

# Statistical Report

All patients diagnosed with HIV during the study period (n=1717)

The FREQ Procedure

sex				
sex	Frequency	Percent	Cumulative Frequency	Cumulative Percent
female	688	40.07	688	40.07
male	1029	59.93	1717	100.00

Frequency Missing = 57

Variable: age (age)

sex=male

Moments			
N	1018	Sum Weights	1018
Mean	33.7043222	Sum Observations	34311
Std Deviation	12.3134257	Variance	151.620453
Skewness	0.44102847	Kurtosis	0.14824806

Quantiles (Definition 5)	
Quantile	Estimate
100% Max	73
99%	65
95%	57
90%	50
75% Q3	42
50% Median	32
25% Q1	24
10%	20
5%	18
1%	0
0% Min	0

The UNIVARIATE Procedure

Variable: age (age)

sex=female

<b>Moments</b>			
<b>N</b>	678	<b>Sum Weights</b>	678
<b>Mean</b>	33.939528	<b>Sum Observations</b>	23011
<b>Std Deviation</b>	13.3768534	<b>Variance</b>	178.940208
<b>Skewness</b>	0.10574218	<b>Kurtosis</b>	0.17126393

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	81
<b>99%</b>	65
<b>95%</b>	57
<b>90%</b>	53
<b>75% Q3</b>	43
<b>50% Median</b>	33
<b>25% Q1</b>	25
<b>10%</b>	19
<b>5%</b>	16
<b>1%</b>	0
<b>0% Min</b>	0

The UNIVARIATE Procedure

The NPAR1WAY Procedure

<b>Wilcoxon Scores (Rank Sums) for Variable age Classified by Variable sex</b>					
<b>sex</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>female</b>	678	583619.50	575283.0	9876.14920	860.795723
<b>male</b>	1018	855436.50	863773.0	9876.14920	840.310904

Average scores were used for ties.

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	583619.5000
<b>Normal Approximation</b>	
<b>Z</b>	0.8441

<b>Wilcoxon Two-Sample Test</b>	
<b>One-Sided Pr &gt; Z</b>	0.1993
<b>Two-Sided Pr &gt;  Z </b>	0.3986

**All patients who were tested for HBV and HCV (n=1259)**

The SAS System

The FREQ Procedure

<b>HIV_HBV</b>				
<b>HIV_HBV</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>non_reactive</b>	1218	96.90	1218	96.90
<b>reagent</b>	39	3.10	1257	100.00

Frequency Missing = 2

<b>HIV_HCV</b>				
<b>HIV_HCV</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>non_reactive</b>	1218	96.90	1218	96.90
<b>reagent</b>	39	3.10	1257	100.00

Frequency Missing = 2

<b>HIV_HBV_HCV</b>				
<b>HIV_HBV_HCV</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>non_reactive</b>	1218	99.84	1218	99.84
<b>reagent</b>	2	0.16	1220	100.00

Frequency Missing = 2

<b>hepatitis</b>				
<b>hepatitis</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>non_reactive</b>	1179	93.65	1179	93.65
<b>reagent</b>	80	6.35	1259	100.00

The SAS System

The FREQ Procedure  
HIV-monoinfected\_and\_HIV\_hepatitis

<b>sex</b>				
<b>sex</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>female</b>	528	41.94	528	41.94
<b>male</b>	731	58.06	1259	100.00

The SAS System

The UNIVARIATE Procedure  
Variable: age (age)  
sex=male

<b>Moments</b>			
<b>N</b>	731	<b>Sum Weights</b>	731
<b>Mean</b>	33.1340629		
<b>Std Deviation</b>	12.1459651	<b>Variance</b>	147.524468
<b>Skewness</b>	0.47682548	<b>Kurtosis</b>	0.08956363

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	70
<b>99%</b>	63
<b>95%</b>	56
<b>90%</b>	50
<b>75% Q3</b>	41
<b>50% Median</b>	31
<b>25% Q1</b>	24
<b>10%</b>	20
<b>5%</b>	18
<b>1%</b>	0
<b>0% Min</b>	0

The SAS System

The UNIVARIATE Procedure

Variable: age (age)

sex=female

<b>Moments</b>			
<b>N</b>	524	<b>Sum Weights</b>	524
<b>Mean</b>	33.7232824		
<b>Std Deviation</b>	13.1700799	<b>Variance</b>	173.451006
<b>Skewness</b>	0.20688551	<b>Kurtosis</b>	0.25039397

**Quantiles (Definition 5)**

<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	81
<b>99%</b>	65
<b>95%</b>	56
<b>90%</b>	53
<b>75% Q3</b>	42
<b>50% Median</b>	32
<b>25% Q1</b>	25
<b>10%</b>	19
<b>5%</b>	17
<b>1%</b>	0
<b>0% Min</b>	0

The SAS System

The NPAR1WAY Procedure

**Wilcoxon Scores (Rank Sums) for Variable age  
Classified by Variable sex**

<b>sex</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>female</b>	524	336564.0	329072.0	6329.47918	642.297710
<b>male</b>	731	451576.0	459068.0	6329.47918	617.751026

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	336564.0000
<b>Z</b>	1.1836
<b>One-Sided Pr &gt; Z</b>	0.1183
<b>Two-Sided Pr &gt;  Z </b>	0.2366

The SAS System

The UNIVARIATE Procedure  
 Variable: age (age)  
 hepatitis=reagent sex=male

<b>Moments</b>			
<b>N</b>	51	<b>Sum Weights</b>	51
<b>Mean</b>	35.8823529	<b>Sum Observations</b>	1830
<b>Std Deviation</b>	10.8658125	<b>Variance</b>	118.065882
<b>Skewness</b>	0.1261037	<b>Kurtosis</b>	-0.6400289

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	58
<b>99%</b>	58
<b>95%</b>	54
<b>90%</b>	52
<b>75% Q3</b>	44
<b>50% Median</b>	35
<b>25% Q1</b>	28
<b>10%</b>	23
<b>5%</b>	18
<b>1%</b>	14
<b>0% Min</b>	14

The SAS System

The UNIVARIATE Procedure  
 Variable: age (age)  
 hepatitis=reagent sex=female

<b>Moments</b>			
<b>N</b>	29	<b>Sum Weights</b>	29
<b>Mean</b>	32.7931034	<b>Sum Observations</b>	951
<b>Std Deviation</b>	10.523373	<b>Variance</b>	110.741379
<b>Skewness</b>	0.68765047	<b>Kurtosis</b>	0.08078095

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	60
<b>99%</b>	60
<b>95%</b>	51
<b>90%</b>	47
<b>75% Q3</b>	39
<b>50% Median</b>	33
<b>25% Q1</b>	25
<b>10%</b>	20
<b>5%</b>	18
<b>1%</b>	18
<b>0% Min</b>	18

The SAS System

The NPARIWAY Procedure  
hepatitis=reagent

<b>Wilcoxon Scores (Rank Sums) for Variable age Classified by Variable sex</b>					
<b>sex</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>female</b>	29	1041.50	1174.50	99.834206	35.913793
<b>male</b>	51	2198.50	2065.50	99.834206	43.107843

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	1041.5000
<b>Normal Approximation</b>	
<b>Z</b>	-1.3272

<b>Wilcoxon Two-Sample Test</b>	
<b>One-Sided Pr &lt; Z</b>	0.0922
<b>Two-Sided Pr &gt;  Z </b>	0.1844

**Table 1.** Socio-demographic characteristics of HIV-infected patients

The SAS System

The FREQ Procedure

<b>Frequency</b>
<b>Col Pct</b>

<b>Table of Age of HIV_monoinfected_vs_HIV_HBV</b>			
	<b>HIV_HBV</b>	<b>HIV_monoinfected</b>	<b>Total</b>
<b>18-29</b>	10	476	486
	25.64	40.51	
<b>30-49</b>	23	505	528
	58.98	42.98	
<b>0-17</b>	0	55	55
	0.00	4.68	
<b>&gt;_50</b>	6	139	145
	15.38	11.83	
<b>Total</b>	39	1175	1214
<b>Frequency Missing = 45</b>			

Statistics for Table of transformed\_age\_1 by HIV\_monoinfected\_vs\_HIV\_HBV

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	3	6.5303	0.0885
<b>Likelihood Ratio Chi-Square</b>	3	8.3443	0.0394
<b>Mantel-Haenszel Chi-Square</b>	1	1.2403	0.2654
<b>Phi Coefficient</b>		0.0733	
<b>Contingency Coefficient</b>		0.0731	
<b>Cramer's V</b>		0.0733	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			



Effective Sample Size = 1214  
 Frequency Missing = 45

**Frequency**  
**Col Pct**

**Table of sex by HIV\_monoinfected\_vs\_HIV\_HBV**

	<b>HIV_HBV</b>	<b>HIV_monoinfected</b>	<b>Total</b>
<b>female</b>	17	499	516
	43.59	42.32	
<b>male</b>	22	680	702
	56.41	57.68	
<b>Total</b>	39	1179	1218

**Frequency Missing = 41**

Statistics for Table of sex by HIV\_monoinfected\_vs\_HIV\_HBV

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	1	0.0248	0.8749
<b>Likelihood Ratio Chi-Square</b>	1	0.0247	0.8751
<b>Continuity Adj. Chi-Square</b>	1	0.0000	1.0000
<b>Mantel-Haenszel Chi-Square</b>	1	0.0247	0.8750
<b>Phi Coefficient</b>		0.0045	
<b>Contingency Coefficient</b>		0.0045	
<b>Cramer's V</b>		0.0045	

**Fisher's Exact Test**

<b>Cell (1,1) Frequency (F)</b>	17
<b>Left-sided Pr &lt;= F</b>	0.6288
<b>Right-sided Pr &gt;= F</b>	0.4997
<b>Table Probability (P)</b>	0.1285
<b>Two-sided Pr &lt;= P</b>	0.8708

Effective Sample Size = 1218  
 Frequency Missing = 41

**Frequency**  
**Col Pct**

□

<b>Table of marital_status by HIV_monoinfected_vs_HIV_HBV</b>			
	<b>HIV_HBV</b>	<b>HIV_monoinfected</b>	<b>Total</b>
<b>divorced</b>	4	100	104
	10.26	8.54	
<b>married</b>	17	459	476
	43.59	39.20	
<b>single</b>	18	562	580
	46.15	47.99	
<b>widower</b>	0	50	50
	0.00	4.27	
<b>Total</b>	39	1171	1210
<b>Frequency Missing = 49</b>			

Statistics for Table of marital\_status by HIV\_monoinfected\_vs\_HIV\_HBV

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	3	2.0064	0.5711
<b>Likelihood Ratio Chi-Square</b>	3	3.6042	0.3075
<b>Mantel-Haenszel Chi-Square</b>	1	1.0948	0.2954
<b>Phi Coefficient</b>		0.0407	
<b>Contingency Coefficient</b>		0.0407	
<b>Cramer's V</b>		0.0407	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Effective Sample Size = 1210

Frequency Missing = 49

**Frequency**  
**Col Pct**

<b>Table of ethnicity by HIV_monoinfected_vs_HIV_HBV</b>			
	<b>HIV_HBV</b>	<b>HIV_monoinfected</b>	<b>Total</b>
<b>black</b>	2	42	44
	5.56	3.65	
<b>brown</b>	14	400	414
	38.89	34.72	
<b>other</b>	0	6	6
	0.00	0.52	
<b>white</b>	20	704	724
	55.55	61.11	
<b>Total</b>	36	1152	1188
<b>Frequency Missing = 71</b>			

Statistics for Table of ethnicity by HIV\_monoinfected\_vs\_HIV\_HBV

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	3	0.8820	0.8298
<b>Likelihood Ratio Chi-Square</b>	3	1.0181	0.7969
<b>Mantel-Haenszel Chi-Square</b>	1	0.5971	0.4397
<b>Phi Coefficient</b>		0.0272	
<b>Contingency Coefficient</b>		0.0272	
<b>Cramer's V</b>		0.0272	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Effective Sample Size = 1188

Frequency Missing = 71

**Frequency**  
**Col Pct**

Table of transformed\_degree\_level by HIV\_monoinfected\_vs\_HIV\_HBV

	HIV_HBV	HIV_monoinfected	Total
≤ 8 years	25 65.79	551 48.72	576
> 8 years	13 34.21	580 51.28	593
<b>Total</b>	38	1131	1169

**Frequency Missing = 90**

Statistics for Table of degree\_level by HIV\_monoinfected\_vs\_HIV\_HBV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	4.2868	0.0384
<b>Likelihood Ratio Chi-Square</b>	1	4.3516	0.0370
<b>Continuity Adj. Chi-Square</b>	1	3.6310	0.0567
<b>Mantel-Haenszel Chi-Square</b>	1	4.2831	0.0385
<b>Phi Coefficient</b>		0.0606	
<b>Contingency Coefficient</b>		0.0604	
<b>Cramer's V</b>		0.0606	

<b>Fisher's Exact Test</b>	
<b>Cell (1,1) Frequency (F)</b>	25
<b>Left-sided Pr ≤ F</b>	0.9878
<b>Right-sided Pr ≥ F</b>	0.0278
<b>Table Probability (P)</b>	0.0156
<b>Two-sided Pr ≤ P</b>	0.0471

Effective Sample Size = 1169  
Frequency Missing = 90

**Frequency**  
**Col Pct**

□

<b>Table of sexual_behavior by HIV_monoinfected_vs_HIV_HBV</b>			
	<b>HIV_HBV</b>	<b>HIV_monoinfected</b>	<b>Total</b>
<b>heterosexual</b>	32	880	912
	86.49	77.13	
<b>homosexual/bisexual</b>	5	261	266
	13.51	22.87	
<b>Total</b>	37	1141	1178
<b>Frequency Missing = 81</b>			

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	1	1.7965	0.1801
<b>Likelihood Ratio Chi-Square</b>	1	2.0103	0.1562
<b>Continuity Adj. Chi-Square</b>	1	1.3009	0.2541
<b>Mantel-Haenszel Chi-Square</b>	1	1.7949	0.1803
<b>Phi Coefficient</b>		0.0391	
<b>Contingency Coefficient</b>		0.0390	
<b>Cramer's V</b>		0.0391	

<b>Fisher's Exact Test</b>	
<b>Cell (1,1) Frequency (F)</b>	32
<b>Left-sided Pr &lt;= F</b>	0.9457
<b>Right-sided Pr &gt;= F</b>	0.1241
<b>Table Probability (P)</b>	0.0698
<b>Two-sided Pr &lt;= P</b>	0.2312

Effective Sample Size = 1178  
Frequency Missing = 81

**Frequency**  
**Col Pct**

WARNING: 20% of the data are missing.

<b>Table of way_of_acquiring_virus by HIV_monoinfected_vs_HIV_HBV</b>			
	<b>HIV_HBV</b>	<b>HIV_monoinfected</b>	<b>Total</b>
<b>OUTROS</b>	0	27	27
	0.00	2.31	
<b>sexual</b>	38	1144	1182
	100.00	97.69	
<b>Total</b>	38	1171	1209

**Frequency Missing = 50**

Statistics for Table of way\_of\_acquiring\_virus by HIV\_monoinfected\_vs\_HIV\_HBV

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	1	0.8962	0.3438
<b>Likelihood Ratio Chi-Square</b>	1	1.7444	0.1866
<b>Continuity Adj. Chi-Square</b>	1	0.1513	0.6973
<b>Mantel-Haenszel Chi-Square</b>	1	0.8954	0.3440
<b>Phi Coefficient</b>		-0.0272	
<b>Contingency Coefficient</b>		0.0272	
<b>Cramer's V</b>		-0.0272	

**WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.**

<b>Fisher's Exact Test</b>	
<b>Cell (1,1) Frequency (F)</b>	0
<b>Left-sided Pr &lt;= F</b>	0.4182
<b>Right-sided Pr &gt;= F</b>	1.0000
<b>Table Probability (P)</b>	0.4182
<b>Two-sided Pr &lt;= P</b>	1.0000

Effective Sample Size = 1209  
Frequency Missing = 50

**Frequency**

**Col Pct**

□ **Table of diagnostic\_period by HIV\_monoinfected\_vs\_HIV\_HBV**

	<b>HIV_HBV</b>	<b>HIV_monoinfected</b>	<b>Total</b>
<b>&gt;_5</b>	21 53.85	528 44.79	549
<b>3 to 5</b>	7 17.95	227 19.25	234
<b>&lt;_3</b>	11 28.20	424 35.96	435
<b>Total</b>	39	1179	1218

**Frequency Missing = 41**

Statistics for Table of diagnostic\_period by HIV\_monoinfected\_vs\_HIV\_HBV

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	1.3574	0.5073
<b>Likelihood Ratio Chi-Square</b>	2	1.3671	0.5048
<b>Mantel-Haenszel Chi-Square</b>	1	1.3354	0.2478
<b>Phi Coefficient</b>		0.0334	
<b>Contingency Coefficient</b>		0.0334	
<b>Cramer's V</b>		0.0334	

Effective Sample Size = 1218

Frequency Missing = 41

**Frequency**

**Col Pct**

□ **Table of Alcohol by HIV\_monoinfected\_vs\_HIV\_HBV**

	<b>HIV_HBV</b>	<b>HIV_monoinfected</b>	<b>Total</b>
<b>no</b>	29	900	929
	76.32	81.45	
<b>yes</b>	9	205	214
	23.68	18.55	
<b>Total</b>	38	1105	1143

**Frequency Missing = 116**

Statistics for Table of Alcohol by HIV\_monoinfected\_vs\_HIV\_HBV

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	1	0.6359	0.4252
<b>Likelihood Ratio Chi-Square</b>	1	0.5993	0.4389
<b>Continuity Adj. Chi-Square</b>	1	0.3433	0.5579
<b>Mantel-Haenszel Chi-Square</b>	1	0.6353	0.4254
<b>Phi Coefficient</b>		-0.0236	
<b>Contingency Coefficient</b>		0.0236	
<b>Cramer's V</b>		-0.0236	

<b>Fisher's Exact Test</b>	
<b>Cell (1,1) Frequency (F)</b>	29
<b>Left-sided Pr &lt;= F</b>	0.2701
<b>Right-sided Pr &gt;= F</b>	0.8439
<b>Table Probability (P)</b>	0.1140
<b>Two-sided Pr &lt;= P</b>	0.4015

Effective Sample Size = 1143

Frequency Missing = 116



Frequency Col Pct	Table of injectable_drugs by HIV_monoinfected_vs_HIV_HBV		
	HIV_HBV	HIV_monoinfected	Total
<b>no</b>	36	1104	1140
	100.00	99.28	
<b>yes</b>	0	8	8
	0.00	0.72	
<b>Total</b>	36	1112	1148
<b>Frequency Missing = 111</b>			

Statistics for Table of injectable\_drugs by HIV\_monoinfected\_vs\_HIV\_HBV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	0.2608	0.6096
<b>Likelihood Ratio Chi-Square</b>	1	0.5116	0.4745
<b>Continuity Adj. Chi-Square</b>	1	0.0000	1.0000
<b>Mantel-Haenszel Chi-Square</b>	1	0.2606	0.6097
<b>Phi Coefficient</b>		0.0151	
<b>Contingency Coefficient</b>		0.0151	
<b>Cramer's V</b>		0.0151	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	36
<b>Left-sided Pr &lt;= F</b>	1.0000
<b>Right-sided Pr &gt;= F</b>	0.7744
<b>Table Probability (P)</b>	0.7744
<b>Two-sided Pr &lt;= P</b>	1.0000

Effective Sample Size = 1148  
Frequency Missing = 111

Frequency Col Pct	Table of tattoo by HIV_monoinfected_vs_HIV_HBV		
	HIV_HBV	HIV_monoinfected	Total
<b>no</b>	31 91.18	986 86.11	1017
<b>yes</b>	3 8.82	159 13.89	162
<b>Total</b>	34	1145	1179
<b>Frequency Missing = 80</b>			

Statistics for Table of tattoo by HIV\_monoinfected\_vs\_HIV\_HBV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	0.7141	0.3981
<b>Likelihood Ratio Chi-Square</b>	1	0.8000	0.3711
<b>Continuity Adj. Chi-Square</b>	1	0.3508	0.5536
<b>Mantel-Haenszel Chi-Square</b>	1	0.7135	0.3983
<b>Phi Coefficient</b>		0.0246	
<b>Contingency Coefficient</b>		0.0246	
<b>Cramer's V</b>		0.0246	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	31
<b>Left-sided Pr &lt;= F</b>	0.8681
<b>Right-sided Pr &gt;= F</b>	0.2907
<b>Table Probability (P)</b>	0.1588
<b>Two-sided Pr &lt;= P</b>	0.6114

Effective Sample Size = 1179  
Frequency Missing = 80

*HIV\_monoinfected\_vs.\_HIV\_HCV*

The SAS System

The FREQ Procedure

Frequency Col Pct	Table of transformed_age_1 by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>18-29</b>	16	476	492
	41.02	40.51	
<b>30-49</b>	17	505	522
	43.59	42.98	
<b>0-17</b>	2	55	57
	5.13	4.68	
<b>&gt;_50</b>	4	139	143
	10.26	11.83	
<b>Total</b>	39	1175	1214
	<b>Frequency Missing = 45</b>		

Statistics for Table of transformed\_age\_1 by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	3	0.1012	0.9917
<b>Likelihood Ratio Chi-Square</b>	3	0.1042	0.9913
<b>Mantel-Haenszel Chi-Square</b>	1	0.0429	0.8359
<b>Phi Coefficient</b>		0.0091	
<b>Contingency Coefficient</b>		0.0091	
<b>Cramer's V</b>		0.0091	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Effective Sample Size = 1214

Frequency Missing = 45

Frequency Col Pct	Table of sex by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>female</b>	12	499	511
	30.77	42.32	
<b>male</b>	27	680	707
	69.23	57.68	
<b>Total</b>	39	1179	1218
<b>Frequency Missing = 41</b>			

Statistics for Table of sex by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	2.0697	0.1503
<b>Likelihood Ratio Chi-Square</b>	1	2.1401	0.1435
<b>Continuity Adj. Chi-Square</b>	1	1.6224	0.2028
<b>Mantel-Haenszel Chi-Square</b>	1	2.0680	0.1504
<b>Phi Coefficient</b>		-0.0412	
<b>Contingency Coefficient</b>		0.0412	
<b>Cramer's V</b>		-0.0412	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	12
<b>Left-sided Pr &lt;= F</b>	0.1001
<b>Right-sided Pr &gt;= F</b>	0.9477
<b>Table Probability (P)</b>	0.0478
<b>Two-sided Pr &lt;= P</b>	0.1870

Effective Sample Size = 1218

Frequency Missing = 41

Frequency Col Pct	Table of marital_status by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>divorced</b>	2 5.41	100 8.54	102
<b>married</b>	16 43.24	459 39.20	475
<b>single</b>	18 48.65	562 47.99	580
<b>widower</b>	1 2.70	50 4.27	51
<b>Total</b>	37	1171	1208

**Frequency Missing = 51**

Statistics for Table of marital\_status by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	3	0.7785	0.8546
<b>Likelihood Ratio Chi-Square</b>	3	0.8634	0.8342
<b>Mantel-Haenszel Chi-Square</b>	1	0.0031	0.9559
<b>Phi Coefficient</b>		0.0254	
<b>Contingency Coefficient</b>		0.0254	
<b>Cramer's V</b>		0.0254	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Effective Sample Size = 1208

Frequency Missing = 51

Frequency Col Pct	Table of ethnicity by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>black</b>	0	42	42
	0.00	3.65	
<b>brown</b>	13	400	413
	35.14	34.72	
<b>other</b>	0	6	6
	0.00	0.52	
<b>white</b>	24	704	728
	64.86	61.11	
<b>Total</b>	37	1152	1189
<b>Frequency Missing = 70</b>			

Statistics for Table of ethnicity by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	3	1.6259	0.6535
<b>Likelihood Ratio Chi-Square</b>	3	3.1176	0.3739
<b>Mantel-Haenszel Chi-Square</b>	1	0.3809	0.5371
<b>Phi Coefficient</b>		0.0370	
<b>Contingency Coefficient</b>		0.0370	
<b>Cramer's V</b>		0.0370	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Effective Sample Size = 1189

Frequency Missing = 70

Frequency Col Pct	Table of degree_level by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>≤ 8 years</b>	22 61.11	551 48.72	573
<b>&gt; 8 years</b>	14 38.89	580 51.28	594
<b>Total</b>	36	1131	1167
<b>Frequency Missing = 92</b>			

Statistics for Table of degree\_level by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	2.1442	0.1431
<b>Likelihood Ratio Chi-Square</b>	1	2.1585	0.1418
<b>Continuity Adj. Chi-Square</b>	1	1.6770	0.1953
<b>Mantel-Haenszel Chi-Square</b>	1	2.1423	0.1433
<b>Phi Coefficient</b>		0.0429	
<b>Contingency Coefficient</b>		0.0428	
<b>Cramer's V</b>		0.0429	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	22
<b>Left-sided Pr ≤ F</b>	0.9493
<b>Right-sided Pr ≥ F</b>	0.0975
<b>Table Probability (P)</b>	0.0467
<b>Two-sided Pr ≤ P</b>	0.1755

Effective Sample Size = 1167

Frequency Missing = 92

Frequency Col Pct	Table of sexual_behavior by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>heterosexual</b>	32	880	912
	86.49	77.13	
<b>homosexual/bisexual</b>	5	261	266
	13.51	22.87	
<b>Total</b>	37	1141	1178
<b>Frequency Missing = 81</b>			

Statistics for Table of sexual\_behavior by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	1.7965	0.1801
<b>Likelihood Ratio Chi-Square</b>	1	2.0103	0.1562
<b>Continuity Adj. Chi-Square</b>	1	1.3009	0.2541
<b>Mantel-Haenszel Chi-Square</b>	1	1.7949	0.1803
<b>Phi Coefficient</b>		0.0391	
<b>Contingency Coefficient</b>		0.0390	
<b>Cramer's V</b>		0.0391	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	32
<b>Left-sided Pr &lt;= F</b>	0.9457
<b>Right-sided Pr &gt;= F</b>	0.1241
<b>Table Probability (P)</b>	0.0698
<b>Two-sided Pr &lt;= P</b>	0.2312

Effective Sample Size = 1178

Frequency Missing = 81



WARNING: 21% of the data are missing.

Frequency Col Pct	Table of way_of_acquiring_virus by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>OUTROS</b>	0	27	27
	0.00	2.31	
<b>sexual</b>	38	1144	1182
	100.00	97.69	
<b>Total</b>	38	1171	1209
<b>Frequency Missing = 50</b>			

Statistics for Table of way\_of\_acquiring\_virus by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	0.8962	0.3438
<b>Likelihood Ratio Chi-Square</b>	1	1.7444	0.1866
<b>Continuity Adj. Chi-Square</b>	1	0.1513	0.6973
<b>Mantel-Haenszel Chi-Square</b>	1	0.8954	0.3440
<b>Phi Coefficient</b>		-0.0272	
<b>Contingency Coefficient</b>		0.0272	
<b>Cramer's V</b>		-0.0272	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	0
<b>Left-sided Pr &lt;= F</b>	0.4182
<b>Right-sided Pr &gt;= F</b>	1.0000
<b>Table Probability (P)</b>	0.4182
<b>Two-sided Pr &lt;= P</b>	1.0000

Effective Sample Size = 1209

Frequency Missing = 50

Frequency Col Pct	Table of diagnostic_period by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
> 5 years	31 79.49	528 44.79	559
3-5 years	5 12.82	227 19.25	232
>5 years	3 7.69	424 35.96	427
<b>Total</b>	39	1179	1218
<b>Frequency Missing = 41</b>			

Statistics for Table of diagnostic\_period by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	2	19.3328	<.0001
<b>Likelihood Ratio Chi-Square</b>	2	21.6126	<.0001
<b>Mantel-Haenszel Chi-Square</b>	1	18.7505	<.0001
<b>Phi Coefficient</b>		0.1260	
<b>Contingency Coefficient</b>		0.1250	
<b>Cramer's V</b>		0.1260	

Effective Sample Size = 1218  
Frequency Missing = 41

Frequency Col Pct	Table of Alcohol by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>no</b>	19 59.38	900 81.45	919
<b>yes</b>	13 40.63	205 18.55	218
<b>Total</b>	32	1105	1137
<b>Frequency Missing = 122</b>			

Statistics for Table of Alcohol by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	9.7774	0.0018
<b>Likelihood Ratio Chi-Square</b>	1	8.0795	0.0045
<b>Continuity Adj. Chi-Square</b>	1	8.4049	0.0037
<b>Mantel-Haenszel Chi-Square</b>	1	9.7688	0.0018
<b>Phi Coefficient</b>		-0.0927	
<b>Contingency Coefficient</b>		0.0923	
<b>Cramer's V</b>		-0.0927	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	19
<b>Left-sided Pr &lt;= F</b>	0.0037
<b>Right-sided Pr &gt;= F</b>	0.9989
<b>Table Probability (P)</b>	0.0026
<b>Two-sided Pr &lt;= P</b>	0.0046

Effective Sample Size = 1137

Frequency Missing = 122

Frequency Col Pct	Table of injectable_drugs by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>no</b>	30 90.91	1104 99.28	1134
<b>yes</b>	3 9.09	8 0.72	11
<b>Total</b>	33	1112	1145
<b>Frequency Missing = 114</b>			

Statistics for Table of injectable\_drugs by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	23.6061	<.0001
<b>Likelihood Ratio Chi-Square</b>	1	9.0899	0.0026
<b>Continuity Adj. Chi-Square</b>	1	15.6274	<.0001
<b>Mantel-Haenszel Chi-Square</b>	1	23.5854	<.0001
<b>Phi Coefficient</b>		-0.1436	
<b>Contingency Coefficient</b>		0.1421	
<b>Cramer's V</b>		-0.1436	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	30
<b>Left-sided Pr &lt;= F</b>	0.0031
<b>Right-sided Pr &gt;= F</b>	0.9998
<b>Table Probability (P)</b>	0.0029
<b>Two-sided Pr &lt;= P</b>	0.0031

Effective Sample Size = 1145  
Frequency Missing = 114

Frequency Col Pct	Table of tattoo by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>no</b>	29 90.63	986 86.11	1015
<b>yes</b>	3 9.38	159 13.89	162
<b>Total</b>	32	1145	1177
<b>Frequency Missing = 82</b>			

Statistics for Table of tattoo by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	0.5338	0.4650
<b>Likelihood Ratio Chi-Square</b>	1	0.5896	0.4426
<b>Continuity Adj. Chi-Square</b>	1	0.2214	0.6380
<b>Mantel-Haenszel Chi-Square</b>	1	0.5334	0.4652
<b>Phi Coefficient</b>		0.0213	
<b>Contingency Coefficient</b>		0.0213	
<b>Cramer's V</b>		0.0213	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	29
<b>Left-sided Pr &lt;= F</b>	0.8397
<b>Right-sided Pr &gt;= F</b>	0.3373
<b>Table Probability (P)</b>	0.1770
<b>Two-sided Pr &lt;= P</b>	0.6082

Effective Sample Size = 1177  
Frequency Missing = 82

*HIV\_monoinfected\_vs\_Hepatitis*

The SAS System

The FREQ Procedure

Frequency Col Pct	Table of transformed_age_1 by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>18-29</b>	476 40.51	26 32.50	502
<b>30-49</b>	505 42.98	42 52.50	547
<b>0-17</b>	55 4.68	2 2.50	57
<b>&gt;_50</b>	139 11.83	10 12.50	149
<b>Total</b>	1175	80	1255
<b>Frequency Missing = 4</b>			

Statistics for Table of transformed\_age\_1 by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	3	3.5721	0.3115
<b>Likelihood Ratio Chi-Square</b>	3	3.7031	0.2954
<b>Mantel-Haenszel Chi-Square</b>	1	0.4244	0.5147
<b>Phi Coefficient</b>		0.0534	
<b>Contingency Coefficient</b>		0.0533	
<b>Cramer's V</b>		0.0534	

Effective Sample Size = 1255

Frequency Missing = 4

Frequency Col Pct	Table of sex by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>female</b>	499	29	528
	42.32	36.25	
<b>male</b>	680	51	731
	57.68	63.75	
<b>Total</b>	1179	80	1259

Statistics for Table of sex by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	1.1351	0.2867
<b>Likelihood Ratio Chi-Square</b>	1	1.1511	0.2833
<b>Continuity Adj. Chi-Square</b>	1	0.8993	0.3430
<b>Mantel-Haenszel Chi-Square</b>	1	1.1342	0.2869
<b>Phi Coefficient</b>		0.0300	
<b>Contingency Coefficient</b>		0.0300	
<b>Cramer's V</b>		0.0300	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	499
<b>Left-sided Pr &lt;= F</b>	0.8820
<b>Right-sided Pr &gt;= F</b>	0.1716
<b>Table Probability (P)</b>	0.0537
<b>Two-sided Pr &lt;= P</b>	0.2952

Sample Size = 1259

Frequency Col Pct	Table of marital_status by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>divorced</b>	100	6	106
	8.54	7.79	
<b>married</b>	459	34	493
	39.20	44.16	
<b>single</b>	562	36	598
	47.99	46.75	
<b>widower</b>	50	1	51
	4.27	1.30	
<b>Total</b>	1171	77	1248
<b>Frequency Missing = 11</b>			

Statistics for Table of marital\_status by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	3	2.0811	0.5557
<b>Likelihood Ratio Chi-Square</b>	3	2.6018	0.4572
<b>Mantel-Haenszel Chi-Square</b>	1	0.5966	0.4399
<b>Phi Coefficient</b>		0.0408	
<b>Contingency Coefficient</b>		0.0408	
<b>Cramer's V</b>		0.0408	

Effective Sample Size = 1248

Frequency Missing = 11



Frequency Col Pct	Table of ethnicity by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>black</b>	42	2	44
	3.65	2.67	
<b>brown</b>	400	28	428
	34.72	37.33	
<b>other</b>	6	0	6
	0.52	0.00	
<b>white</b>	704	45	749
	61.11	60.00	
<b>Total</b>	1152	75	1227
<b>Frequency Missing = 32</b>			

Statistics for Table of ethnicity by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	3	0.7308	0.8659
<b>Likelihood Ratio Chi-Square</b>	3	1.1115	0.7743
<b>Mantel-Haenszel Chi-Square</b>	1	0.0205	0.8861
<b>Phi Coefficient</b>		0.0244	
<b>Contingency Coefficient</b>		0.0244	
<b>Cramer's V</b>		0.0244	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Effective Sample Size = 1227

Frequency Missing = 32

Frequency Col Pct	Table of degree_level by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
$\leq 8$ years	551 48.72	48 64.00	599
$> 8$ years	580 51.28	27 36.00	607
<b>Total</b>	1131	75	1206
<b>Frequency Missing = 53</b>			

Statistics for Table of degree\_level by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	6.5708	0.0104
<b>Likelihood Ratio Chi-Square</b>	1	6.6500	0.0099
<b>Continuity Adj. Chi-Square</b>	1	5.9737	0.0145
<b>Mantel-Haenszel Chi-Square</b>	1	6.5654	0.0104
<b>Phi Coefficient</b>		-0.0738	
<b>Contingency Coefficient</b>		0.0736	
<b>Cramer's V</b>		-0.0738	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	551
<b>Left-sided Pr <math>\leq F</math></b>	0.0070
<b>Right-sided Pr <math>\geq F</math></b>	0.9965
<b>Table Probability (P)</b>	0.0036
<b>Two-sided Pr <math>\leq P</math></b>	0.0120

Effective Sample Size = 1206

Frequency Missing = 53

Frequency Col Pct	Table of sexual_behavior by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>heterosexual</b>	880	66	946
	77.13	86.84	
<b>homosexual/bisexual</b>	261	10	271
	22.87	13.16	
<b>Total</b>	1141	76	1217
<b>Frequency Missing = 42</b>			

Statistics for Table of sexual\_behavior by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	3.8866	0.0487
<b>Likelihood Ratio Chi-Square</b>	1	4.3473	0.0371
<b>Continuity Adj. Chi-Square</b>	1	3.3455	0.0674
<b>Mantel-Haenszel Chi-Square</b>	1	3.8834	0.0488
<b>Phi Coefficient</b>		-0.0565	
<b>Contingency Coefficient</b>		0.0564	
<b>Cramer's V</b>		-0.0565	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	880
<b>Left-sided Pr &lt;= F</b>	0.0289
<b>Right-sided Pr &gt;= F</b>	0.9868
<b>Table Probability (P)</b>	0.0156
<b>Two-sided Pr &lt;= P</b>	0.0472

Effective Sample Size = 1217

Frequency Missing = 42

WARNING: 18% of the data are missing.

Frequency Col Pct	Table of way_of_acquiring_virus by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>OUTROS</b>	27	0	27
	2.31	0.00	
<b>sexual</b>	1144	78	1222
	97.69	100.00	
<b>Total</b>	1171	78	1249
<b>Frequency Missing = 10</b>			

Statistics for Table of way\_of\_acquiring\_virus by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	1.8382	0.1752
<b>Likelihood Ratio Chi-Square</b>	1	3.5217	0.0606
<b>Continuity Adj. Chi-Square</b>	1	0.9097	0.3402
<b>Mantel-Haenszel Chi-Square</b>	1	1.8367	0.1753
<b>Phi Coefficient</b>		0.0384	
<b>Contingency Coefficient</b>		0.0383	
<b>Cramer's V</b>		0.0384	
<b>WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.</b>			

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	27
<b>Left-sided Pr &lt;= F</b>	1.0000
<b>Right-sided Pr &gt;= F</b>	0.1720
<b>Table Probability (P)</b>	0.1720
<b>Two-sided Pr &lt;= P</b>	0.4072

Effective Sample Size = 1249

Frequency Missing = 10

Frequency Col Pct	Table of _diagnostic_period by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
> 5 years	528 44.78	54 67.50	582
3-5 years	227 19.25	12 15.00	239
< 3 years	424 35.96	14 17.50	438
<b>Total</b>	<b>1179</b>	<b>80</b>	<b>1259</b>

Statistics for Table of transformed\_diagnostic\_period by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	2	16.4173	0.0003
<b>Likelihood Ratio Chi-Square</b>	2	17.0364	0.0002
<b>Mantel-Haenszel Chi-Square</b>	1	15.9250	<.0001
<b>Phi Coefficient</b>		0.1142	
<b>Contingency Coefficient</b>		0.1135	
<b>Cramer's V</b>		0.1142	

Sample Size = 1259

Frequency Col Pct	Table of Alcohol by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>no</b>	900 81.45	50 69.44	950
<b>yes</b>	205 18.55	22 30.56	227
<b>Total</b>	1105	72	1177

**Frequency Missing = 82**

Statistics for Table of Alcohol by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	6.2566	0.0124
<b>Likelihood Ratio Chi-Square</b>	1	5.5950	0.0180
<b>Continuity Adj. Chi-Square</b>	1	5.5093	0.0189
<b>Mantel-Haenszel Chi-Square</b>	1	6.2513	0.0124
<b>Phi Coefficient</b>		0.0729	
<b>Contingency Coefficient</b>		0.0727	
<b>Cramer's V</b>		0.0729	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	900
<b>Left-sided Pr &lt;= F</b>	0.9943
<b>Right-sided Pr &gt;= F</b>	0.0122
<b>Table Probability (P)</b>	0.0065
<b>Two-sided Pr &lt;= P</b>	0.0196

Effective Sample Size = 1177

Frequency Missing = 82

Frequency Col Pct	Table of injectable_drugs by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>no</b>	1104 99.28	68 95.77	1172
<b>yes</b>	8 0.72	3 4.23	11
<b>Total</b>	1112	71	1183

**Frequency Missing = 76**

Statistics for Table of injectable\_drugs by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	8.9050	0.0028
<b>Likelihood Ratio Chi-Square</b>	1	5.0618	0.0245
<b>Continuity Adj. Chi-Square</b>	1	5.5058	0.0190
<b>Mantel-Haenszel Chi-Square</b>	1	8.8975	0.0029
<b>Phi Coefficient</b>		0.0868	
<b>Contingency Coefficient</b>		0.0864	
<b>Cramer's V</b>		0.0868	

**WARNING: 25% of the cells have expected counts less than 5. Chi-Square may not be a valid test.**

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	1104
<b>Left-sided Pr &lt;= F</b>	0.9971
<b>Right-sided Pr &gt;= F</b>	0.0241
<b>Table Probability (P)</b>	0.0213
<b>Two-sided Pr &lt;= P</b>	0.0241

Effective Sample Size = 1183  
Frequency Missing = 76

Frequency Col Pct	Table of tattoo by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>no</b>	986 86.11	62 91.18	1048
<b>yes</b>	159 13.89	6 8.82	165
<b>Total</b>	1145	68	1213
<b>Frequency Missing = 46</b>			

Statistics for Table of tattoo by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	1	1.4000	0.2367
<b>Likelihood Ratio Chi-Square</b>	1	1.5601	0.2117
<b>Continuity Adj. Chi-Square</b>	1	1.0024	0.3167
<b>Mantel-Haenszel Chi-Square</b>	1	1.3989	0.2369
<b>Phi Coefficient</b>		-0.0340	
<b>Contingency Coefficient</b>		0.0340	
<b>Cramer's V</b>		-0.0340	

Fisher's Exact Test	
<b>Cell (1,1) Frequency (F)</b>	986
<b>Left-sided Pr &lt;= F</b>	0.1579
<b>Right-sided Pr &gt;= F</b>	0.9208
<b>Table Probability (P)</b>	0.0787
<b>Two-sided Pr &lt;= P</b>	0.2784

Effective Sample Size = 1213

Frequency Missing = 46



The FREQ Procedure

HIV\_monoinfected

<b>HIV_monoinfected</b>				
<b>HIV_monoinfected</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>HIV_monoinfected</b>	1179	100.00	1179	100.00

<b>anti_HBs</b>				
<b>anti_HBs</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>non_reactive</b>	605	53.73	605	53.73
<b>reagent</b>	521	46.27	1126	100.00

<b>anti_HBC</b>				
<b>anti_HBC</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>non_reactive</b>	900	83.26	900	83.26
<b>reagent</b>	181	16.74	1181	100.00

Frequency Missing = 108

The SAS System

The FREQ Procedure

HIV\_HCV=reagent

<b>HIV_HCV</b>				
<b>HIV_HCV</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>reagent</b>	39	100.00	39	100.00

<b>anti_HBs</b>				
<b>anti_HBs</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>non_reactive</b>	22	68.75	22	68.75
<b>reagent</b>	10	31.25	32	100.00

<b>anti_HBC</b>				
<b>anti_HBC</b>	<b>Frequency</b>	<b>Percent</b>	<b>Cumulative Frequency</b>	<b>Cumulative Percent</b>
<b>non_reactive</b>	25	78.24	25	78.24
<b>reagent</b>	7	21.86	32	100.00

The SAS System

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_monoinfected\_female

<b>Moments</b>			
<b>N</b>	476	<b>Sum Weights</b>	476
<b>Mean</b>	503.2164		
<b>Std Deviation</b>	333.5076	<b>Variance</b>	111227.3
<b>Skewness</b>	1.188964	<b>Kurtosis</b>	5.550357

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1636
<b>99%</b>	1636
<b>95%</b>	1115
<b>90%</b>	894
<b>75% Q3</b>	688.5
<b>50% Median</b>	450.5
<b>25% Q1</b>	265.5
<b>10%</b>	120
<b>5%</b>	62
<b>1%</b>	20

The SAS System

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_monoinfected\_male

<b>Moments</b>			
<b>N</b>	627	<b>Sum Weights</b>	627
<b>Mean</b>	408.7177		

<b>Moments</b>			
<b>Std Deviation</b>	269.8876	<b>Variance</b>	72839.32
<b>Skewness</b>	1.136779	<b>Kurtosis</b>	5.630798

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1248
<b>99%</b>	1248
<b>95%</b>	918
<b>90%</b>	764
<b>75% Q3</b>	538
<b>50% Median</b>	379
<b>25% Q1</b>	213
<b>10%</b>	91
<b>5%</b>	43
<b>1%</b>	9

<b>Wilcoxon Scores (Rank Sums) for Variable cd4_diagnosis (cd4_diagnosis) Classified by Variable HIV_monoinfected female vs male</b>		
<b>female vs male</b>	<b>N</b>	<b>Sum of Scores</b>
<b>female</b>	476	476
<b>male</b>	627	627

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	
<b>U</b>	124944.00
<b>Z</b>	4.6340
<b>One-Sided Pr &lt; Z</b>	<0.0001
<b>Two-Sided Pr &gt;  Z </b>	<0.0001

The SAS System  
The UNIVARIATE Procedure  
Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_HBV\_female

<b>Moments</b>			
<b>N</b>	16	<b>Sum Weights</b>	16
<b>Mean</b>	504.6875		
<b>Std Deviation</b>	223.0048	<b>Variance</b>	49731.16
<b>Skewness</b>	1.231659	<b>Kurtosis</b>	5.26303

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1138
<b>99%</b>	1138
<b>95%</b>	1138
<b>90%</b>	718
<b>75% Q3</b>	590.5
<b>50% Median</b>	475
<b>25% Q1</b>	383.5
<b>10%</b>	257
<b>5%</b>	161
<b>1%</b>	161

The SAS System

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_HBV\_male

<b>Moments</b>			
<b>N</b>	21	<b>Sum Weights</b>	21
<b>Mean</b>	376.1905		
<b>Std Deviation</b>	214.5259	<b>Variance</b>	46021.36
<b>Skewness</b>	0.7132155	<b>Kurtosis</b>	3.619756

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	938
<b>99%</b>	938
<b>95%</b>	701
<b>90%</b>	638
<b>75% Q3</b>	477
<b>50% Median</b>	336

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>25% Q1</b>	258
<b>10%</b>	156
<b>5%</b>	74
<b>1%</b>	17

<b>Wilcoxon Scores (Rank Sums) for Variable cd4_diagnosis (cd4_diagnosis) Classified by Variable HIV_HBV female vs male</b>		
<b>female vs male</b>	<b>N</b>	<b>Sum of Scores</b>
<b>Female</b>	16	16
<b>male</b>	21	21

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	
<b>U</b>	108.00
<b>Z</b>	1.8394
<b>One-Sided Pr &lt; Z</b>	0.0329
<b>Two-Sided Pr &gt;  Z </b>	0.0659

The SAS System

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_HCV\_female

<b>Moments</b>			
<b>N</b>	<b>11</b>	<b>Sum Weights</b>	<b>11</b>
<b>Mean</b>	360.2727		
<b>Std Deviation</b>	184.5487	<b>Variance</b>	34058.22
<b>Skewness</b>	0.1671016	<b>Kurtosis</b>	1.675045

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	666
<b>99%</b>	666
<b>95%</b>	666

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>90%</b>	568
<b>75% Q3</b>	506
<b>50% Median</b>	410
<b>25% Q1</b>	185
<b>10%</b>	141
<b>5%</b>	134
<b>1%</b>	134

The SAS System

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_HCV\_male

<b>Moments</b>			
<b>N</b>	25	<b>Sum Weights</b>	25
<b>Mean</b>	421.15		
<b>Std Deviation</b>	249.9246	<b>Variance</b>	62462.31
<b>Skewness</b>	1.359291	<b>Kurtosis</b>	5.100264

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1211
<b>99%</b>	1211
<b>95%</b>	815
<b>90%</b>	726
<b>75% Q3</b>	522
<b>50% Median</b>	345
<b>25% Q1</b>	243
<b>10%</b>	154
<b>5%</b>	147
<b>1%</b>	98

**Wilcoxon Scores (Rank Sums) for Variable cd4\_diagnosis (cd4\_diagnosis) Classified by Variable HIV\_HCV female vs male**

female vs male	N	Sum of Scores
female	11	11
male	25	25

**Wilcoxon Two-Sample Test**

**Statistic**

<b>U</b>	119
<b>Z</b>	0.6353
<b>One-Sided Pr &lt; Z</b>	0.2626
<b>Two-Sided Pr &gt;  Z </b>	0.5252

The SAS System

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) All\_cd4 ≤ 18\_age

**Moments**

<b>N</b>	71	<b>Sum Weights</b>	71
<b>Mean</b>	660.0423		
<b>Std Deviation</b>	379.138	<b>Variance</b>	143745.6
<b>Skewness</b>	1.312528	<b>Kurtosis</b>	4.967981

**Quantiles (Definition 5)**

Quantile	Estimate
<b>100% Max</b>	2001
<b>99%</b>	2001
<b>95%</b>	1402
<b>90%</b>	1175
<b>75% Q3</b>	764
<b>50% Median</b>	582
<b>25% Q1</b>	423
<b>10%</b>	265
<b>5%</b>	156
<b>1%</b>	97

The SAS System

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) All\_cd4  $\geq$  50\_age

<b>Moments</b>			
<b>N</b>	139	<b>Sum Weights</b>	139
<b>Mean</b>	374.3094		
<b>Std Deviation</b>	263.7228	<b>Variance</b>	69549.71
<b>Skewness</b>	1.321624	<b>Kurtosis</b>	5.486856

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1361
<b>99%</b>	1361
<b>95%</b>	838
<b>90%</b>	756
<b>75% Q3</b>	507
<b>50% Median</b>	333
<b>25% Q1</b>	168
<b>10%</b>	97
<b>5%</b>	54
<b>1%</b>	20

<b>Wilcoxon Scores (Rank Sums) for Variable cd4_diagnosis (cd4_diagnosis) Classified by Variable All <math>\leq</math> 18_age_vs <math>\geq</math> 50_age</b>		
<b><math>\leq</math> 18_age_vs <math>\geq</math> 50_age</b>	<b>N</b>	<b>Sum of Scores</b>
<b><math>\leq</math> 18_age</b>	71	71
<b><math>\geq</math> 50_age</b>	139	139

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	
<b>U</b>	2447.50
<b>Z</b>	5.9702
<b>One-Sided Pr &lt; Z</b>	<0.0001
<b>Two-Sided Pr &gt;  Z </b>	<0.0001



**Table 3.** CD4+ T-cell count and HIV viral load and their association with HBV and HCV in HIV-infected patients at baseline

The SAS System

Frequency Col Pct	Table of cd4_diagnosis_transformed by HIV_monoinfected_vs_HIV_HBV		
	HIV_HBV	HIV_monoinfected	Total
<b>200-500</b>	20 54.06	473 42.88	493
<b>&lt;200</b>	5 13.51	232 21.03	237
<b>&gt;500</b>	12 32.43	398 36.08	410
<b>Total</b>	37	1103	1140
<b>Frequency Missing = 119</b>			

Statistics for Table of cd4\_diagnosis\_transformed by HIV\_monoinfected\_vs\_HIV\_HBV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	2	2.1395	0.3371
<b>Likelihood Ratio Chi-Square</b>	2	2.2042	0.3322
<b>Mantel-Haenszel Chi-Square</b>	1	0.9987	0.3176
<b>Phi Coefficient</b>		0.0433	
<b>Contingency Coefficient</b>		0.0433	
<b>Cramer's V</b>		0.0433	

Effective Sample Size = 1140

Frequency Missing = 119

Frequency Col Pct	Table of cd4_diagnosis_transformed by HIV_monoinfected_vs_HIV_HCV		
	HIV_HCV	HIV_monoinfected	Total
<b>200-500</b>	20 55.55	473 42.88	493
<b>&lt;200</b>	6 16.67	232 21.03	238
<b>&gt;500</b>	10 27.78	398 36.08	408
<b>Total</b>	36	1103	1139
<b>Frequency Missing = 120</b>			

Statistics for Table of cd4\_diagnosis\_transformed by HIV\_monoinfected\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	2	2.2830	0.3143
<b>Likelihood Ratio Chi-Square</b>	2	2.2572	0.3235
<b>Mantel-Haenszel Chi-Square</b>	1	1.9515	0.1624
<b>Phi Coefficient</b>		0.0448	
<b>Contingency Coefficient</b>		0.0447	
<b>Cramer's V</b>		0.0448	

Effective Sample Size = 1139

Frequency Missing = 120

Frequency Col Pct	Table of cd4_diagnosis_transformed by HIV_monoinfected_vs_Hepatitis		
	HIV_monoinfected	Hepatitis	Total
<b>200-500</b>	473	42	515
	42.88	56.00	
<b>&lt;200</b>	232	11	243
	21.03	14.67	
<b>&gt;500</b>	398	22	420
	36.08	29.33	
<b>Total</b>	1103	75	1178
<b>Frequency Missing = 81</b>			

Statistics for Table of cd4\_diagnosis\_transformed by HIV\_monoinfected\_vs\_Hepatitis

Statistic	DF	Value	Prob
<b>Chi-Square</b>	2	5.0412	0.0774
<b>Likelihood Ratio Chi-Square</b>	2	5.0293	0.0809
<b>Mantel-Haenszel Chi-Square</b>	1	3.5180	0.0607
<b>Phi Coefficient</b>		0.0654	
<b>Contingency Coefficient</b>		0.0653	
<b>Cramer's V</b>		0.0654	

Effective Sample Size = 1178

Frequency Missing = 81

Frequency Col Pct	Table of cd4_diagnosis_transformed by HIV_HBV_vs_HIV_HCV		
	HIV_HBV	HIV_HCV	Total
<b>200-500</b>	5 13.51	6 16.67	11
<b>&lt;200</b>	20 54.05	20 55.56	40
<b>&gt;500</b>	12 32.43	10 27.78	22
<b>Total</b>	37	36	73

Statistics for Table of cd4\_diagnosis\_transformed by HIV\_HBV\_vs\_HIV\_HCV

Statistic	DF	Value	Prob
<b>Chi-Square</b>	2	2.2591	0.8790
<b>Likelihood Ratio Chi-Square</b>	2	2.0293	0.8809
<b>Mantel-Haenszel Chi-Square</b>	1	2.5180	0.8607
<b>Phi Coefficient</b>		0.0234	
<b>Contingency Coefficient</b>		0.0356	
<b>Cramer's V</b>		0.0346	

The UNIVARIATE Procedure  
 Variable: cd4\_diagnosis (cd4\_diagnosis)  
 HIV\_monoinfected

<b>Moments</b>			
<b>N</b>	1103	<b>Sum Weights</b>	1103
<b>Mean</b>	449.499547		
<b>Std Deviation</b>	302.509462	<b>Variance</b>	91511.9744
<b>Skewness</b>	1.24370692	<b>Kurtosis</b>	2.96352292

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	2283
<b>99%</b>	1408
<b>95%</b>	974
<b>90%</b>	828
<b>75% Q3</b>	608
<b>50% Median</b>	405
<b>25% Q1</b>	234
<b>10%</b>	99
<b>5%</b>	54
<b>1%</b>	14
<b>0% Min</b>	2

The SAS System

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_HBV

<b>Moments</b>			
<b>N</b>	37	<b>Sum Weights</b>	37
<b>Mean</b>	431.756757		
<b>Std Deviation</b>	224.619234	<b>Variance</b>	50453.8003
<b>Skewness</b>	0.90223774	<b>Kurtosis</b>	1.86296389

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1138
<b>99%</b>	1138
<b>95%</b>	938
<b>90%</b>	701
<b>75% Q3</b>	557
<b>50% Median</b>	387
<b>25% Q1</b>	287
<b>10%</b>	161
<b>5%</b>	74
<b>1%</b>	17

The NPAR1WAY Procedure

<b>Wilcoxon Scores (Rank Sums) for Variable cd4_diagnosis Classified by Variable HIV_monoinfected_vs_HIV_HBV</b>		
<b>HIV_monoinfected_vs_HIV_HBV</b>	<b>N</b>	<b>Sum of Scores</b>
<b>HIV_monoinfected</b>	1103	629075.0
<b>HIV_HBV</b>	37	21295.0

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	21295.0000
<b>Normal Approximation</b>	
<b>Z</b>	0.0944
<b>One-Sided Pr &gt; Z</b>	0.4624

**Wilcoxon Two-Sample Test**

**Two-Sided Pr > |Z|** 0.9248

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_HCV

<b>Moments</b>			
<b>N</b>	36	<b>Sum Weights</b>	36
<b>Mean</b>	402.5556		
<b>Std Deviation</b>	231.0224	<b>Variance</b>	53371.34
<b>Skewness</b>	1.289724	<b>Kurtosis</b>	5.348068

**Quantiles (Definition 5)**

<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1211
<b>99%</b>	1211
<b>95%</b>	815
<b>90%</b>	697
<b>75% Q3</b>	514
<b>50% Median</b>	377.5
<b>25% Q1</b>	228.5
<b>10%</b>	147
<b>5%</b>	134
<b>1%</b>	98

The SAS System

The NPAR1WAY Procedure

**Wilcoxon Scores (Rank Sums) for Variable cd4\_diagnosis  
Classified by Variable HIV\_monoinfected\_vs\_HIV\_HCV**

<b>HIV_monoinfected_vs_HIV_HCV</b>	<b>N</b>	<b>Sum of Scores</b>
<b>HIV_monoinfected</b>	1103	630146
<b>HIV_HCV</b>	36	19084

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

**Statistic** 18418.000

**Normal Approximation**

<b>Wilcoxon Two-Sample Test</b>	
<b>Z</b>	0.7394
<b>One-Sided Pr &gt; Z</b>	0.2298
<b>Two-Sided Pr &gt;  Z </b>	0.4597

The UNIVARIATE Procedure

Variable: cd4\_diagnosis (cd4\_diagnosis) HIV\_hepatitis

<b>Moments</b>			
<b>N</b>	75	<b>Sum Weights</b>	75
<b>Mean</b>	417.3733		
<b>Std Deviation</b>	223.6863	<b>Variance</b>	50035.56
<b>Skewness</b>	1.084653	<b>Kurtosis</b>	4.970194

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1211
<b>99%</b>	1211
<b>95%</b>	815
<b>90%</b>	697
<b>75% Q3</b>	522
<b>50% Median</b>	387
<b>25% Q1</b>	257
<b>10%</b>	156
<b>5%</b>	134
<b>1%</b>	17

The NPAR1WAY Procedure

<b>Wilcoxon Scores (Rank Sums) for Variable cd4_diagnosis Classified by Variable HIV_monoinfected_vs_HIV_hepatitis</b>		
<b>HIV_monoinfected_vs_HIV_HCV</b>	<b>N</b>	<b>Sum of Scores</b>
<b>HIV_monoinfected</b>	1103	651430.5
<b>HIV_hepatitis</b>	75	43000.5
<b>Average scores were used for ties.</b>		

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	40150.50



<b>Wilcoxon Two-Sample Test</b>	
<b>Normal Approximation</b>	
<b>Z</b>	0.4251
<b>One-Sided Pr &gt; Z</b>	0.3354
<b>Two-Sided Pr &gt;  Z </b>	0.6707

The NPAR1WAY Procedure

<b>Wilcoxon Scores (Rank Sums) for Variable cd4_diagnosis Classified by Variable HIV_HBV_vs_HIV_HCV</b>		
<b>HIV_HBV_vs_HIV_HCV</b>	<b>N</b>	<b>Sum of Scores</b>
<b>HIV_HBV</b>	37	1452.5
<b>HIV_HCV</b>	36	1248.5
<b>Average scores were used for ties.</b>		

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	582.50
<b>Normal Approximation</b>	
<b>Z</b>	0.9213
<b>One-Sided Pr &gt; Z</b>	0.1784
<b>Two-Sided Pr &gt;  Z </b>	0.3569

The UNIVARIATE Procedure  
 Variable: HIV\_viral\_load\_LOG (HIV\_viral\_load\_LOG)  
 HIV\_monoinfected\_

<b>Moments</b>			
<b>N</b>	1098	<b>Sum Weights</b>	1098
<b>Mean</b>	4.02112754		
<b>Std Deviation</b>	0.61235927	<b>Variance</b>	0.37498388
<b>Skewness</b>	-0.7940259	<b>Kurtosis</b>	0.56704082

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	4.99397
<b>99%</b>	4.96757
<b>95%</b>	4.87880
<b>90%</b>	4.75727
<b>75% Q3</b>	4.44988
<b>50% Median</b>	4.09915
<b>25% Q1</b>	3.64205
<b>10%</b>	3.11926
<b>5%</b>	2.85673
<b>1%</b>	2.26007
<b>0% Min</b>	1.66745

The UNIVARIATE Procedure

Variable: HIV\_viral\_load\_LOG (HIV\_viral\_load\_LOG)

HIV\_monoinfected\_vs\_HIV\_HBV=HIV\_HBV

<b>Moments</b>			
<b>N</b>	34	<b>Sum Weights</b>	34
<b>Mean</b>	3.80586445		
<b>Std Deviation</b>	1.09463913	<b>Variance</b>	1.19823482
<b>Skewness</b>	-0.69925	<b>Kurtosis</b>	-0.4538951

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	5.32314
<b>99%</b>	5.32314
<b>95%</b>	5.18959
<b>90%</b>	5.08189
<b>75% Q3</b>	4.49868
<b>50% Median</b>	4.11798
<b>25% Q1</b>	3.17026
<b>10%</b>	1.86332
<b>5%</b>	1.60206
<b>1%</b>	1.60206
<b>0% Min</b>	1.60206

<b>Wilcoxon Scores (Rank Sums) for Variable HIV_viral_load_LOG</b>		
<b>Classified by Variable HIV_monoinfected_vs_HIV_HBV</b>		
<b>HIV_monoinfected_vs_HIV_HBV</b>	<b>N</b>	<b>Sum of Scores</b>
<b>HIV_monoinfected</b>	1098	622804.0
<b>HIV_HBV</b>	34	18474.0

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	18474.0000
<b>Z</b>	-0.4189
<b>One-Sided Pr &lt; Z</b>	0.3376
<b>Two-Sided Pr &gt;  Z </b>	0.6753

The SAS System

The UNIVARIATE Procedure

Variable: HIV\_viral\_load\_LOG\_HIV\_HCV

<b>Moments</b>			
<b>N</b>	27	<b>Sum Weights</b>	27
<b>Mean</b>	4.236667		
<b>Std Deviation</b>	.9510237	<b>Variance</b>	.904446
<b>Skewness</b>	-.5194416	<b>Kurtosis</b>	3.514769

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	5.95
<b>99%</b>	5.95
<b>95%</b>	5.68
<b>90%</b>	5.68
<b>75% Q3</b>	4.81
<b>50% Median</b>	4.24
<b>25% Q1</b>	3.77
<b>10%</b>	2.83
<b>5%</b>	2.65
<b>1%</b>	1.7

<b>Wilcoxon Scores (Rank Sums) for Variable HIV_viral_load_LOG Classified by Variable HIV_monoinfected_vs_HIV_HCV</b>		
<b>HIV_monoinfected_vs_HIV_HCV</b>	<b>N</b>	<b>Sum of Scores</b>
<b>HIV_monoinfected</b>	1098	615581.5
<b>HIV_HCV</b>	27	17793.5

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	12230.50
<b>Normal Approximation</b>	
<b>Z</b>	1.5544
<b>One-Sided Pr &lt; Z</b>	0.0600
<b>Two-Sided Pr &gt;  Z </b>	0.1201

The SAS System

The UNIVARIATE Procedure

Variable: HIV\_viral\_load\_LOG HIV\_hepatitis

<b>Moments</b>			
<b>N</b>	63	<b>Sum Weights</b>	63
<b>Mean</b>	3.984444		
<b>Std Deviation</b>	1.042074	<b>Variance</b>	1.085919
<b>Skewness</b>	0.6399357	<b>Kurtosis</b>	2.940407

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	5.95
<b>99%</b>	5.95
<b>95%</b>	5.32
<b>90%</b>	5.14
<b>75% Q3</b>	4.70
<b>50% Median</b>	4.12
<b>25% Q1</b>	3.47
<b>10%</b>	2.65
<b>5%</b>	1.72
<b>1%</b>	1.6

**Wilcoxon Scores (Rank Sums) for Variable HIV\_viral\_load\_LOG  
Classified by Variable HIV\_monoinfected\_vs\_HIV\_hepatitis**

<b>HIV_monoinfected_vs_HIV_hepatitis</b>	<b>N</b>	<b>Sum of Scores</b>
<b>HIV_monoinfected</b>	1098	636424
<b>HIV_hepatitis</b>	63	38117

**Wilcoxon Two-Sample Test**

<b>Statistic</b>	33073
<b>Normal Approximation</b>	
<b>Z</b>	0.5850
<b>One-Sided Pr &lt; Z</b>	0.2793
<b>Two-Sided Pr &gt;  Z </b>	0.5586

**Wilcoxon Scores (Rank Sums) for Variable HIV\_viral\_load\_LOG  
Classified by Variable HIV\_HBV\_vs\_HIV\_HCV**

<b>HIV_HBV_vs_HIV_HCV</b>	<b>N</b>	<b>Sum of Scores</b>
<b>HIV_HBV</b>	30	777.0
<b>HIV_HCV</b>	27	876.0

**Wilcoxon Two-Sample Test**

<b>Statistic</b>	312.00
<b>Normal Approximation</b>	
<b>Z</b>	14863
<b>One-Sided Pr &lt; Z</b>	0.0686
<b>Two-Sided Pr &gt;  Z </b>	0.1372

**Table 4.** Effect of cART on CD4+ T-cell count, CD4/CD8 ratio and HIV viral load in HIV-coinfected patients

The SAS System

The UNIVARIATE Procedure  
 Variable: HBV (HBV)  
 cd\_4\_naïve

<b>Moments</b>			
<b>N</b>	24	<b>Sum Weights</b>	24
<b>Mean</b>	347.541667	<b>Sum Observations</b>	8341
<b>Std Deviation</b>	198.75569	<b>Variance</b>	39503.8243
<b>Skewness</b>	0.83695443	<b>Kurtosis</b>	0.84732151
<b>Uncorrected SS</b>	3807433	<b>Corrected SS</b>	908587.958
<b>Coeff Variation</b>	57.1890248	<b>Std Error Mean</b>	40.5708353

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	869.0
<b>99%</b>	869.0
<b>95%</b>	701.0
<b>90%</b>	595.0
<b>75% Q3</b>	465.5
<b>50% Median</b>	316.0
<b>25% Q1</b>	226.5
<b>10%</b>	128.0
<b>5%</b>	74.0
<b>1%</b>	30.0
<b>0% Min</b>	30.0

The UNIVARIATE Procedure

Variable: HBV (HBV)

Treat=cd\_4\_time\_12\_months

<b>Moments</b>			
<b>N</b>	24	<b>Sum Weights</b>	24
<b>Mean</b>	514.625	<b>Sum Observations</b>	12351
<b>Std Deviation</b>	230.574861	<b>Variance</b>	53164.7663
<b>Skewness</b>	1.16483499	<b>Kurtosis</b>	1.96882452
<b>Uncorrected SS</b>	7578923	<b>Corrected SS</b>	1222789.63
<b>Coeff Variation</b>	44.8044422	<b>Std Error Mean</b>	47.0658963

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1103.0
<b>99%</b>	1103.0
<b>95%</b>	1101.0
<b>90%</b>	678.0
<b>75% Q3</b>	612.5
<b>50% Median</b>	512.0
<b>25% Q1</b>	346.0
<b>10%</b>	246.0
<b>5%</b>	243.0
<b>1%</b>	194.0
<b>0% Min</b>	194.0



The SAS System  
The NPAR1WAY Procedure  
Treat=cd\_4 count

<b>Wilcoxon Scores (Rank Sums) for Variable HBV</b>					
<b>Classified by Variable time_0_vs_12</b>					
<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	24	462.0	588.0	48.497423	19.250
<b>12</b>	24	714.0	588.0	48.497423	29.750

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	462.0000
<b>Normal Approximation</b>	
<b>Z</b>	-2.5878
<b>One-Sided Pr &lt; Z</b>	0.0048
<b>Two-Sided Pr &gt;  Z </b>	0.0097
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0064
<b>Two-Sided Pr &gt;  Z </b>	0.0128
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	6.7500
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0094

The UNIVARIATE Procedure

Variable: HBV (HBV)

Treat=cd\_4\_time\_24\_months

<b>Moments</b>			
<b>N</b>	24	<b>Sum Weights</b>	24
<b>Mean</b>	597.5	<b>Sum Observations</b>	14340
<b>Std Deviation</b>	229.043759	<b>Variance</b>	52461.0435
<b>Skewness</b>	0.13367518	<b>Kurtosis</b>	-0.0836305
<b>Uncorrected SS</b>	9774754	<b>Corrected SS</b>	1206604
<b>Coeff Variation</b>	38.3336835	<b>Std Error Mean</b>	46.7533615

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1127.0
<b>99%</b>	1127.0
<b>95%</b>	908.0
<b>90%</b>	835.0
<b>75% Q3</b>	747.5
<b>50% Median</b>	613.0
<b>25% Q1</b>	423.0
<b>10%</b>	312.0
<b>5%</b>	280.0
<b>1%</b>	170.0
<b>0% Min</b>	170.0

The NPAR1WAY Procedure  
Treat=cd\_4 cell count

<b>Wilcoxon Scores (Rank Sums) for Variable HBV Classified by Variable time_0_vs_24</b>					
<b>time_0_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	24	414.50	588.0	48.494790	17.270833
<b>24</b>	24	761.50	588.0	48.494790	31.729167

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	414.5000
<b>Normal Approximation</b>	
<b>Z</b>	-3.5674
<b>One-Sided Pr &lt; Z</b>	0.0002
<b>Two-Sided Pr &gt;  Z </b>	0.0004
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0004
<b>Two-Sided Pr &gt;  Z </b>	0.0008
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	12.8000
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0003

The NPAR1WAY Procedure  
Treat=cd\_4\_count

**Wilcoxon Scores (Rank Sums) for Variable HBV  
Classified by Variable time\_12\_vs\_24**

<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>12</b>	24	503.50	588.0	48.494790	20.979167
<b>24</b>	24	672.50	588.0	48.494790	28.020833

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

**Statistic** 503.5000

**Normal Approximation**

**Z** -1.7321

**One-Sided Pr < Z** 0.0416

**Two-Sided Pr > |Z|** 0.0832

**t Approximation**

**One-Sided Pr < Z** 0.0449

**Two-Sided Pr > |Z|** 0.0898

**Z includes a continuity correction  
of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 3.0362

**DF** 1

**Pr > Chi-Square** 0.0814

The UNIVARIATE Procedure  
 Variable: HBV (HBV)  
 cd\_4:cd\_8\_naïve

<b>Moments</b>			
<b>N</b>	24	<b>Sum Weights</b>	24
<b>Mean</b>	0.40958333	<b>Sum Observations</b>	9.83
<b>Std Deviation</b>	0.35201876	<b>Variance</b>	0.12391721
<b>Skewness</b>	1.92805077	<b>Kurtosis</b>	4.49438498
<b>Uncorrected SS</b>	6.8763	<b>Corrected SS</b>	2.85009583
<b>Coeff Variation</b>	85.9455782	<b>Std Error Mean</b>	0.07185553

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.590
<b>99%</b>	1.590
<b>95%</b>	0.960
<b>90%</b>	0.900
<b>75% Q3</b>	0.585
<b>50% Median</b>	0.295
<b>25% Q1</b>	0.185
<b>10%</b>	0.120
<b>5%</b>	0.080
<b>1%</b>	0.040
<b>0% Min</b>	0.040

The UNIVARIATE Procedure  
 Variable: HBV (HBV)  
 Treat=cd\_4:cd\_8 time\_12

<b>Moments</b>			
<b>N</b>	24	<b>Sum Weights</b>	24
<b>Mean</b>	0.58375	<b>Sum Observations</b>	14.01
<b>Std Deviation</b>	0.40338581	<b>Variance</b>	0.16272011
<b>Skewness</b>	1.60015192	<b>Kurtosis</b>	2.77777703
<b>Uncorrected SS</b>	11.9209	<b>Corrected SS</b>	3.7425625
<b>Coeff Variation</b>	69.1024936	<b>Std Error Mean</b>	0.08234078

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.81
<b>99%</b>	1.81
<b>95%</b>	1.39
<b>90%</b>	1.08
<b>75% Q3</b>	0.70
<b>50% Median</b>	0.48
<b>25% Q1</b>	0.30
<b>10%</b>	0.20
<b>5%</b>	0.19
<b>1%</b>	0.11
<b>0% Min</b>	0.11

The NPAR1WAY Procedure  
Treat=cd\_4/cd\_8

<b>Wilcoxon Scores (Rank Sums) for Variable HBV Classified by Variable time_0_vs_12</b>					
<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	24	491.50	588.0	48.486892	20.479167
<b>12</b>	24	684.50	588.0	48.486892	28.520833

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	491.5000
<b>Normal Approximation</b>	
<b>Z</b>	-1.9799
<b>One-Sided Pr &lt; Z</b>	0.0239
<b>Two-Sided Pr &gt;  Z </b>	0.0477
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0268
<b>Two-Sided Pr &gt;  Z </b>	0.0536
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	3.9610
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0466

The UNIVARIATE Procedure

Variable: HBV (HBV)

Treat=cd\_4:cd\_8 time\_24\_months

<b>Moments</b>			
<b>N</b>	24	<b>Sum Weights</b>	24
<b>Mean</b>	0.71291667	<b>Sum Observations</b>	17.11
<b>Std Deviation</b>	0.35799537	<b>Variance</b>	0.12816069
<b>Skewness</b>	1.03779198	<b>Kurtosis</b>	1.85730225
<b>Uncorrected SS</b>	15.1457	<b>Corrected SS</b>	2.94769583
<b>Coeff Variation</b>	50.215599	<b>Std Error Mean</b>	0.0730755

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.740
<b>99%</b>	1.740
<b>95%</b>	1.390
<b>90%</b>	0.970
<b>75% Q3</b>	0.885
<b>50% Median</b>	0.655
<b>25% Q1</b>	0.480
<b>10%</b>	0.330
<b>5%</b>	0.230
<b>1%</b>	0.170
<b>0% Min</b>	0.170



The NPAR1WAY Procedure  
Treat=cd\_4/cd\_8

**Wilcoxon Scores (Rank Sums) for Variable HBV  
Classified by Variable time\_0\_vs\_24**

<b>time_0_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	24	430.0	588.0	48.481626	17.916667
<b>24</b>	24	746.0	588.0	48.481626	31.083333

Average scores were used for ties.

**Wilcoxon Two-Sample Test**

**Statistic** 430.0000

**Normal Approximation**

**Z** -3.2487

**One-Sided Pr < Z** 0.0006

**Two-Sided Pr > |Z|** 0.0012

**t Approximation**

**One-Sided Pr < Z** 0.0011

**Two-Sided Pr > |Z|** 0.0021

**Z includes a continuity correction of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 10.6209

**DF** 1

**Pr > Chi-Square** 0.0011

The NPAR1WAY Procedure  
Treat=cd\_4/cd\_8

**Wilcoxon Scores (Rank Sums) for Variable HBV  
Classified by Variable time\_12\_vs\_24**

<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
12	24	508.0	588.0	48.481626	21.166667
24	24	668.0	588.0	48.481626	27.833333

Average scores were used for ties.

**Wilcoxon Two-Sample Test**

**Statistic** 508.0000

**Normal Approximation**

**Z** -1.6398

**One-Sided Pr < Z** 0.0505

**Two-Sided Pr > |Z|** 0.1010

**t Approximation**

**One-Sided Pr < Z** 0.0539

**Two-Sided Pr > |Z|** 0.1077

**Z includes a continuity correction of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 2.7229

**DF** 1

**Pr > Chi-Square** 0.0989

The UNIVARIATE Procedure

Variable: HBV (HBV)

cv\_log\_naïve

<b>Moments</b>			
<b>N</b>	20	<b>Sum Weights</b>	20
<b>Mean</b>	3.90788235	<b>Sum Observations</b>	78.157647
<b>Std Deviation</b>	1.38726974	<b>Variance</b>	1.92451734
<b>Skewness</b>	-0.643462	<b>Kurtosis</b>	-1.0758625
<b>Uncorrected SS</b>	341.996718	<b>Corrected SS</b>	36.5658295
<b>Coeff Variation</b>	35.499271	<b>Std Error Mean</b>	0.31020295

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	5.49958
<b>99%</b>	5.49958
<b>95%</b>	5.46713
<b>90%</b>	5.37892
<b>75% Q3</b>	5.10267
<b>50% Median</b>	4.25391
<b>25% Q1</b>	2.64960
<b>10%</b>	1.69897
<b>5%</b>	1.69897
<b>1%</b>	1.69897
<b>0% Min</b>	1.69897

The UNIVARIATE Procedure

Variable: HBV (HBV)

Treat=cv\_log time\_12\_months

<b>Moments</b>			
<b>N</b>	24	<b>Sum Weights</b>	24
<b>Mean</b>	1.84097179	<b>Sum Observations</b>	44.183323
<b>Std Deviation</b>	0.36142809	<b>Variance</b>	0.13063026
<b>Skewness</b>	1.39137635	<b>Kurtosis</b>	2.04905513
<b>Uncorrected SS</b>	84.3447474	<b>Corrected SS</b>	3.00449606
<b>Coeff Variation</b>	19.6324621	<b>Std Error Mean</b>	0.0737762

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	2.80889
<b>99%</b>	2.80889
<b>95%</b>	2.60206
<b>90%</b>	2.55630
<b>75% Q3</b>	1.90224
<b>50% Median</b>	1.69897
<b>25% Q1</b>	1.69897
<b>10%</b>	1.65321
<b>5%</b>	1.30103
<b>1%</b>	1.30103
<b>0% Min</b>	1.30103

The NPAR1WAY Procedure  
Treat=cv\_log

<b>Wilcoxon Scores (Rank Sums) for Variable HBV Classified by Variable time_0_vs_12</b>					
<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	20	627.0	450.0	41.739148	31.3500
<b>12</b>	24	363.0	540.0	41.739148	15.1250

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	627.0000
<b>Normal Approximation</b>	
<b>Z</b>	4.2286
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>t Approximation</b>	
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	0.0001
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	17.9829
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	<.0001

The UNIVARIATE Procedure

Variable: HBV (HBV)

Treat=cv\_log time\_24\_months

<b>Moments</b>			
<b>N</b>	24	<b>Sum Weights</b>	24
<b>Mean</b>	1.64018853	<b>Sum Observations</b>	39.3645247
<b>Std Deviation</b>	0.34071471	<b>Variance</b>	0.11608651
<b>Skewness</b>	2.68138938	<b>Kurtosis</b>	10.8251556
<b>Uncorrected SS</b>	67.2352315	<b>Corrected SS</b>	2.6699898
<b>Coeff Variation</b>	20.7728992	<b>Std Error Mean</b>	0.0695481

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	2.99056
<b>99%</b>	2.99056
<b>95%</b>	1.83251
<b>90%</b>	1.75587
<b>75% Q3</b>	1.69897
<b>50% Median</b>	1.69897
<b>25% Q1</b>	1.39794
<b>10%</b>	1.30103
<b>5%</b>	1.30103
<b>1%</b>	1.30103
<b>0% Min</b>	1.30103

The NPAR1WAY Procedure  
Treat=cv\_log

**Wilcoxon Scores (Rank Sums) for Variable HBV  
Classified by Variable time\_0\_vs\_24**

<b>time_0_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	20	650.0	450.0	41.151502	32.500000
<b>24</b>	24	340.0	540.0	41.151502	14.166667

Average scores were used for ties.

**Wilcoxon Two-Sample Test**

**Statistic** 650.0000

**Normal Approximation**

**Z** 4.8479

**One-Sided Pr > Z** <.0001

**Two-Sided Pr > |Z|** <.0001

**t Approximation**

**One-Sided Pr > Z** <.0001

**Two-Sided Pr > |Z|** <.0001

**Z includes a continuity correction  
of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 23.6205

**DF** 1

**Pr > Chi-Square** <.0001

The NPAR1WAY Procedure  
Treat=cv\_log

**Wilcoxon Scores (Rank Sums) for Variable HBV  
Classified by Variable time\_12\_vs\_24**

<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
12	24	692.0	588.0	45.672283	28.833333
24	24	484.0	588.0	45.672283	20.166667

Average scores were used for ties.

**Wilcoxon Two-Sample Test**

**Statistic** 692.0000

**Normal Approximation**

**Z** 2.2661

**One-Sided Pr > Z** 0.0117

**Two-Sided Pr > |Z|** 0.0234

**t Approximation**

**One-Sided Pr > Z** 0.0140

**Two-Sided Pr > |Z|** 0.0281

**Z includes a continuity correction of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 5.1851

**DF** 1

**Pr > Chi-Square** 0.0228



The UNIVARIATE Procedure

Variable: HCV (HCV)

cd\_4\_naive

<b>Moments</b>			
<b>N</b>	23	<b>Sum Weights</b>	23
<b>Mean</b>	364.521739	<b>Sum Observations</b>	8384
<b>Std Deviation</b>	269.03884	<b>Variance</b>	72381.8972
<b>Skewness</b>	1.82338367	<b>Kurtosis</b>	3.35485997
<b>Uncorrected SS</b>	4648552	<b>Corrected SS</b>	1592401.74
<b>Coeff Variation</b>	73.8059794	<b>Std Error Mean</b>	56.098476

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1171
<b>99%</b>	1171
<b>95%</b>	977
<b>90%</b>	697
<b>75% Q3</b>	478
<b>50% Median</b>	304
<b>25% Q1</b>	154
<b>10%</b>	141
<b>5%</b>	140
<b>1%</b>	67
<b>0% Min</b>	67

The UNIVARIATE Procedure  
 Variable: HCV (HCV)  
 Treat=cd\_4 time\_12\_months

<b>Moments</b>			
<b>N</b>	23	<b>Sum Weights</b>	23
<b>Mean</b>	407.934633	<b>Sum Observations</b>	9382.49657
<b>Std Deviation</b>	208.678839	<b>Variance</b>	43546.8577
<b>Skewness</b>	1.43441038	<b>Kurtosis</b>	1.88947398
<b>Uncorrected SS</b>	4785476.17	<b>Corrected SS</b>	958030.87
<b>Coeff Variation</b>	51.1549699	<b>Std Error Mean</b>	43.5125458

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1009.33
<b>99%</b>	1009.33
<b>95%</b>	786.00
<b>90%</b>	706.00
<b>75% Q3</b>	555.50
<b>50% Median</b>	344.00
<b>25% Q1</b>	252.00
<b>10%</b>	209.00
<b>5%</b>	208.00
<b>1%</b>	191.00
<b>0% Min</b>	191.00

The NPAR1WAY Procedure  
Treat=cd\_4

**Wilcoxon Scores (Rank Sums) for Variable HCV  
Classified by Variable time\_0\_vs\_12**

<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	23	477.50	540.50	45.516908	20.760870
<b>12</b>	23	603.50	540.50	45.516908	26.239130

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

<b>Statistic</b>	477.5000
<b>Normal Approximation</b>	
<b>Z</b>	-1.3731
<b>One-Sided Pr &lt; Z</b>	0.0849
<b>Two-Sided Pr &gt;  Z </b>	0.1697
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0883
<b>Two-Sided Pr &gt;  Z </b>	0.1765
<b>Z includes a continuity correction of 0.5.</b>	

**Kruskal-Wallis Test**

<b>Chi-Square</b>	1.9157
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.1663

The UNIVARIATE Procedure  
 Variable: HCV (HCV)  
 Treat=cd\_4 time\_24\_months

<b>Moments</b>			
<b>N</b>	23	<b>Sum Weights</b>	23
<b>Mean</b>	521.094893	<b>Sum Observations</b>	11985.1825
<b>Std Deviation</b>	179.657242	<b>Variance</b>	32276.7245
<b>Skewness</b>	1.35136353	<b>Kurtosis</b>	2.22099672
<b>Uncorrected SS</b>	6955505.35	<b>Corrected SS</b>	710087.94
<b>Coeff Variation</b>	34.4768763	<b>Std Error Mean</b>	37.4611245

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1029.50
<b>99%</b>	1029.50
<b>95%</b>	898.25
<b>90%</b>	755.00
<b>75% Q3</b>	561.00
<b>50% Median</b>	485.00
<b>25% Q1</b>	406.00
<b>10%</b>	357.00
<b>5%</b>	312.60
<b>1%</b>	260.00
<b>0% Min</b>	260.00

The NPAR1WAY Procedure  
Treat=cd\_4

<b>Wilcoxon Scores (Rank Sums) for Variable HCV Classified by Variable time_0_vs_24</b>					
<b>time_0_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	23	389.0	540.50	45.518311	16.913043
<b>24</b>	23	692.0	540.50	45.518311	30.086957

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	389.0000
<b>Normal Approximation</b>	
<b>Z</b>	-3.3173
<b>One-Sided Pr &lt; Z</b>	0.0005
<b>Two-Sided Pr &gt;  Z </b>	0.0009
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0009
<b>Two-Sided Pr &gt;  Z </b>	0.0018
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	11.0778
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0009

The NPAR1WAY Procedure  
Treat=cd\_4

<b>Wilcoxon Scores (Rank Sums) for Variable HCV Classified by Variable time_12_vs_24</b>					
<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>12</b>	23	430.0	540.50	45.518311	18.695652
<b>24</b>	23	651.0	540.50	45.518311	28.304348

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	430.0000
<b>Normal Approximation</b>	
<b>Z</b>	-2.4166
<b>One-Sided Pr &lt; Z</b>	0.0078
<b>Two-Sided Pr &gt;  Z </b>	0.0157
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0099
<b>Two-Sided Pr &gt;  Z </b>	0.0198
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	5.8932
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0152

The UNIVARIATE Procedure

Variable: HCV (HCV)

cd\_4:cd\_8\_naïve

<b>Moments</b>			
<b>N</b>	21	<b>Sum Weights</b>	21
<b>Mean</b>	0.37952381	<b>Sum Observations</b>	7.97
<b>Std Deviation</b>	0.30292699	<b>Variance</b>	0.09176476
<b>Skewness</b>	1.4991632	<b>Kurtosis</b>	1.90482872
<b>Uncorrected SS</b>	4.8601	<b>Corrected SS</b>	1.83529524
<b>Coeff Variation</b>	79.8176513	<b>Std Error Mean</b>	0.06610409

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.20
<b>99%</b>	1.20
<b>95%</b>	0.93
<b>90%</b>	0.93
<b>75% Q3</b>	0.47
<b>50% Median</b>	0.32
<b>25% Q1</b>	0.16
<b>10%</b>	0.13
<b>5%</b>	0.06
<b>1%</b>	0.04
<b>0% Min</b>	0.04

The UNIVARIATE Procedure

Variable: HCV (HCV)

Treat=cd\_4:cd\_8\_time\_12\_months

<b>Moments</b>			
<b>N</b>	23	<b>Sum Weights</b>	23
<b>Mean</b>	0.46947705	<b>Sum Observations</b>	10.797972
<b>Std Deviation</b>	0.25162292	<b>Variance</b>	0.06331409
<b>Skewness</b>	1.14999088	<b>Kurtosis</b>	2.20756216
<b>Uncorrected SS</b>	6.46231007	<b>Corrected SS</b>	1.39291006
<b>Coeff Variation</b>	53.5964265	<b>Std Error Mean</b>	0.05246701

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.202538
<b>99%</b>	1.202538
<b>95%</b>	0.929559
<b>90%</b>	0.697546
<b>75% Q3</b>	0.561721
<b>50% Median</b>	0.481043
<b>25% Q1</b>	0.288509
<b>10%</b>	0.181119
<b>5%</b>	0.167385
<b>1%</b>	0.105423
<b>0% Min</b>	0.105423



The NPAR1WAY Procedure  
Treat=cd\_4\_8

**Wilcoxon Scores (Rank Sums) for Variable HCV  
Classified by Variable time\_0\_vs\_12**

<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	21	395.0	472.50	42.555784	18.809524
<b>12</b>	23	595.0	517.50	42.555784	25.869565

Average scores were used for ties.

**Wilcoxon Two-Sample Test**

**Statistic** 395.0000

**Normal Approximation**

**Z** -1.8094

**One-Sided Pr < Z** 0.0352

**Two-Sided Pr > |Z|** 0.0704

**t Approximation**

**One-Sided Pr < Z** 0.0387

**Two-Sided Pr > |Z|** 0.0774

**Z includes a continuity correction  
of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 3.3165

**DF** 1

**Pr > Chi-Square** 0.0686

The UNIVARIATE Procedure

Variable: HCV (HCV)

Treat=cd\_4:cd\_8\_time\_24\_months

<b>Moments</b>			
<b>N</b>	23	<b>Sum Weights</b>	23
<b>Mean</b>	0.5903869	<b>Sum Observations</b>	13.5788988
<b>Std Deviation</b>	0.24704325	<b>Variance</b>	0.06103037
<b>Skewness</b>	1.09898667	<b>Kurtosis</b>	2.00357549
<b>Uncorrected SS</b>	9.3594721	<b>Corrected SS</b>	1.34266812
<b>Coeff Variation</b>	41.8442977	<b>Std Error Mean</b>	0.05151208

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.313138
<b>99%</b>	1.313138
<b>95%</b>	0.935780
<b>90%</b>	0.872832
<b>75% Q3</b>	0.708029
<b>50% Median</b>	0.496272
<b>25% Q1</b>	0.433754
<b>10%</b>	0.332751
<b>5%</b>	0.317933
<b>1%</b>	0.185464
<b>0% Min</b>	0.185464

The NPAR1WAY Procedure  
Treat=cd\_4\_8

**Wilcoxon Scores (Rank Sums) for Variable HCV  
Classified by Variable time\_0\_vs\_24**

<b>time_0_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	21	348.0	472.50	42.555784	16.571429
<b>24</b>	23	642.0	517.50	42.555784	27.913043

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

<b>Statistic</b>	348.0000
<b>Normal Approximation</b>	
<b>Z</b>	-2.9138
<b>One-Sided Pr &lt; Z</b>	0.0018
<b>Two-Sided Pr &gt;  Z </b>	0.0036
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0028
<b>Two-Sided Pr &gt;  Z </b>	0.0056
<b>Z includes a continuity correction of 0.5.</b>	

**Kruskal-Wallis Test**

<b>Chi-Square</b>	8.5590
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0034

The NPAR1WAY Procedure  
Treat=cd\_4\_8

**Wilcoxon Scores (Rank Sums) for Variable HCV  
Classified by Variable time\_12\_vs\_24**

<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>12</b>	23	463.0	540.50	45.518311	20.130435
<b>24</b>	23	618.0	540.50	45.518311	26.869565

**Wilcoxon Two-Sample Test**

**Statistic** 463.0000

**Normal Approximation**

**Z** -1.6916

**One-Sided Pr < Z** 0.0454

**Two-Sided Pr > |Z|** 0.0907

**t Approximation**

**One-Sided Pr < Z** 0.0488

**Two-Sided Pr > |Z|** 0.0976

**Z includes a continuity correction  
of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 2.8989

**DF** 1

**Pr > Chi-Square** 0.0886

The UNIVARIATE Procedure  
 Variable: HCV (HCV)  
 cv\_log\_naïve

<b>Moments</b>			
<b>N</b>	19	<b>Sum Weights</b>	19
<b>Mean</b>	3.84786664	<b>Sum Observations</b>	73.1094661
<b>Std Deviation</b>	1.63562441	<b>Variance</b>	2.6752672
<b>Skewness</b>	-0.1319716	<b>Kurtosis</b>	-1.4781717
<b>Uncorrected SS</b>	329.470285	<b>Corrected SS</b>	48.1548095
<b>Coeff Variation</b>	42.5073049	<b>Std Error Mean</b>	0.37523797

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	6.00000
<b>99%</b>	6.00000
<b>95%</b>	6.00000
<b>90%</b>	6.00000
<b>75% Q3</b>	5.35218
<b>50% Median</b>	4.05258
<b>25% Q1</b>	1.69897
<b>10%</b>	1.69897
<b>5%</b>	1.69897
<b>1%</b>	1.69897
<b>0% Min</b>	1.69897

The SAS System

The UNIVARIATE Procedure  
Variable: HCV (HCV)  
Treat=cv\_log time\_12\_months

<b>Moments</b>			
<b>N</b>	23	<b>Sum Weights</b>	23
<b>Mean</b>	1.8309835	<b>Sum Observations</b>	42.1126206
<b>Std Deviation</b>	0.51657329	<b>Variance</b>	0.26684797
<b>Skewness</b>	2.457642	<b>Kurtosis</b>	6.03110656
<b>Uncorrected SS</b>	82.9781689	<b>Corrected SS</b>	5.87065527
<b>Coeff Variation</b>	28.2128862	<b>Std Error Mean</b>	0.10771298

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	3.58422
<b>99%</b>	3.58422
<b>95%</b>	2.83948
<b>90%</b>	2.73640
<b>75% Q3</b>	1.69897
<b>50% Median</b>	1.69897
<b>25% Q1</b>	1.69897
<b>10%</b>	1.39794
<b>5%</b>	1.39794
<b>1%</b>	1.30103
<b>0% Min</b>	1.30103

The NPAR1WAY Procedure  
Treat=cv\_log

<b>Wilcoxon Scores (Rank Sums) for Variable HCV Classified by Variable time_0_vs_12</b>					
<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	19	567.0	408.50	37.016974	29.842105
<b>12</b>	23	336.0	494.50	37.016974	14.608696

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	567.0000
<b>Normal Approximation</b>	
<b>Z</b>	4.2683
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>t Approximation</b>	
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	0.0001
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	18.3340
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	<.0001

The UNIVARIATE Procedure

Variable: HCV (HCV)

Treat=cv\_log time\_24\_months

<b>Moments</b>			
<b>N</b>	23	<b>Sum Weights</b>	23
<b>Mean</b>	1.63978574	<b>Sum Observations</b>	37.7150719
<b>Std Deviation</b>	0.32821062	<b>Variance</b>	0.10772221
<b>Skewness</b>	2.58708864	<b>Kurtosis</b>	10.3701658
<b>Uncorrected SS</b>	64.2145257	<b>Corrected SS</b>	2.36988866
<b>Coeff Variation</b>	20.0154577	<b>Std Error Mean</b>	0.06843665

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	2.91116
<b>99%</b>	2.91116
<b>95%</b>	1.77085
<b>90%</b>	1.69897
<b>75% Q3</b>	1.69897
<b>50% Median</b>	1.69897
<b>25% Q1</b>	1.39794
<b>10%</b>	1.30103
<b>5%</b>	1.30103
<b>1%</b>	1.30103
<b>0% Min</b>	1.30103



The NPAR1WAY Procedure  
Treat=cv\_log

<b>Wilcoxon Scores (Rank Sums) for Variable HCV Classified by Variable time_0_vs_24</b>					
<b>time_0_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	19	583.50	408.50	37.949589	30.710526
<b>24</b>	23	319.50	494.50	37.949589	13.891304

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	583.5000
<b>Normal Approximation</b>	
<b>Z</b>	4.5982
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>t Approximation</b>	
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	21.2648
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	<.0001

The NPAR1WAY Procedure  
Treat=cv\_log

**Wilcoxon Scores (Rank Sums) for Variable HCV  
Classified by Variable time\_12\_vs\_24**

<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
12	23	597.50	540.50	39.336864	25.978261
24	23	483.50	540.50	39.336864	21.021739

Average scores were used for ties.

**Wilcoxon Two-Sample Test**

**Statistic** 597.5000

**Normal Approximation**

**Z** 1.4363

**One-Sided Pr > Z** 0.0755

**Two-Sided Pr > |Z|** 0.1509

**t Approximation**

**One-Sided Pr > Z** 0.0789

**Two-Sided Pr > |Z|** 0.1578

**Z includes a continuity correction  
of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 2.0997

**DF** 1

**Pr > Chi-Square** 0.1473

The UNIVARIATE Procedure

Variable: hepatitis (hepatitis)

cd\_4\_naïve

<b>Moments</b>			
<b>N</b>	47	<b>Sum Weights</b>	47
<b>Mean</b>	353.893617	<b>Sum Observations</b>	16633
<b>Std Deviation</b>	235.479182	<b>Variance</b>	55450.445
<b>Skewness</b>	1.50883137	<b>Kurtosis</b>	2.83466909
<b>Uncorrected SS</b>	8437033	<b>Corrected SS</b>	2550720.47
<b>Coeff Variation</b>	66.5395391	<b>Std Error Mean</b>	34.3481688

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1171
<b>99%</b>	1171
<b>95%</b>	869
<b>90%</b>	697
<b>75% Q3</b>	467
<b>50% Median</b>	312
<b>25% Q1</b>	202
<b>10%</b>	128
<b>5%</b>	67
<b>1%</b>	30
<b>0% Min</b>	30

The UNIVARIATE Procedure  
 Variable: hepatitis (hepatitis)  
 Treat=cd\_4\_time\_12\_months

<b>Moments</b>			
<b>N</b>	47	<b>Sum Weights</b>	47
<b>Mean</b>	462.387289	<b>Sum Observations</b>	21732.2026
<b>Std Deviation</b>	224.311503	<b>Variance</b>	50315.6504
<b>Skewness</b>	1.20229569	<b>Kurtosis</b>	1.52505047
<b>Uncorrected SS</b>	12363214.1	<b>Corrected SS</b>	2314519.92
<b>Coeff Variation</b>	48.5116067	<b>Std Error Mean</b>	32.7191955

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1103.000
<b>99%</b>	1103.000
<b>95%</b>	1009.330
<b>90%</b>	706.000
<b>75% Q3</b>	568.500
<b>50% Median</b>	435.333
<b>25% Q1</b>	266.330
<b>10%</b>	233.000
<b>5%</b>	208.000
<b>1%</b>	191.000
<b>0% Min</b>	191.000

The NPAR1WAY Procedure  
Treat=cd\_4

**Wilcoxon Scores (Rank Sums) for Variable hepatitis  
Classified by Variable time\_0\_vs\_12**

<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	47	1877.0	2232.50	132.240932	39.936170
<b>12</b>	47	2588.0	2232.50	132.240932	55.063830

Average scores were used for ties.

**Wilcoxon Two-Sample Test**

**Statistic** 1877.0000

**Normal Approximation**

**Z** -2.6845

**One-Sided Pr < Z** 0.0036

**Two-Sided Pr > |Z|** 0.0073

**t Approximation**

**One-Sided Pr < Z** 0.0043

**Two-Sided Pr > |Z|** 0.0086

**Z includes a continuity correction  
of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 7.2268

**DF** 1

**Pr > Chi-Square** 0.0072

The UNIVARIATE Procedure  
 Variable: hepatitis (hepatitis)  
 Treat=cd\_4\_time\_24\_months

<b>Moments</b>			
<b>N</b>	47	<b>Sum Weights</b>	47
<b>Mean</b>	560.042394	<b>Sum Observations</b>	26321.9925
<b>Std Deviation</b>	207.748795	<b>Variance</b>	43159.562
<b>Skewness</b>	0.61617295	<b>Kurtosis</b>	0.24658514
<b>Uncorrected SS</b>	16726771.6	<b>Corrected SS</b>	1985339.85
<b>Coeff Variation</b>	37.0951909	<b>Std Error Mean</b>	30.3032763

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1127.00
<b>99%</b>	1127.00
<b>95%</b>	908.00
<b>90%</b>	834.66
<b>75% Q3</b>	662.00
<b>50% Median</b>	554.00
<b>25% Q1</b>	406.00
<b>10%</b>	312.60
<b>5%</b>	279.50
<b>1%</b>	170.00
<b>0% Min</b>	170.00

The NPAR1WAY Procedure  
Treat=cd\_4

**Wilcoxon Scores (Rank Sums) for Variable hepatitis  
Classified by Variable time\_0\_vs\_24**

time_0_vs_24	N	Sum of Scores	Expected Under H0	Std Dev Under H0	Mean Score
0	47	1596.0	2232.50	132.241409	33.957447
24	47	2869.0	2232.50	132.241409	61.042553

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

<b>Statistic</b>	1596.0000
<b>Normal Approximation</b>	
<b>Z</b>	-4.8094
<b>One-Sided Pr &lt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>Z includes a continuity correction of 0.5.</b>	

**Kruskal-Wallis Test**

<b>Chi-Square</b>	23.1666
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	<.0001

The NPAR1WAY Procedure  
Treat=cd\_4

**Wilcoxon Scores (Rank Sums) for Variable hepatitis  
Classified by Variable time\_12\_vs\_24**

<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>12</b>	47	1903.50	2232.50	132.240932	40.50
<b>24</b>	47	2561.50	2232.50	132.240932	54.50

Average scores were used for ties.

**Wilcoxon Two-Sample Test**

**Statistic** 1903.5000

**Normal Approximation**

**Z** -2.4841

**One-Sided Pr < Z** 0.0065

**Two-Sided Pr > |Z|** 0.0130

**t Approximation**

**One-Sided Pr < Z** 0.0074

**Two-Sided Pr > |Z|** 0.0148

**Z includes a continuity correction  
of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 6.1896

**DF** 1

**Pr > Chi-Square** 0.0129



The UNIVARIATE Procedure  
 Variable: hepatitis (hepatitis)  
 cd\_4:cd\_8\_naïve

<b>Moments</b>			
<b>N</b>	47	<b>Sum Weights</b>	47
<b>Mean</b>	0.39957447	<b>Sum Observations</b>	18.78
<b>Std Deviation</b>	0.3236474	<b>Variance</b>	0.10474764
<b>Skewness</b>	1.69613778	<b>Kurtosis</b>	3.2478416
<b>Uncorrected SS</b>	12.3224	<b>Corrected SS</b>	4.81839149
<b>Coeff Variation</b>	80.9980187	<b>Std Error Mean</b>	0.04720883

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.59
<b>99%</b>	1.59
<b>95%</b>	0.96
<b>90%</b>	0.93
<b>75% Q3</b>	0.50
<b>50% Median</b>	0.30
<b>25% Q1</b>	0.17
<b>10%</b>	0.12
<b>5%</b>	0.06
<b>1%</b>	0.04
<b>0% Min</b>	0.04

The UNIVARIATE Procedure  
 Variable: hepatitis (hepatitis)  
 Treat=cd\_4:cd\_8\_time\_12\_months

<b>Moments</b>			
<b>N</b>	47	<b>Sum Weights</b>	47
<b>Mean</b>	0.52791123	<b>Sum Observations</b>	24.8118277
<b>Std Deviation</b>	0.33865149	<b>Variance</b>	0.11468483
<b>Skewness</b>	1.7241357	<b>Kurtosis</b>	3.90472742
<b>Uncorrected SS</b>	18.3739449	<b>Corrected SS</b>	5.2755024
<b>Coeff Variation</b>	64.1493259	<b>Std Error Mean</b>	0.0493974

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.807882
<b>99%</b>	1.807882
<b>95%</b>	1.202538
<b>90%</b>	0.992703
<b>75% Q3</b>	0.632539
<b>50% Median</b>	0.481043
<b>25% Q1</b>	0.291814
<b>10%</b>	0.188291
<b>5%</b>	0.167385
<b>1%</b>	0.105423
<b>0% Min</b>	0.105423

The NPAR1WAY Procedure  
Treat=cd\_4\_8

**Wilcoxon Scores (Rank Sums) for Variable hepatitis  
Classified by Variable time\_0\_vs\_12**

<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	47	1901.0	2232.50	132.236154	40.446809
<b>12</b>	47	2564.0	2232.50	132.236154	54.553191

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

<b>Statistic</b>	1901.0000
<b>Normal Approximation</b>	
<b>Z</b>	-2.5031
<b>One-Sided Pr &lt; Z</b>	0.0062
<b>Two-Sided Pr &gt;  Z </b>	0.0123
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0070
<b>Two-Sided Pr &gt;  Z </b>	0.0141
<b>Z includes a continuity correction of 0.5.</b>	

**Kruskal-Wallis Test**

<b>Chi-Square</b>	6.2844
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0122

The UNIVARIATE Procedure

Variable: hepatitis (hepatitis)

Treat=cd\_4:cd\_8 time\_24\_months

<b>Moments</b>			
<b>N</b>	47	<b>Sum Weights</b>	47
<b>Mean</b>	0.65347821	<b>Sum Observations</b>	30.713476
<b>Std Deviation</b>	0.31192005	<b>Variance</b>	0.09729412
<b>Skewness</b>	1.20759219	<b>Kurtosis</b>	2.36382749
<b>Uncorrected SS</b>	24.5461169	<b>Corrected SS</b>	4.4755295
<b>Coeff Variation</b>	47.7322804	<b>Std Error Mean</b>	0.04549822

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	1.739198
<b>99%</b>	1.739198
<b>95%</b>	1.313138
<b>90%</b>	0.964548
<b>75% Q3</b>	0.855786
<b>50% Median</b>	0.614629
<b>25% Q1</b>	0.458201
<b>10%</b>	0.331034
<b>5%</b>	0.232814
<b>1%</b>	0.173293
<b>0% Min</b>	0.173293

The NPAR1WAY Procedure  
Treat=cd\_4\_8

**Wilcoxon Scores (Rank Sums) for Variable hepatitis  
Classified by Variable time\_0\_24**

<b>time_0_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	47	1650.0	2232.50	132.236154	35.106383
<b>24</b>	47	2815.0	2232.50	132.236154	59.893617

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

**Statistic** 1650.0000

**Normal Approximation**

**Z** -4.4012

**One-Sided Pr < Z** <.0001

**Two-Sided Pr > |Z|** <.0001

**t Approximation**

**One-Sided Pr < Z** <.0001

**Two-Sided Pr > |Z|** <.0001

**Z includes a continuity correction  
of 0.5.**

**Kruskal-Wallis Test**

**Chi-Square** 19.4040

**DF** 1

**Pr > Chi-Square** <.0001

The NPAR1WAY Procedure  
Treat=cd\_4\_8

<b>Wilcoxon Scores (Rank Sums) for Variable hepatitis Classified by Variable time_12_vs_24</b>					
<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>12</b>	47	1912.0	2232.50	132.241887	40.680851
<b>24</b>	47	2553.0	2232.50	132.241887	54.319149

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	1912.0000
<b>Normal Approximation</b>	
<b>Z</b>	-2.4198
<b>One-Sided Pr &lt; Z</b>	0.0078
<b>Two-Sided Pr &gt;  Z </b>	0.0155
<b>t Approximation</b>	
<b>One-Sided Pr &lt; Z</b>	0.0087
<b>Two-Sided Pr &gt;  Z </b>	0.0175
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	5.8738
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0154

The UNIVARIATE Procedure  
 Variable: hepatitis (hepatitis)  
 cv\_log\_naïve

<b>Moments</b>			
<b>N</b>	39	<b>Sum Weights</b>	39
<b>Mean</b>	3.20837	<b>Sum Observations</b>	77.0008801
<b>Std Deviation</b>	1.1555229	<b>Variance</b>	1.33523317
<b>Skewness</b>	-0.2632938	<b>Kurtosis</b>	-1.6299519
<b>Uncorrected SS</b>	277.757677	<b>Corrected SS</b>	30.7103629
<b>Coeff Variation</b>	36.0158865	<b>Std Error Mean</b>	0.23587012

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	4.69512
<b>99%</b>	4.69512
<b>95%</b>	4.60977
<b>90%</b>	4.53667
<b>75% Q3</b>	4.21188
<b>50% Median</b>	3.55601
<b>25% Q1</b>	1.69897
<b>10%</b>	1.69897
<b>5%</b>	1.69897
<b>1%</b>	1.69897
<b>0% Min</b>	1.69897

The UNIVARIATE Procedure  
 Variable: hepatitis (hepatitis)  
 Treat=cv\_log time\_12\_months

<b>Moments</b>			
<b>N</b>	47	<b>Sum Weights</b>	47
<b>Mean</b>	1.83608391	<b>Sum Observations</b>	86.2959436
<b>Std Deviation</b>	0.43927616	<b>Variance</b>	0.19296354
<b>Skewness</b>	2.18154879	<b>Kurtosis</b>	5.34636267
<b>Uncorrected SS</b>	167.322916	<b>Corrected SS</b>	8.87632305
<b>Coeff Variation</b>	23.9246234	<b>Std Error Mean</b>	0.06407501

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	3.58422
<b>99%</b>	3.58422
<b>95%</b>	2.80889
<b>90%</b>	2.60206
<b>75% Q3</b>	1.77815
<b>50% Median</b>	1.69897
<b>25% Q1</b>	1.69897
<b>10%</b>	1.39794
<b>5%</b>	1.30103
<b>1%</b>	1.30103
<b>0% Min</b>	1.30103



The NPAR1WAY Procedure  
Treat=cv\_log

**Wilcoxon Scores (Rank Sums) for Variable hepatitis  
Classified by Variable time\_0\_vs\_12**

<b>time_0_vs_12</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	39	1233.0	864.0	78.027963	51.375000
<b>12</b>	47	1323.0	1692.0	78.027963	28.148936

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

<b>Statistic</b>	1233.0000
<b>Normal Approximation</b>	
<b>Z</b>	4.7227
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>t Approximation</b>	
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>Z includes a continuity correction of 0.5.</b>	

**Kruskal-Wallis Test**

<b>Chi-Square</b>	22.3641
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	<.0001

The UNIVARIATE Procedure

Variable: hepatitis (hepatitis)

Treat=cv\_log time\_24\_months

<b>Moments</b>			
<b>N</b>	47	<b>Sum Weights</b>	47
<b>Mean</b>	1.63999142	<b>Sum Observations</b>	77.0795966
<b>Std Deviation</b>	0.33100244	<b>Variance</b>	0.10956262
<b>Skewness</b>	2.55031731	<b>Kurtosis</b>	9.3591623
<b>Uncorrected SS</b>	131.449757	<b>Corrected SS</b>	5.03988036
<b>Coeff Variation</b>	20.1831814	<b>Std Error Mean</b>	0.04828167

<b>Quantiles (Definition 5)</b>	
<b>Quantile</b>	<b>Estimate</b>
<b>100% Max</b>	2.99056
<b>99%</b>	2.99056
<b>95%</b>	1.83251
<b>90%</b>	1.75587
<b>75% Q3</b>	1.69897
<b>50% Median</b>	1.69897
<b>25% Q1</b>	1.39794
<b>10%</b>	1.30103
<b>5%</b>	1.30103
<b>1%</b>	1.30103
<b>0% Min</b>	1.30103

The NPAR1WAY Procedure  
Treat=cv\_log

**Wilcoxon Scores (Rank Sums) for Variable hepatitis  
Classified by Variable time\_0\_vs\_24**

<b>time_0_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>0</b>	39	1296.0	864.0	77.896960	54.000000
<b>24</b>	47	1260.0	1692.0	77.896960	26.808511

**Average scores were used for ties.**

**Wilcoxon Two-Sample Test**

<b>Statistic</b>	1296.0000
<b>Normal Approximation</b>	
<b>Z</b>	5.5394
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>t Approximation</b>	
<b>One-Sided Pr &gt; Z</b>	<.0001
<b>Two-Sided Pr &gt;  Z </b>	<.0001
<b>Z includes a continuity correction of 0.5.</b>	

**Kruskal-Wallis Test**

<b>Chi-Square</b>	30.7558
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	<.0001

The NPAR1WAY Procedure  
Treat=cv\_log

<b>Wilcoxon Scores (Rank Sums) for Variable hepatitis Classified by Variable time_12_vs_24</b>					
<b>time_12_vs_24</b>	<b>N</b>	<b>Sum of Scores</b>	<b>Expected Under H0</b>	<b>Std Dev Under H0</b>	<b>Mean Score</b>
<b>12</b>	47	2552.0	2232.50	120.312575	54.297872
<b>24</b>	47	1913.0	2232.50	120.312575	40.702128

**Average scores were used for ties.**

<b>Wilcoxon Two-Sample Test</b>	
<b>Statistic</b>	2552.0000
<b>Normal Approximation</b>	
<b>Z</b>	2.6514
<b>One-Sided Pr &gt; Z</b>	0.0040
<b>Two-Sided Pr &gt;  Z </b>	0.0080
<b>t Approximation</b>	
<b>One-Sided Pr &gt; Z</b>	0.0047
<b>Two-Sided Pr &gt;  Z </b>	0.0094
<b>Z includes a continuity correction of 0.5.</b>	

<b>Kruskal-Wallis Test</b>	
<b>Chi-Square</b>	7.0521
<b>DF</b>	1
<b>Pr &gt; Chi-Square</b>	0.0079

Frequency Col Pct	Table of cd4_diagnosis_transformed by naïve			
	<200	200-500	>500	Total
<b>HIV-HBV</b>	5 20.84	15 62.50	4 16.66	24
<b>HIV-HCV</b>	6 26.09	13 56.52	4 17.39	23
<b>HIV-hepatitis</b>	11 23.41	28 59.57	8 17.02	47
<b>Total</b>	22	56	16	94

Frequency Col Pct	Table of cd4_diagnosis_transformed by 12 months			
	<200	200-500	>500	Total
<b>HIV-HBV</b>	1 4.17	11 45.83	12 50.00	24
<b>HIV-HCV</b>	1 4.17	16 69.57	6 26.08	23
<b>HIV-hepatitis</b>	2 4.26	27 57.45	18 38.29	47
<b>Total</b>	4	54	36	94

Frequency Col Pct	Table of cd4_diagnosis_transformed by 24 months			
	<200	200-500	>500	Total
<b>HIV-HBV</b>	1 4.17	7 29.17	16 66.66	24
<b>HIV-HCV</b>	0 0.00	12 52.17	11 47.83	23
<b>HIV-hepatitis</b>	1 2.13	19 40.43	27 57.44	47
<b>Total</b>	4	54	36	94

Statistics for Table of cd4\_diagnosis  
HIV-HBV time naïve vs 12

Statistic	DF	Value	Prob
<b>Chi-Square</b>	2	7.282	0.0262

Statistics for Table of cd4\_diagnosis

HIV-HBV time naïve vs 24

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	12.776	0.0017

Statistics for Table of cd4\_diagnosis

HIV-HBV time 12 vs 24

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	1.460	0.4818

Statistics for Table of cd4\_diagnosis

HIV-HCV time naïve vs 12

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	4.282	0.1176

Statistics for Table of cd4\_diagnosis

HIV-HCV time naïve vs 24

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	9.307	0.0095

Statistics for Table of cd4\_diagnosis

HIV-HCV time 12 vs 24

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	3.042	0.2185

Statistics for Table of cd4\_diagnosis

hepatitis time naïve vs 12

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	10.095	0.0064

Statistics for Table of cd4\_diagnosis

hepatitis time naïve vs 24

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	20.371	<.0001

Statistics for Table of cd4\_diagnosis

hepatitis time 12 vs 24

<b>Statistic</b>	<b>DF</b>	<b>Value</b>	<b>Prob</b>
<b>Chi-Square</b>	2	3.525	0.1716