

Lampiran 1

Uji Normalitas

One-Sample Kolmogorov-Smirnov Test

		pH_Saliva
N		21
Normal Parameters ^{a,b}	Mean	5.6305
	Std. Deviation	.38517
Most Extreme Differences	Absolute	.358
	Positive	.358
	Negative	-.225
Kolmogorov-Smirnov Z		.720
Asymp. Sig. (2-tailed)		.678

a. Test distribution is Normal.

b. Calculated from data.

Correlations

Correlations

		DK	pH_Saliva
DK	Pearson Correlation	1	-.995**
	Sig. (2-tailed)	.	.000
	N	21	21
pH_Saliva	Pearson Correlation	-.995**	1
	Sig. (2-tailed)	.000	.
	N	21	21

** . Correlation is significant at the 0.01 level (2-tailed).

Regression

Model Summary

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.995 ^a	.991	.990	.03825

a. Predictors: (Constant), DK

ANOVA^b

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	2.939	1	2.939	2008.674	.000 ^a
	Residual	.028	19	.001		
	Total	2.967	20			

a. Predictors: (Constant), DK

b. Dependent Variable: pH_Saliva

Coefficients^c

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	6.505	.021		306.408	.000
	DK	-.704	.016	-.995	-44.818	.000

a. Dependent Variable: pH_Saliva

Oneway

Descriptives

pH_Saliva

	N	Mean	Std. Deviation	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
					Lower Bound	Upper Bound		
					K Neg	3		
P 2	3	5.6200	.01000	.00577	5.5952	5.6448	5.61	5.63
P 3	3	5.5500	.01000	.00577	5.5252	5.5748	5.54	5.56
P 4	3	5.4900	.01000	.00577	5.4652	5.5148	5.48	5.50
P 5	3	5.4533	.01528	.00882	5.4154	5.4913	5.44	5.47
P 6	3	5.4067	.00577	.00333	5.3923	5.4210	5.40	5.41
P 7	3	5.3633	.02082	.01202	5.3116	5.4150	5.34	5.38
Total	21	5.6305	.38517	.08405	5.4551	5.8058	5.34	6.54

Test of Homogeneity of Variances

pH_Saliva

Levene Statistic	df 1	df 2	Sig.
1.179	6	14	.372

ANOVA

pH_Saliva

	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	2.965	6	.494	3144.586	.000
Within Groups	.002	14	.000		
Total	2.967	20			



Post Hoc Tests

Multiple Comparisons

Dependent Variable: pH_Saliva
Tukey HSD

(I) Kelompok	(J) Kelompok	Mean Difference (I-J)	Std. Error	Sig.	95% Confidence Interval	
					Lower Bound	Upper Bound
K Neg	P 2	.91000*	.01024	.000	.8751	.9449
	P 3	.98000*	.01024	.000	.9451	1.0149
	P 4	1.04000*	.01024	.000	1.0051	1.0749
	P 5	1.07667*	.01024	.000	1.0417	1.1116
	P 6	1.12333*	.01024	.000	1.0884	1.1583
	P 7	1.16667*	.01024	.000	1.1317	1.2016
	P 2	K Neg	-.91000*	.01024	.000	-.9449
P 3		.07000*	.01024	.000	.0351	.1049
P 4		.13000*	.01024	.000	.0951	.1649
P 5		.16667*	.01024	.000	.1317	.2016
P 6		.21333*	.01024	.000	.1784	.2483
P 7		.25667*	.01024	.000	.2217	.2916
P 3		K Neg	-.98000*	.01024	.000	-1.0149
	P 2	-.07000*	.01024	.000	-.1049	-.0351
	P 4	.06000*	.01024	.001	.0251	.0949
	P 5	.09667*	.01024	.000	.0617	.1316
	P 6	.14333*	.01024	.000	.1084	.1783
	P 7	.18667*	.01024	.000	.1517	.2216
	P 4	K Neg	-1.04000*	.01024	.000	-1.0749
P 2		-.13000*	.01024	.000	-.1649	-.0951
P 3		-.06000*	.01024	.001	-.0949	-.0251
P 5		.03667*	.01024	.037	.0017	.0716
P 6		.08333*	.01024	.000	.0484	.1183
P 7		.12667*	.01024	.000	.0917	.1616
P 5		K Neg	-1.07667*	.01024	.000	-1.1116
	P 2	-.16667*	.01024	.000	-.2016	-.1317
	P 3	-.09667*	.01024	.000	-.1316	-.0617
	P 4	-.03667*	.01024	.037	-.0716	-.0017
	P 6	.04667*	.01024	.006	.0117	.0816
	P 7	.09000*	.01024	.000	.0551	.1249
	P 6	K Neg	-1.12333*	.01024	.000	-1.1583
P 2		-.21333*	.01024	.000	-.2483	-.1784
P 3		-.14333*	.01024	.000	-.1783	-.1084
P 4		-.08333*	.01024	.000	-.1183	-.0484
P 5		-.04667*	.01024	.006	-.0816	-.0117
P 7		.04333*	.01024	.011	.0084	.0783
P 7		K Neg	-1.16667*	.01024	.000	-1.2016
	P 2	-.25667*	.01024	.000	-.2916	-.2217
	P 3	-.18667*	.01024	.000	-.2216	-.1517
	P 4	-.12667*	.01024	.000	-.1616	-.0917
	P 5	-.09000*	.01024	.000	-.1249	-.0551
	P 6	-.04333*	.01024	.011	-.0783	-.0084

*. The mean difference is significant at the .05 level.

Homogeneous Subsets

pH_Saliva

Tukey HSD

Kelompok	N	Subset for alpha = .05						
		1	2	3	4	5	6	7
P 7	3	5.3633						
P 6	3		5.4067					
P 5	3			5.4533				
P 4	3				5.4900			
P 3	3					5.5500		
P 2	3						5.6200	
K Neg	3							6.5300
Sig.		1.000	1.000	1.000	1.000	1.000	1.000	1.000

Means for groups in homogeneous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Means Plots

