

Supplementary material 6 – Complete PCA, Kruskal-Wallis Test and Dunn's test results for Study 2

Complete results for Purchase attitudes and Willingness to buy

Principal Components Analysis

Call: principal(r = data_standardized_attitudes, nfactors = 1, scores = TRUE)

Standardized loadings (pattern matrix) based upon correlation matrix

	PC1	h2	u2	com
PA1	0.77	0.60	0.40	1
PA2	0.84	0.70	0.30	1
PA3	0.82	0.67	0.33	1
PA4	0.70	0.49	0.51	1
PA5	0.75	0.56	0.44	1

PC1

SS loadings 3.01

Proportion Var 0.60

Mean item complexity = 1

Test of the hypothesis that 1 component is sufficient.

The root mean square of the residuals (RMSR) is 0.1

with the empirical chi square 109.93 with prob < 4.2e-22

Fit based upon off diagonal values = 0.96> pca_willingness

Principal Components Analysis

Call: principal(r = data_standardized_willingness, nfactors = 1, scores = TRUE)

Standardized loadings (pattern matrix) based upon correlation matrix

	PC1	h2	u2	com
WTB1	0.89	0.80	0.20	1
WTB2	0.91	0.84	0.16	1
WTB3	0.91	0.83	0.17	1

PC1
SS loadings 2.46
Proportion Var 0.82

Mean item complexity = 1
Test of the hypothesis that 1 component is sufficient.

The root mean square of the residuals (RMSR) is 0.09
with the empirical chi square 25.42 with prob < NA

Fit based upon off diagonal values = 0.98>
> data\$PC_Attitudes <- pca_attitudes\$scores
> data\$PC_Willingness <- pca_willingness\$scores
> data\$PCAt <- as.numeric(data\$PC_Attitudes)
> data\$PCWill <- as.numeric(data\$PC_Willingness)
>
> # Kruskal-Wallis Test
> kw_test_attitudes <- kruskal.test(PCAt ~ Pub, data = data)
> kw_test_willingness <- kruskal.test(PCWill ~ Pub, data = data)
>
> # Convert Pub to a factor if it's not already
> data\$Pub <- as.factor(data\$Pub)

```
>
> # Dunn's test for attitudes
> dunn_test_attitudes <- dunn.test(x = data$PCAt, g = data$Pub, method = "bonferroni")
Kruskal-Wallis rank sum test
```

data: x and group

Kruskal-Wallis chi-squared = 21.3127, df = 4, p-value = 0

Comparison of x by group (Bonferroni)				
Col Mean-				
Row Mean	Alt	Bio	Control	Ego
-----+-----				
Bio	1.931282			
	0.2672			
Control	4.339686	2.331670		
	0.0001*	0.0986		
Ego	2.858277	0.892723	-1.445608	
	0.0213*	1.0000	0.7414	
Hed	3.348065	1.298132	-1.123112	0.375738
	0.0041*	0.9712	1.0000	1.0000

alpha = 0.05

Reject Ho if $p \leq \alpha/2$

>

> # Dunn's test for willingness

> dunn_test_willingness <- dunn.test(x = data\$PCWill, g = data\$Pub, method = "bonferroni")

Kruskal-Wallis rank sum test

data: x and group

Kruskal-Wallis chi-squared = 15.2108, df = 4, p-value = 0

Comparison of x by group (Bonferroni)

Col Mean-

Row Mean | Alt Bio Control Ego

-----+-----

Bio | 2.289425

| 0.1103

|

Control | 3.485362 1.130809

| 0.0025* 1.0000

|

Ego | 2.082712 -0.227398 -1.374926

| 0.1864 1.0000 0.8458

|

Hed | 3.291103 0.872184 -0.296714 1.120726

| 0.0050* 1.0000 1.0000 1.0000

alpha = 0.05

Reject H_0 if $p \leq \alpha/2$

>

> # Print the results

> print(dunn_test_attitudes)

\$chi2

[1] 21.31274

\$Z

[1] 1.9312823 4.3396862 2.3316700 2.8582771 0.8927230

[6] -1.4456082 3.3480656 1.2981323 -1.1231122 0.3757385

\$P

[1] 2.672408e-02 7.134317e-06 9.859029e-03 2.129741e-03

[5] 1.860028e-01 7.414356e-02 4.068887e-04 9.712094e-02

[9] 1.306949e-01 3.535557e-01

\$P.adjusted

[1] 2.672408e-01 7.134317e-05 9.859029e-02 2.129741e-02

[5] 1.000000e+00 7.414356e-01 4.068887e-03 9.712094e-01

[9] 1.000000e+00 1.000000e+00

\$comparisons

[1] "Alt - Bio" "Alt - Control" "Bio - Control"

[4] "Alt - Ego" "Bio - Ego" "Control - Ego"

[7] "Alt - Hed" "Bio - Hed" "Control - Hed"

[10] "Ego - Hed"

> print(dunn_test_willingness)

\$chi2

[1] 15.21077

\$Z

[1] 2.2894259 3.4853623 1.1308094 2.0827128 -0.2273980

[6] -1.3749270 3.2911035 0.8721842 -0.2967149 1.1207261

\$P

[1] 0.0110273087 0.0002457356 0.1290676690 0.0186387028

[5] 0.4100571280 0.0845770428 0.0004989760 0.1915539560

[9] 0.3833420732 0.1312022418

\$P.adjusted

[1] 0.110273087 0.002457356 1.000000000 0.186387028 1.000000000

[6] 0.845770428 0.004989760 1.000000000 1.000000000 1.000000000

\$comparisons

[1] "Alt - Bio" "Alt - Control" "Bio - Control"

[4] "Alt - Ego" "Bio - Ego" "Control - Ego"

[7] "Alt - Hed" "Bio - Hed" "Control - Hed"

[10] "Ego - Hed"

Complete results for Consumption attitudes and Willingness to consume

Principal Components Analysis

Call: principal(r = data_standardized_attitudes2, nfactors = 1, scores = TRUE)

Standardized loadings (pattern matrix) based upon correlation matrix

	PC1	h2	u2	com
CA1	0.88	0.78	0.22	1
CA2	0.89	0.80	0.20	1
CA3	0.87	0.75	0.25	1
CA4	0.80	0.64	0.36	1
CA5	0.84	0.70	0.30	1

PC1

SS loadings 3.67

Proportion Var 0.73

Mean item complexity = 1

Test of the hypothesis that 1 component is sufficient.

The root mean square of the residuals (RMSR) is 0.07

with the empirical chi square 49.62 with prob < 1.7e-09

Fit based upon off diagonal values = 0.99> pca_willingness2

Principal Components Analysis

Call: principal(r = data_standardized_willingness2, nfactors = 1, scores = TRUE)

Standardized loadings (pattern matrix) based upon correlation matrix

	PC1	h2	u2	com
WTC1	0.92	0.85	0.15	1

WTC2 0.94 0.89 0.11 1

WTC3 0.94 0.87 0.13 1

PC1

SS loadings 2.61

Proportion Var 0.87

Mean item complexity = 1

Test of the hypothesis that 1 component is sufficient.

The root mean square of the residuals (RMSR) is 0.07
with the empirical chi square 13.72 with prob < NA

Fit based upon off diagonal values = 0.99>

```
> data$PC_Attitudes2 <- pca_attitudes2$scores
```

```
> data$PC_willingness2 <- pca_willingness2$scores
```

```
> data$PCAt2 <- as.numeric(data$PC_Attitudes2)
```

```
> data$PCWill2 <- as.numeric(data$PC_willingness2)
```

```
>
```

```
> # Kruskal-Wallis Test
```

```
> kw_test_attitudes2 <- kruskal.test(PCAt2 ~ Pub, data = data)
```

```
> kw_test_willingness2 <- kruskal.test(PCWill2 ~ Pub, data = data)
```

```
>
```

```
> kw_test_attitudes2
```

Kruskal-Wallis rank sum test

data: PCAt2 by Pub

Kruskal-Wallis chi-squared = 1.5515, df = 4, p-value =
0.8175

> kw_test_willingness2

Kruskal-Wallis rank sum test

data: PCWill2 by Pub

Kruskal-Wallis chi-squared = 1.1816, df = 4, p-value =
0.8811