

GET

```
FILE = 'C:\Users\hasim.capar\Desktop\Masaüstü\2021 Masaüstü\Askeri Hoca COVID-19 Aşı Karşıtlığı Çalışması\AŞI KARARI.sav'
```

```
DATASET NAME DataSet1 WINDOW = FRONT.
```

```
USE ALL.
```

```
COMPUTE filter_$ = (Hangi_aşı > = 2).
```

```
VARIABLE LABELS filter_$ 'Hangi_aşı > = 2 (FILTER)'
```

```
VALUE LABELS filter_$ 0 'Not Selected' 1 'Selected'.
```

```
FORMATS filter_$ (f1.0).
```

```
FILTER BY filter_$.
```

```
EXECUTE.
```

```
FREQUENCIES VARIABLES = filter_$
```

```
 /ORDER = ANALYSIS.
```

## Frequencies

Notes		
Output Created	15-FEB-2023 09:56:17	
Comments		
Input	Data	C:\Users\hasim.capar\Desktop\Masaüstü\2021 Masaüstü\Askeri Hoca COVID-19 Aşı Karşıtlığı Çalışması\AŞI KARARI.sav
	Active Dataset	DataSet1
	Filter	Hangi_aşı > = 2 (FILTER)
	Weight	< none >
	Split File	< none >
	N of Rows in Working Data File	51
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics are based on all cases with valid data.
Syntax		FREQUENCIES VARIABLES = filter_\$ /ORDER = ANALYSIS.
Resources	Processor Time	00:00:00,02
	Elapsed Time	00:00:00,01

[DataSet1] C:\Users\hasim.capar\Desktop\Masaüstü\2021 Masaüstü\Askeri Hoca COVID-19 Aşı Karşıtlığı Çalışması\AŞI KARARI.sav

Statistics		
Hangi_aşı > = 2 (FILTER)		
N	Valid	51
	Missing	0

Hangi_aşı > = 2 (FILTER)					
		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Sinovac	19	37.3	37.3	37.3
	BioNTech	24	47.1	47.1	84.3
	Sinovac+BioNTech	8	15.7	15.7	100.0
	Total	51	100.0	100.0	

ONEWAY Total\_Scale\_Score Lack\_of\_Confidence Risk BY filter\_\$

```
 /STATISTICS DESCRIPTIVES HOMOGENEITY
```

```
 /MISSING ANALYSIS
```

```
 /POSTHOC = BONFERRONI ALPHA(0.05).
```

## Oneway

Notes		
Output Created		15-FEB-2023 09:57:09
Comments		
Input	Data	C:\Users\hasim.capar\Desktop\Masaüstü\2021 Masaüstü\Askeri Hoca COVID-19 Aşı Karşıtlığı Çalışması\AŞI KARARI.sav
	Active Dataset	DataSet1
	Filter	Hangi_aşı > = 2 (FILTER)
	Weight	< none >
	Split File	< none >
	N of Rows in Working Data File	51
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each analysis are based on cases with no missing data for any variable in the analysis.
Syntax	ONEWAY Total_Scale_Score Lack_of_Confidence Risk BY filter_Ş	
	/STATISTICS DESCRIPTIVES HOMOGENEITY	
	/MISSING ANALYSIS	
	/POSTHOC = BONFERRONI ALPHA(0.05).	
Resources	Processor Time	00:00:00.00
	Elapsed Time	00:00:00.04

Descriptives								
		N	Mean	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
					Lower Bound	Upper Bound		
Total_Scale_Score	Sinovac	19	24.1053	1.28166	21.4126	26.7979	15.00	37.00
	BioNTech	24	26.3750	0.91448	24.4833	28.2667	20.00	38.00
	Sinovac+BioNTech	8	25.5000	1.30931	22.4040	28.5960	19.00	30.00
	Total	51	25.3922	0.67857	24.0292	26.7551	15.00	38.00
Lack_of_Confidence	Sinovac	19	18.4211	1.23619	15.8239	21.0182	10.00	31.00
	BioNTech	24	20.2083	0.78014	18.5945	21.8222	12.00	29.00
	Sinovac+BioNTech	8	18.6250	1.08459	16.0603	21.1897	14.00	22.00
	Total	51	19.2941	0.61440	18.0601	20.5282	10.00	31.00
Risk	Sinovac	19	5.6842	0.41922	4.8035	6.5650	2.00	8.00
	BioNTech	24	6.1667	0.40229	5.3345	6.9989	2.00	9.00
	Sinovac+BioNTech	8	6.8750	0.44068	5.8330	7.9170	5.00	8.00
	Total	51	6.0980	0.25706	5.5817	6.6144	2.00	9.00

Test of Homogeneity of Variances					
		Levene Statistic	df1	df2	Sig.
Total_Scale_Score	Based on Mean	1.205	2	48	0.309
	Based on Median	0.844	2	48	0.436
	Based on Median and with adjusted df	0.844	2	43.036	0.437
Lack_of_Confidence	Based on trimmed mean	1.253	2	48	0.295
	Based on Mean	1.989	2	48	0.148
	Based on Median	1.977	2	48	0.150
Risk	Based on Median and with adjusted df	1.977	2	43.479	0.151
	Based on trimmed mean	1.955	2	48	0.153
	Based on Mean	1.124	2	48	0.333
Risk	Based on Median	1.124	2	48	0.333
	Based on Median and with adjusted df	1.124	2	39.269	0.335
	Based on trimmed mean	1.181	2	48	0.316

ANOVA						
		Sum of Squares	df	Mean Square	F	Sig.
Total_Scale_Score	Between Groups	54.742	2	27.371	1.174	0.318
	Within Groups	1119.414	48	23.321		
	Total	1174.157	50			
Lack_of_Confidence	Between Groups	38.123	2	19.062	0.990	0.379
	Within Groups	924.465	48	19.260		
	Total	962.588	50			
Risk	Between Groups	8.196	2	4.098	1.227	0.302
	Within Groups	160.314	48	3.340		
	Total	168.510	50			

## Post Hoc Tests

Multiple Comparisons							
Bonferroni							
Dependent Variable	(I) Hangi_aşı > = 2 (FILTER)	(J) Hangi_aşı > = 2 (FILTER)	Mean Difference (I-J)	Std. Error	Sig.	95% Confidence Interval	
						Lower Bound	Upper Bound
Total_Scale_Score	Sinovac	BioNTech	-2.26974	1.48295	0.397	-5.9486	1.4091
		Sinovac+BioNTech	-1.39474	2.03533	1.000	-6.4440	3.6545
	BioNTech	Sinovac	2.26974	1.48295	0.397	-1.4091	5.9486
		Sinovac+BioNTech	0.87500	1.97151	1.000	-4.0159	5.7659
	Sinovac+BioNTech	Sinovac	1.39474	2.03533	1.000	-3.6545	6.4440
		BioNTech	-0.87500	1.97151	1.000	-5.7659	4.0159
Lack_of_Confidence	Sinovac	BioNTech	-1.78728	1.34765	0.573	-5.1305	1.5559
		Sinovac+BioNTech	-0.20395	1.84963	1.000	-4.7925	4.3846
	BioNTech	Sinovac	1.78728	1.34765	0.573	-1.5559	5.1305
		Sinovac+BioNTech	1.58333	1.79163	1.000	-2.8613	6.0280
	Sinovac+BioNTech	Sinovac	0.20395	1.84963	1.000	-4.3846	4.7925
		BioNTech	-1.58333	1.79163	1.000	-6.0280	2.8613
Risk	Sinovac	BioNTech	-0.48246	0.56120	1.000	-1.8747	0.9098
		Sinovac+BioNTech	-1.19079	0.77024	0.386	-3.1016	0.7200
	BioNTech	Sinovac	0.48246	0.56120	1.000	-0.9098	1.8747
		Sinovac+BioNTech	-0.70833	0.74609	1.000	-2.5592	1.1425
	Sinovac+BioNTech	Sinovac	1.19079	0.77024	0.386	-0.7200	3.1016
		BioNTech	0.70833	0.74609	1.000	-1.1425	2.5592