect Size	Lower	Upper	
Limberger	Epoisses	Student's t	1.75791	9.00000	0.11263	3.56000	2.02513	Cohen's d	0.55590	-0.12654	1.21236
	Blue Stilton	Student's t	5.95141	9.00000	0.00021	12.87000	2.16251	Cohen's d	1.88200	0.80914	2.92113
	Roquefort	Student's t	2.52350	9.00000	0.03258	4.30000	1.70398	Cohen's d	0.79800	0.06351	1.50021
Epoisses	Blue Stilton	Student's t	6.83114	9.00000	0.00008	9.31000	1.36288	Cohen's d	2.16020	0.98413	3.30482
	Roquefort	Student's t	0.38622	9.00000	0.70831	0.74000	1.91603	Cohen's d	0.12213	-0.50347	0.74110
Blue Stilton		Student's t	-5.10074	9.00000	0.00064	-8.57000	1.68015	Cohen's d	-1.61300	-2.55522	-0.63526

Descriptives

	N	Mean	Median	SD	SE
Limberger	10	21.55000	21.00000	5.67416	1.79433
Epoisses	10	17.99000	17.95000	4.17598	1.32056
Limberger	10	21.55000	21.00000	5.67416	1.79433
Blue Stilton	10	8.68000	8.20000	4.92585	1.55769
Limberger	10	21.55000	21.00000	5.67416	1.79433
Roquefort	10	17.25000	17.25000	4.00895	1.26774
Epoisses	10	17.99000	17.95000	4.17598	1.32056
Blue Stilton	10	8.68000	8.20000	4.92585	1.55769
Epoisses	10	17.99000	17.95000	4.17598	1.32056
Roquefort	10	17.25000	17.25000	4.00895	1.26774
Blue Stilton	10	8.68000	8.20000	4.92585	1.55769
Roquefort	10	17.25000	17.25000	4.00895	1.26774

Note that having run these comparisons as separate *t*-tests the standard errors associated with each comparison are different. They are based on errors derived from each pairing rather than the whole model.

Step 5a – Finding the components for reporting the omnibus results

We've now run all the things we need to write up our one-way repeated measures ANOVA results, complete with post hoc pairwise comparisons. Let's pull it all together.

Firstly, let's report our omnibus results.

The components we obtain here are:

1. The F statistic, dfs and p value – the omnibus ANOVA result
2. An effect size in the form of η^2

Repeated Measures ANOVA

Within Subjects Effects						
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Note. Type 3 Sums of Squares

The Write Up (Part 1):

Ten mosquitoes were exposed to four types of Stinky Cheese (Limberger, Epoisses, Blue Stilton and Roquefort) for thirty seconds to determine their cheese preference. The number of seconds within each thirty second block that the mosquitoes hovered within one centimetre of the cheese was recorded by sensors. A one-way repeated measures ANOVA revealed that seconds spent hovering differed depending on the variety of cheese contained in the box, $F(3,27) = 17.84, p < .001, \eta^2 = .52$.

Tip: In APA format we report our p value to three decimal places. Where our p value would round to .000 we report this as $p < .001$ and not $p = .000$ which would imply our p value is equal to zero when it is not.

Step 5b – Finding the components for reporting the post hoc comparisons.

The next stage of the write-up is to present the post hoc comparisons results that reveal which particular differences in cheese hovering are significant. We'll use the paired *t*-test results to put this part of our write up together.

The elements needed for the post hoc section of our write up are:

1. **Post hoc comparison results** – to determine which group means are significant from each other. It is sufficient to report the *p* value for this.
2. **An effect size** for each post hoc comparison in the form of **Cohen's *d* and its associated confidence interval**.
3. **Means and standard deviations** – to help describe the pattern of these differences.

Paired Samples T-Test

Paired Samples T-Test

			statistic	df	p	Mean difference	SE difference		95% Confidence Interval		
								Effect Size	Lower	Upper	
Limberger	Epoisses	Student's t	1.75791	9.00000	0.11263	3.56000	2.02513	Cohen's d	0.55590	-0.12654	1.21236
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Roquefort	10	17.25000	17.25000	4.00895	1.26774

The continuation of the write up could go as follows:

The Write Up (Part 2):

Unadjusted post hoc comparisons revealed that mosquitoes spent significantly fewer seconds hovering over the Blue Stilton ($M = 8.68$, $SD = 4.93$) than the Limberger ($M = 21.55$, $SD = 5.67$, $p < .001$, $d = 1.88$, 95% CI [0.81, 2.92]), Epoisses ($M = 17.99$, $SD = 4.18$, $p < .001$, $d = 2.16$, 95% CI [0.98, 3.30]) and the Roquefort ($M = 17.25$, $SD = 4.01$, $p = .001$, $d = 1.61$, 95% CI [0.64, 2.56]). Further, mosquitoes were also seen to hover over the Roquefort cheese for significantly fewer seconds than the Limberger ($p = .033$, $d = 0.80$, 95% CI [0.06, 1.50]). No other differences were significant ($ps > .113$, $ds < 0.56$).

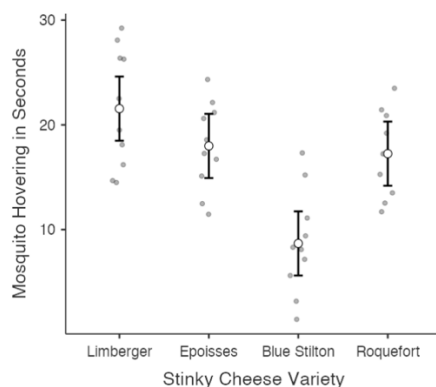
Potential addition of plot:

You could also add the plot we obtained to help illustrate the pattern of results. You might add a sentence like the following if you choose to include the plot:

Figure 1 below provides a visual demonstration of mosquito preference for Stinky Cheese Varieties.

Figure 1

Mosquito Cheese Hovering as a function of Stinky Cheese Variety



Note. Error bars represent 95% confidence intervals.

Created by Janine Lurie in consultation with the Statistics Working Group within the School of Psychology, University of Queensland ²

Based on *jamovi* v.1.8.4 ³

² The Statistics Working Group was formed in November 2020 to review the use of statistical packages in teaching across the core undergraduate statistics unit. The working group is led by Winnifred Louis and Philip Grove, with contributions from Timothy Ballard, Stefanie Becker, Jo Brown, Jenny Burt, Nathan Evans, Mark Horswill, David Sewell, Eric Vanman, Bill von Hippel, Courtney von Hippel, Zoe Walter, and Brendan Zietsch.

³ The jamovi project (2021). jamovi (Version 1.8.4) [Computer Software]. Retrieved from <https://www.jamovi.org>