## Results

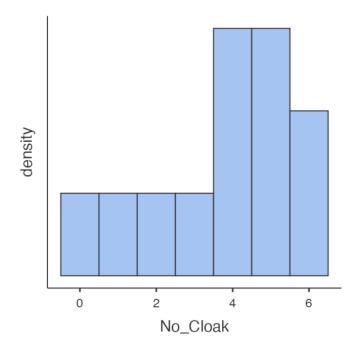
# **Descriptives**

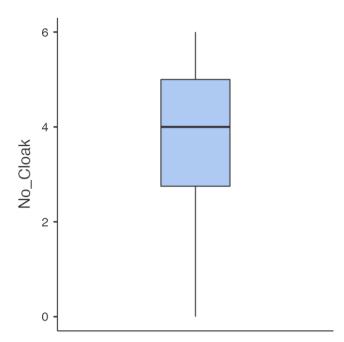
### Descriptives

	No_Cloak	Cloak
N	12	12
Missing	0	0
Mean	3.75	5.00
Median	4.00	5.00
Standard deviation	1.91	1.65
Minimum	0.00	2.00
Maximum	6.00	8.00
Skewness	-0.789	0.00
Std. error skewness	0.637	0.637
Kurtosis	-0.229	0.161
Std. error kurtosis	1.23	1.23
Shapiro-Wilk W	0.913	0.973
Shapiro-Wilk p	0.231	0.936

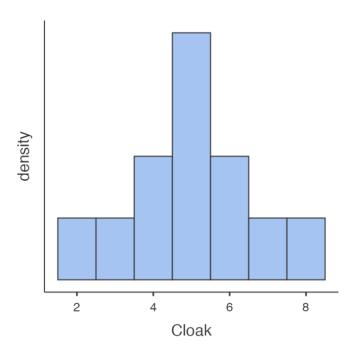
### Plots

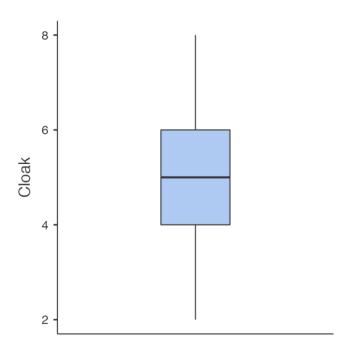
### No\_Cloak





## Cloak





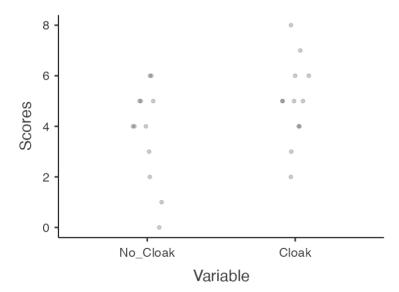
### **Repeated Measurements**

You have entered two related numeric variables. Hence, the <u>paired sample t test</u> seems to be a good option for you! In order to run this test in jamovi, go to: T-Tests > Paired Samples T-Test

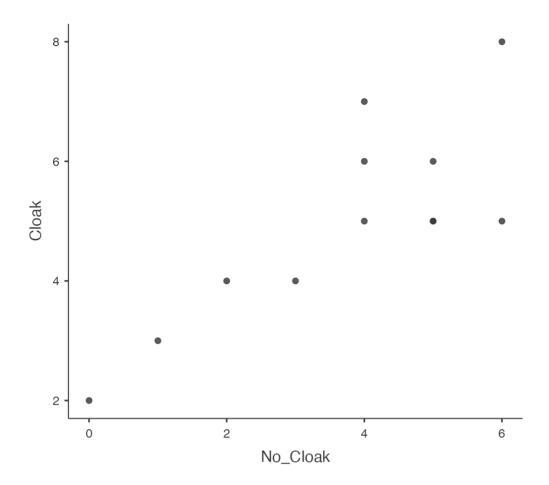
- Drop the two paired variables in the box below Paired Variables, one on the left side of the vertical line and one on the right side of the vertical line
- Under Hypothesis, select your alternative hypothesis

If the normality assumption is violated, you could use the non-parametric Wilcoxon signed rank test. Click on the links to learn more about these tests!

#### **Scatter Plot**



### **Scatterplot**



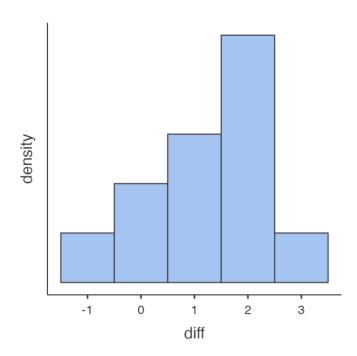
# **Descriptives**

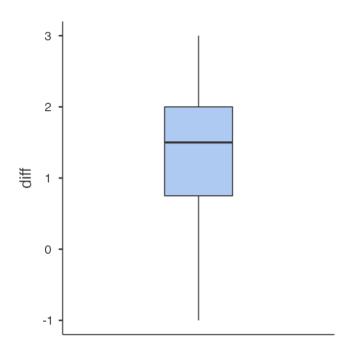
Descriptives

	diff
N	12
Missing	0
Mean	1.25
Median	1.50
Standard deviation	1.14
Minimum	-1.00
Maximum	3.00
Skewness	-0.583
Std. error skewness	0.637
Kurtosis	-0.138
Std. error kurtosis	1.23
Shapiro-Wilk W	0.912
Shapiro-Wilk p	0.228

### Plots

diff





# **Paired Samples T-Test**

Paired Samples T-Test

								Confi	5% dence rval		
			statistic	df	р	Mean difference	SE difference	Lower	Upper	-	Effect Size
No_Cloak	Cloak	Student's t	-3.80	11.0	0.003	-1.25	0.329	-1.97	-0.527	Cohen's d	-1.10

Note.  $H_a \mu_{Measure 1}$  - Measure 2  $\neq$  0

#### Normality Test (Shapiro-Wilk)

			W	р
No_Cloak	-	Cloak	0.912	0.228

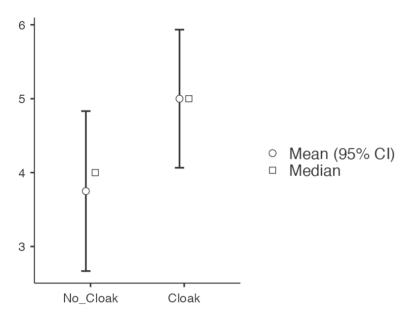
Note. A low p-value suggests a violation of the assumption of normality

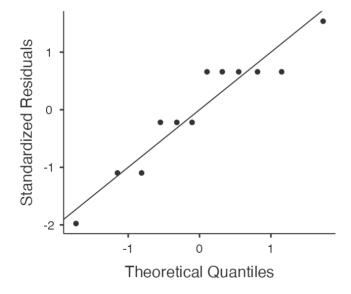
#### Descriptives

	N	Mean	Median	SD	SE
No_Cloak	12	3.75	4.00	1.91	0.552
Cloak	12	5.00	5.00	1.65	0.477

#### **Plots**

#### No\_Cloak - Cloak





## **Robust Paired Samples T-Test**

							95% Confide		
		t	df	р	Mean difference	SE	Lower	Upper	Cohen's d
No_Cloak	Cloak	-2.70	7.00	0.031	-1.00	0.370	-1.87	-0.125	0.398

## **Bayesian Paired Samples T-Test**

Bayesian Paired Samples T-Test

			BF <sub>10</sub>	error %
No_Cloak	-	Cloak	16.3	4.03e-6

[3] [4] [5]

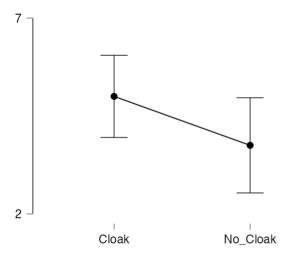
### **Descriptives**

#### Descriptives

					95% Credible Interval	
	N	Mean	SD	SE	Lower	Upper
No_Cloak Cloak	12 12	3.75 5.00	1.91 1.65	0.552 0.477	2.53 3.95	4.97 6.05

#### **Descriptives Plot**

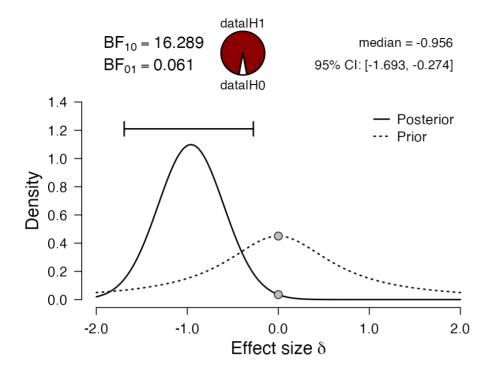
No\_Cloak - Cloak



#### **Inferential Plots**

No\_Cloak - Cloak

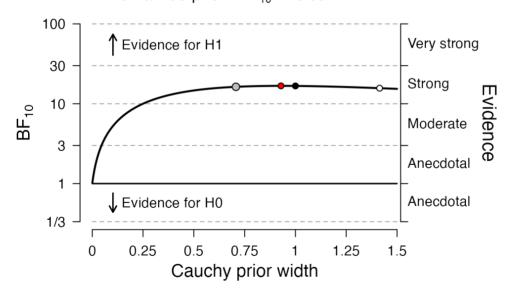
**Prior and Posterior** 



#### **Bayes Factor Robustness Check**

• max BF<sub>10</sub>: 16.738 at r = 0.9288

• wide prior:  $BF_{10} = 16.705$ • user prior:  $BF_{10} = 16.289$ • ultrawide prior:  $BF_{10} = 15.664$ 



#### References

[1] The jamovi project (2022). jamovi. (Version 2.3) [Computer Software]. Retrieved from https://www.jamovi.org.

[2] R Core Team (2021). R: A Language and environment for statistical computing. (Version 4.1) [Computer software]. Retrieved from <a href="https://cran.r-project.org">https://cran.r-project.org</a>. (R packages retrieved from MRAN snapshot 2022-01-01).

[3] JASP Team (2018). JASP. [Computer software]. Retrieved from https://jasp-stats.org.

[4] Morey, R. D., & Rouder, J. N. (2018). *BayesFactor: Computation of Bayes Factors for Common Designs*. [R package]. Retrieved from <a href="https://cran.r-project.org/package=BayesFactor">https://cran.r-project.org/package=BayesFactor</a>.

[5] Rouder, J. N., Speckman, P. L., Sun, D., Morey, R. D., & Iverson, G. (2009). Bayesian t tests for accepting and rejecting the null hypothesis. *Psychonomic Bulletin & Review, 16*, 225-237.