```
>Warning # 853 in column 23. Text: pt_PT
>The LOCALE subcommand of the SET command specifies a locale
>for which collation and translation are not available.

GET
    FILE='/Users/PauloGarcia/Desktop/blendingbox/Analysis/First Study/SPSS Files
/datasets/q10_anal_lab.sav'.

DATASET NAME DataSet1 WINDOW=FRONT.

EXAMINE VARIABLES=distance_expectedC1C2distance_HSV distance_LCh distance_CMY
K distance_RGB distance_Lab
    /PLOT BOXPLOT STEMLEAF NPPLOT
    /COMPARE GROUPS
    /STATISTICS DESCRIPTIVES
    /CINTERVAL 95
```

Explore

/NOTOTAL.

/MISSING LISTWISE

Notes

	Notes	
Output Created		21-SEP-2016 17:32:40
Comments		
Input	Data	/Users/PauloGarcia/Des ktop/blendingbox/Anal ysis/First Study/SPSS Files/datasets/q10_anal _lab.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	14
Missing Value Handling	Definition of Missing	User-defined missing values for dependent variables are treated as missing.
	Cases Used	Statistics are based on cases with no missing values for any dependent variable or factor used.

Notes

Syntax		EXAMINE VARIABLES=distance_ex pectedC1C2 distance_HSV distance_LCh distance_CMYK distance_RGB distance_Lab /PLOT BOXPLOT STEMLEAF NPPLOT /COMPARE GROUPS /STATISTICS DESCRIPTIVES /CINTERVAL 95 /MISSING LISTWISE /NOTOTAL.
Resources	Processor Time	00:00:04,20
	Elapsed Time	00:00:03,00

 $\label{lem:condition} $$[DataSet1] / Users/PauloGarcia/Desktop/blendingbox/Analysis/First Study/SPSS Files/datasets/q10_anal_lab.sav$

Case Processing Summary

		Cases				
	Va	alid	Mis	sing	To	otal
	N	Percent	N	Percent	N	Percent
distance_expected C1C2	14	100,0%	0	0,0%	14	100,0%
distance_HSV	14	100,0%	0	0,0%	14	100,0%
distance_LCh	14	100,0%	0	0,0%	14	100,0%
distance_CMYK	14	100,0%	0	0,0%	14	100,0%
distance_RGB	14	100,0%	0	0,0%	14	100,0%
distance_Lab	14	100,0%	0	0,0%	14	100,0%

Descriptives

			Statistic	Std. Error
distance_expected	Mean		.5671	.04055
C1C2	95% Confidence Lo	wer Bound	.4795	
	Interval for Mean Up	per Bound	.6547	
	5% Trimmed Mean	-	.5646	
	Median		.5950	
	Variance		,023	
	Std. Deviation		.15173	
	Minimum		.35	
	Maximum		.83	
	Range		.48	
	Interquartile Range		.27	
	Skewness		,131	,597
	Kurtosis		-1,052	1,154
distance_HSV	Mean		.3000	.04221
		wer Bound	.2088	
	Interval for Mean Up	per Bound	.3912	
	5% Trimmed Mean		.3083	
	Median		.3950	
	Variance		,025	
	Std. Deviation		.15792	
	Minimum		.02	
	Maximum		.43	
	Range		.41	
	Interquartile Range		.30	
	Skewness		-,776	,597
	Kurtosis		-1,385	1,154
distance_LCh	Mean		.2557	.03111
		wer Bound	.1885	
	Interval for Mean Up	per Bound	.3229	
	5% Trimmed Mean		.2591	
	Median		.2750	
	Variance		,014	
	Std. Deviation		.11640	
	Minimum		.05	
	Maximum		.40	
	Range		.35	
	Interquartile Range		.16	
	Skewness		-,703	,597
	Kurtosis		-,416	1,154
distance_CMYK	Mean		.1314	.01275
		wer Bound	.1039	
	Interval for Mean Up	per Bound	.1590	
	5% Trimmed Mean		.1310	
	Median		.1200	

Descriptives

			Statistic	Std. Error
	Variance		,002	
	Std. Deviation		.04769	
	Minimum		.06	
	Maximum		.21	
	Range		.15	
	Interquartile Range		.08	
	Skewness		,393	,597
	Kurtosis		-1,130	1,154
distance_RGB	Mean		.2107	.01711
	95% Confidence	Lower Bound	.1737	
	Interval for Mean	Upper Bound	.2477	
	5% Trimmed Mean		.2086	
	Median		.2000	
	Variance		,004	
	Std. Deviation		.06403	
	Minimum		.12	
	Maximum		.34	
	Range		.22	
	Interquartile Range		.07	
	Skewness		,936	,597
	Kurtosis		,194	1,154
distance_Lab	Mean		.2043	.02561
	95% Confidence	Lower Bound	.1490	
	Interval for Mean	Upper Bound	.2596	
	5% Trimmed Mean		.2020	
	Median		.1750	
	Variance		,009	
	Std. Deviation		.09581	
	Minimum		.07	
	Maximum		.38	
	Range		.31	
	Interquartile Range		.16	
	Skewness		,784	,597
	Kurtosis		-,599	1,154

Tests of Normality

	Kolmogorov-Smirnov ^a			Shapiro-Wilk		
	Statistic	df	Sig.	Statistic	df	Sig.
distance_expected C1C2	,137	14	,200 [*]	,955	14	,633
distance_HSV	,358	14	,000	,747	14	,001
distance_LCh	,195	14	,157	,890	14	,081
distance_CMYK	,238	14	,031	,923	14	,241
distance_RGB	,228	14	,047	,898	14	,106
distance_Lab	,274	14	,006	,878	14	,054

- *. This is a lower bound of the true significance.
- a. Lilliefors Significance Correction

NPAR TESTS

 $/ {\tt FRIEDMAN\!=} distance_expected C1C2 distance_HSV \ distance_LCh \ distance_CMYK \ distance_RGB \ distance_Lab$

/STATISTICS DESCRIPTIVES QUARTILES

/MISSING LISTWISE.

NPar Tests

Notes

		04 055 0040 45 00 54
Output Created		21-SEP-2016 17:33:54
Comments		
Input	Data	/Users/PauloGarcia/Des ktop/blendingbox/Anal ysis/First Study/SPSS Files/datasets/q10_anal _lab.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	14
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for all tests are based on cases with no missing data for any variables used.

Notes

Syntax		NPAR TESTS
		/FRIEDMAN=distance_e xpectedC1C2 distance_HSV distance_LCh distance_CMYK distance_RGB distance_Lab /STATISTICS DESCRIPTIVES QUARTILES /MISSING LISTWISE.
Resources	Processor Time	00:00:00,01
	Elapsed Time	00:00:00,00
	Number of Cases Allowed ^a	71493

a. Based on availability of workspace memory.

Descriptive Statistics

						Percentile
	N	Mean	Std. Deviation	Minimum	Maximum	25th
distance_expected C1C2	14	.5671	.15173	.35	.83	.4300
distance_HSV	14	.3000	.15792	.02	.43	.1200
distance_LCh	14	.2557	.11640	.05	.40	.1875
distance_CMYK	14	.1314	.04769	.06	.21	.0975
distance_RGB	14	.2107	.06403	.12	.34	.1675
distance_Lab	14	.2043	.09581	.07	.38	.1375

Descriptive Statistics

	Percentiles		
	50th (Median)	75th	
distance_expected C1C2	.5950	.7000	
distance_HSV	.3950	.4200	
distance_LCh	.2750	.3450	
distance_CMYK	.1200	.1750	
distance_RGB	.2000	.2375	
distance_Lab	.1750	.3025	

Friedman Test

Ranks

	Mean Rank
distance_expected C1C2	5,68
distance_HSV	3,89
distance_LCh	3,54
distance_CMYK	1,43
distance_RGB	3,36
distance_Lab	3,11

Test Statistics^a

N	14
Chi-Square	37,700
df	5
Asymp. Sig.	,000

a. Friedman Test

NPAR TESTS

/WILCOXON=distance_HSV distance_HSV distance_HSV distance_LCh d istance_LCh distance_LCh distance_CMYK distance_RGB WITH distance_LCh distance_CMYK distance_RGB distance_Lab distance_CMYK distance_RGB distance_Lab distance_RGB distance_RGB distance_Lab distance_RGB distance_RGB distance_Lab distance_RGB distance_RGB

/STATISTICS DESCRIPTIVES QUARTILES /MISSING ANALYSIS.

NPar Tests

Notes

Comments		
001111101110		
Input D	ata	/Users/PauloGarcia/Des ktop/blendingbox/Anal ysis/First Study/SPSS Files/datasets/q10_anal _lab.sav
A	ctive Dataset	DataSet1
Fi	ilter	<none></none>
W	/eight	<none></none>
S	plit File	<none></none>
	of Rows in /orking Data File	14
	efinition of lissing	User-defined missing values are treated as missing.
С	ases Used	Statistics for each test are based on all cases with valid data for the variable(s) used in that test.
Syntax		NPAR TESTS
		/WILCOXON=distance_H SV distance_HSV distance_HSV distance_HSV distance_LCh distance_LCh distance_LCh distance_CMYK distance_CMYK distance_CMYK distance_CMYK distance_LCh distance_LAB dis
Resources P	rocessor Time	00:00:00,01
	lapsed Time	00:00:00,01
N	umber of Cases llowed ^a	78643

a. Based on availability of workspace memory.

Descriptive Statistics

						Percentile
	N	Mean	Std. Deviation	Minimum	Maximum	25th
distance_HSV	14	.3000	.15792	.02	.43	.1200
distance_LCh	14	.2557	.11640	.05	.40	.1875
distance_CMYK	14	.1314	.04769	.06	.21	.0975
distance_RGB	14	.2107	.06403	.12	.34	.1675
distance_Lab	14	.2043	.09581	.07	.38	.1375

Descriptive Statistics

	Percentiles			
	50th (Median)	75th		
distance_HSV	.3950	.4200		
distance_LCh	.2750	.3450		
distance_CMYK	.1200	.1750		
distance_RGB	.2000	.2375		
distance_Lab	.1750	.3025		

Wilcoxon Signed Ranks Test

Ranks

N Mean Rank Sum of Ranks					
Distance_HSV			N	Mean Rank	Sum of Ranks
Ties	<u> </u>	Negative Ranks	10 ^a	6,40	64,00
Total	distance_HSV	Positive Ranks	4 ^b	10,25	41,00
distance_CMYK - distance_HSV Negative Ranks Positive Ranks Ties Of Total 10 ^d 3,38 13,50 distance_RGB - distance_HSV Negative Ranks Positive Ranks Ties Oi Total 9 ^g 9,06 81,50 distance_HSV Positive Ranks Ties Oi Total 14 distance_Lab - distance_HSV Negative Ranks Positive Ranks Ties Oi Total 9 ^j 8,06 72,50 32,50 distance_HSV Negative Ranks Ties Oi Total 14 distance_CMYK - distance_LCh Negative Ranks Ties Oi Total 12 ^m 8,13 97,50 7,50 distance_LCh Positive Ranks Ties Oo Total 2 ⁿ 3,75 7,50 Ties Total 14 distance_RGB - distance_LCh Negative Ranks Ties Oo Total Adistance_RGB - distance_LCh Negative Ranks Ties Oo Total Adistance_RGB - distance_LCh Negative Ranks Ties Oo Total Adistance_LCh Negative Ranks Ties Oo Total Adistance_LCh Negative Ranks Ties Oo Total		Ties	0 c		
Desitive Ranks Continue Con		Total	14		
Ties 0 f Total 14 distance_RGB - Negative Ranks Positive Ranks Ties 0 i Total 14 distance_Lab - Dositive Ranks Positive Ranks Positive Ranks Ties 0 i Total 14 distance_Lab - Dositive Ranks Positive Ranks Ties 0 i Total 14 distance_Lab - Dositive Ranks Ties 0 i Total 14 distance_CMYK - Negative Ranks 12 i 8,13 97,50 Positive Ranks 2 i 3,75 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i 7,50 Ties 0 i		Negative Ranks	10 ^d	9,15	91,50
Total	distance_HSV	Positive Ranks	4 ^e	3,38	13,50
distance_RGB - distance_HSV Negative Ranks Positive Ranks 9g bh 9,06 bh 81,50 bh Ties Total 0i Total 14 23,50 bh distance_Lab - distance_HSV Negative Ranks Positive Ranks Ties Total 9j bh 8,06 bh 72,50 bh Ties Total 0i Total 14 14 14 distance_CMYK - distance_LCh Negative Ranks Positive Ranks Ties Total 12m bh 8,13 bh 97,50 bh Ties Total 0° Total 3,75 bh 7,50 bh distance_RGB - Total Negative Ranks Bh 9p bh 7,39 bh 66,50 bh		Ties	0 ^f		
distance_HSV Positive Ranks 5h 4,70 23,50 Ties 0i 14 23,50 distance_Lab - distance_HSV Negative Ranks positive Ranks positive Ranks Ties 5k positive Ranks positive Ranks positive Ranks distance_CMYK positive Ranks positive Ranks positive Ranks Ties 12m positive Ranks positive Rank		Total	14		
Ties 0 i		Negative Ranks	9 g	9,06	81,50
Total	distance_HSV	Positive Ranks	5 ^h	4,70	23,50
distance_Lab - distance_HSV Negative Ranks Positive Ranks 9j		Ties	0 ⁱ		
Desitive Ranks Ties O		Total	14		
Ties 0 1 14		Negative Ranks	9 ^j	8,06	72,50
Total		Positive Ranks	5 ^k	6,50	32,50
distance_CMYK - distance_LCh Negative Ranks Positive Ranks Ranks Ties Positive Ranks R		Ties	0 ¹		
distance_LCh Positive Ranks 2 ⁿ 3,75 7,50 Ties 0°		Total	14		
Ties 0° Total 14 distance_RGB - Negative Ranks 9° 7,39 66,50		Negative Ranks	12 ^m	8,13	97,50
Total 14 distance_RGB - Negative Ranks 9 ^p 7,39 66,50	distance_LCh	Positive Ranks	2 ⁿ	3,75	7,50
distance_RGB - Negative Ranks 9 ^p 7,39 66,50		Ties	0°		
distance I Ch		Total	14		
distance_LCh Positive Ranks 4 ^q 6,13 24,50		Negative Ranks	9 ^p	7,39	66,50
	distance_LCh	Positive Ranks	4 ^q	6,13	24,50

Ranks

		N	Mean Rank	Sum of Ranks
	Ties	1 r		
	Total	14		
distance_Lab -	Negative Ranks	8 ^s	9,38	75,00
distance_LCh	Positive Ranks	6 ^t	5,00	30,00
	Ties	0 ^u		
	Total	14		
distance_RGB -	Negative Ranks	0 v	,00	,00
distance_CMYK	Positive Ranks	14 ^w	7,50	105,00
	Ties	0 ^x		
	Total	14		
distance_Lab -	Negative Ranks	0 y	,00	,00
distance_CMYK	Positive Ranks	14 ^z	7,50	105,00
	Ties	0 ^{aa}		
	Total	14		
distance_Lab -	Negative Ranks	9 ^{ab}	5,67	51,00
distance_RGB	Positive Ranks	4 ^{ac}	10,00	40,00
	Ties	1 ^{ad}		
	Total	14		

a. distance_LCh < distance_HSV

b. distance_LCh > distance_HSV

c. distance_LCh = distance_HSV

- d. distance_CMYK < distance_HSV
- e. distance_CMYK > distance_HSV
- f. distance CMYK = distance HSV
- g. distance_RGB < distance_HSV
- h. distance RGB > distance HSV
- i. distance_RGB = distance_HSV
- j. distance_Lab < distance_HSV
- k. distance_Lab > distance_HSV
- I. distance_Lab = distance_HSV
- m. distance CMYK < distance LCh
- n. distance_CMYK > distance_LCh
- o. distance_CMYK = distance_LCh
- p. distance_RGB < distance_LCh
- q. distance_RGB > distance_LCh
- r. distance_RGB = distance_LCh
- s. distance_Lab < distance_LCh
- t. distance_Lab > distance_LCh
- u. distance_Lab = distance_LCh
- v. distance_RGB < distance_CMYK
- w. distance_RGB > distance_CMYK
- x. distance_RGB = distance_CMYK
- y. distance_Lab < distance_CMYK
- z. distance_Lab > distance_CMYK
- aa. distance_Lab = distance_CMYK
- ab. distance_Lab < distance_RGB
- ac. distance_Lab > distance_RGB
- ad. distance_Lab = distance_RGB

Test Statistics^a

	distance_LCh - distance_HSV	distance_CMY K - distance_HSV	distance_RGB - distance_HSV	distance_Lab - distance_HSV	distance_CMY K - distance_LCh
Z	-,723 ^b	-2,451 ^b	-1,826 ^b	-1,260 ^b	-2,826 ^b
Asymp. Sig. (2-tailed)	,470	,014	,068	,208	,005

Test Statistics^a

	distance_RGB - distance_LCh	distance_Lab - distance_LCh	distance_RGB - distance_CMY K	distance_Lab - distance_CMY K	distance_Lab - distance_RGB
Z	-1,469 ^b	-1,414 ^b	-3,306 ^c	-3,301 ^c	-,395 ^b
Asymp. Sig. (2-tailed)	,142	,157	,001	,001	,693

- a. Wilcoxon Signed Ranks Test
- b. Based on positive ranks.
- c. Based on negative ranks.