CROSSTABS
/TABLES=diagPca BY ch6983
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ RISK CMH(1)
/CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

#### Notes

	0	00 1:- 0040 40:40:40
	Output Created	22-lip-2012 12:12:42
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=diagPca BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.031
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	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Va	Valid Missing				tal
	N	N Percent		Percent	N	Percent
diagPca * ch6983	300	75,0%	100	25,0%	400	100,0%

### diagPca \* ch6983 Crosstabulation

#### Count

		ch69		
		normal	Total	
diagPca	no	26	124	150
	yes	17	133	150
	Total	43	257	300

### **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	2,199 <sup>a</sup>	1	,138		
Continuity Correction b	1,737	1	,187		
Likelihood Ratio	2,213	1	,137		
Fisher's Exact Test				,187	,094
Linear-by-Linear Association	2,192	1	,139		
N of Valid Cases	300				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 21,50.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower Uppe		
Odds Ratio for diagPca (no / yes)	1,640	,849	3,169	
For cohort ch6983 = normal	1,529	,867	2,699	
For cohort ch6983 = patology	,932	,850	1,023	
N of Valid Cases	300			

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

## **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	2,199	1	,138
Mantel-Haenszel	1,732	1	,188

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	1,640
		In(Estimate)	,495
		Std. Error of In(Estimate)	,336
		Asymp. Sig. (2-sided)	,141
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,849
Interval		Upper Bound	3,169
	In(Common Odds Ratio)	Lower Bound	-,164
		Upper Bound	1,153

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=cmDiagPca0Kont BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:12:42
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmDiagPca0Kont BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
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	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Valid Missing Total					
	N	Percent	N	Percent	N	Percent
cmDiagPca0Kont * ch6983	250	62,5%	150	37,5%	400	100,0%

## cmDiagPca0Kont \* ch6983 Crosstabulation

## Count

004.14				
		ch69		
		normal	patology	Total
cmDiagPca0Kont	no	26	124	150
	control	26	74	100
	Total	52	198	250

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	2,736 <sup>a</sup>	1	,098		
Continuity Correction b	2,235	1	,135		
Likelihood Ratio	2,696	1	,101		
Fisher's Exact Test				,113	,068
Linear-by-Linear Association	2,725	1	,099		
N of Valid Cases	250				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 20,80.

## **Risk Estimate**

		95% Confidence Interv	
	Value	Lower	Upper
Odds Ratio for cmDiagPca0Kont (no / control)	,597	,323	1,104
For cohort ch6983 = normal	,667	,412	1,079
For cohort ch6983 = patology	1,117	,974	1,282
N of Valid Cases	250		

# Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

b. Computed only for a 2x2 table

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	2,736	1	,098
Mantel-Haenszel	2,226	1	,136

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,597
		In(Estimate)	-,516
		Std. Error of In(Estimate)	,314
		Asymp. Sig. (2-sided)	,100
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,323
Interval		Upper Bound	1,104
	In(Common Odds Ratio)	Lower Bound	-1,131
		Upper Bound	,099

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

### CROSSTABS

/TABLES=cmDiagPca1Kont BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

## **Crosstabs**

	Output Created	22-lip-2012 12:12:42
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
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	Split File	<none></none>
	N of Rows in Working Data File	400

Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmDiagPca1Kont BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.016
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Valid Missing Total					tal
	N Percent N Percent			N	Percent	
cmDiagPca1Kont * ch6983	250	62,5%	150	37,5%	400	100,0%

## cmDiagPca1Kont \* ch6983 Crosstabulation

#### Count

Count					
		ch6983			
				Total	
cmDiagPca1Kont	yes	17	133	150	
	control	26	74	100	
	Total	43	207	250	

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	9,063 <sup>a</sup>	1	,003		
Continuity Correction b	8,062	1	,005		
Likelihood Ratio	8,882	1	,003		
Fisher's Exact Test				,003	,002
Linear-by-Linear Association	9,026	1	,003		
N of Valid Cases	250				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 17,20.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interva	
	Value	Lower	Upper
Odds Ratio for cmDiagPca1Kont (yes / control)	,364	,185	,714
For cohort ch6983 = normal	,436	,250	,760
For cohort ch6983 = patology	1,198	1,053	1,364
N of Valid Cases	250		

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	9,063	1	,003
Mantel-Haenszel	8,030	1	,005

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,364
		In(Estimate)	-1,011
		Std. Error of In(Estimate)	,344
		Asymp. Sig. (2-sided)	,003
Asymp. 95% Confidence Interval	Common Odds Ratio	Lower Bound	,185
	In(Common Odds Ratio)	Upper Bound	,714
		Lower Bound	-1,685
		Upper Bound	-,337

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=kontrol BY ch6983

/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

### **Crosstabs**

	Output Created	22-lip-2012 12:12:42
	Comments	.,,
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=kontrol BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
	Elapsed Time	0:00:00.146
	Dimensions Requested	2
	Cells Available	174762

 $[DataSet1] \ \, U:\ \, Data\ \, My \ \, Folders\ \, Science\ \, WorkCurrent\ \, \_rad\_b01\_x\_dsmbdmvf\ \, Lossyman \ \, Lossyman$ 

# **Case Processing Summary**

	Cases						
	Va	Valid Missing Total					
	N Percent N Percent N Percer					Percent	
kontrol * ch6983	400	TO THE PROPERTY OF THE PROPERT					

### kontrol \* ch6983 Crosstabulation

Count

		ch69		
		normal	patology	Total
kontrol	no control	43	257	300
	control	26	74	100
	Total	69	331	400

### **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	7,152 <sup>a</sup>	1	,007

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 17,25.

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>D</sup>	6,358	1	,012		
Likelihood Ratio	6,670	1	,010		
Fisher's Exact Test				,009	,007
Linear-by-Linear Association	7,134	1	,008		
N of Valid Cases	400				

### b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower Uppe		
Odds Ratio for kontrol (no control / control)	,476	,274	,826	
For cohort ch6983 = normal	,551	,358	,848	
For cohort ch6983 = patology	1,158	1,022	1,312	
N of Valid Cases	400			

### Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	7,152	1	,007
Mantel-Haenszel	6,342	1	,012

Under the conditional independence assumption,
Cochran's statistic is asymptotically distributed as a 1 df
chi-squared distribution, only if the number of strata is
fixed, while the Mantel-Haenszel statistic is always
asymptotically distributed as a 1 df chi-squared
distribution. Note that the continuity correction is removed
from the Mantel-Haenszel statistic when the sum of the
differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,476
		In(Estimate)	-,742
		Std. Error of In(Estimate)	,281
Asymp. 95% Confidence		Asymp. Sig. (2-sided)	,008
	Common Odds Ratio	Lower Bound	,274
Interval		Upper Bound	,826
	In(Common Odds Ratio)	Lower Bound	-1,293
		Upper Bound	-,191

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=cmTStadOnly12 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:12:43
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmTStadOnly12 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.166
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Valid Missing Total				tal	
	N Percent N Percent N Percent					Percent
cmTStadOnly12 * ch6983	100	25,0%	300	75,0%	400	100,0%

## cmTStadOnly12 \* ch6983 Crosstabulation

#### Count

		ch69	983	
		normal	Total	
cmTStadOnly12	T1	2	26	28
	T2	11 61		72
	Total	13 87		100

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	1,180 <sup>a</sup>	1	,277		
Continuity Correction b	,570	1	,450		
Likelihood Ratio	1,308	1	,253		
Fisher's Exact Test				,342	,231
Linear-by-Linear Association	1,168	1	,280		
N of Valid Cases	100				

a. 1 cells (25,0%) have expected count less than 5. The minimum expected count is 3,64.

## **Risk Estimate**

		95% Confidence Interval		
	Value	Lower Upper		
Odds Ratio for cmTStadOnly12 (T1 / T2)	,427	,088	2,061	
For cohort ch6983 = normal	,468	,111	1,977	
For cohort ch6983 = patology	1,096	,951	1,263	
N of Valid Cases	100			

# Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

b. Computed only for a 2x2 table

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	1,180	1	,277
Mantel-Haenszel	,564	1	,453

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,427
		In(Estimate)	-,852
		Std. Error of In(Estimate)	,804
		Asymp. Sig. (2-sided)	,289
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,088
Interval		Upper Bound	2,061
	In(Common Odds Ratio)	Lower Bound	-2,427
		Upper Bound	,723

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

### CROSSTABS

/TABLES=cmTStadOnly13 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

## **Crosstabs**

	Output Created	22-lip-2012 12:12:43
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400

Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmTStadOnly13 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.000
	Elapsed Time	0:00:00.180
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Valid Missing Total				tal	
	N Percent N Percent N				N	Percent
cmTStadOnly13 * ch6983	78	19,5%	322	80,5%	400	100,0%

## cmTStadOnly13 \* ch6983 Crosstabulation

#### Count

Count						
		ch6983				
		normal	patology	Total		
cmTStadOnly13	T1	2	26	28		
	T3,T4	4	46	50		
	Total	6	72	78		

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,019 <sup>a</sup>	1	,892		
Continuity Correction b	,000	1	1,000		
Likelihood Ratio	,019	1	,891		
Fisher's Exact Test				1,000	,632
Linear-by-Linear Association	,018	1	,892		
N of Valid Cases	78				

- a. 2 cells (50,0%) have expected count less than 5. The minimum expected count is 2,15.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for cmTStadOnly13 (T1 / T3, T4)	,885	,152	5,163	
For cohort ch6983 = normal	,893	,174	4,571	
For cohort ch6983 = patology	1,009	,885	1,151	
N of Valid Cases	78			

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,019	1	,892
Mantel-Haenszel	,093	1	,761

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,885
		In(Estimate)	-,123
		Std. Error of In(Estimate)	,900
		Asymp. Sig. (2-sided)	,892
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,152
Interval		Upper Bound	5,163
	In(Common Odds Ratio)	Lower Bound	-1,887
		Upper Bound	1,642

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

## CROSSTABS

/TABLES=cmTStadOnly23 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

## **Crosstabs**

	Output Created	22-lip-2012 12:12:43
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmTStadOnly23 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.000
	Elapsed Time	0:00:00.202
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# **Case Processing Summary**

	Cases					
	Valid Missing Total			tal		
	N	Percent	N	Percent	Ν	Percent
cmTStadOnly23 * ch6983	122	30,5%	278	69,5%	400	100,0%

## cmTStadOnly23 \* ch6983 Crosstabulation

Count

		ch6983		
		normal	Total	
cmTStadOnly23	T2	11	61	72
	T3,T4	4	46	50
	Total	15	107	122

### **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	1,449 <sup>a</sup>	1	,229		
Continuity Correction b	,853	1	,356		
Likelihood Ratio	1,518	1	,218		
Fisher's Exact Test				,273	,179
Linear-by-Linear Association	1,438	1	,231		
N of Valid Cases	122				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,15.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for cmTStadOnly23 (T2 / T3, T4)	2,074	,620	6,931	
For cohort ch6983 = normal	1,910	,645	5,657	
For cohort ch6983 = patology	,921	,811	1,046	
N of Valid Cases	122			

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	1,449	1	,229
Mantel-Haenszel	,846	1	,358

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

Estimate	2,074
In(Estimate)	,729
Std. Error of In(Estimate)	,616
Asymp. Sig. (2-sided)	,236

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

#### **Mantel-Haenszel Common Odds Ratio Estimate**

Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,620
Interval		Upper Bound	6,931
	In(Common Odds Ratio)	Lower Bound	-,477
		Upper Bound	1,936

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=cmPsaLT10vs10to20FonPCA1 BY ch6983

/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmPsaLT10vs10to20Fon PCA1 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

Resources	Processor Time	0:00:00.016
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	Cells Available	174762

 $\label{thm:cond} $$[DataSet1]$ U:\Personal Data\My Folders\Science\WorkCurrent\_rad\_b01_x_dsmbdmvf\rez\SPSS\Stat.sav$ 

## **Case Processing Summary**

	Cases					
	Va	lid	Miss	sing	To	tal
	N	Percent	Ν	Percent	Ν	Percent
cmPsaLT10vs10to20Fon PCA1 * ch6983	88	22,0%	312	78,0%	400	100,0%

#### cmPsaLT10vs10to20FonPCA1 \* ch6983 Crosstabulation

#### Count

			ch6983	
		normal	patology	Total
cmPsaLT10vs10to20Fon	<10	8	41	49
PCA1	10-20	2	37	39
	Total	10	78	88

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	2,704 <sup>a</sup>	1	,100		
Continuity Correction b	1,706	1	,191		
Likelihood Ratio	2,921	1	,087		
Fisher's Exact Test				,175	,094
Linear-by-Linear Association	2,673	1	,102		
N of Valid Cases	88				

- a. 1 cells (25,0%) have expected count less than 5. The minimum expected count is 4,43.
- b. Computed only for a 2x2 table

### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for cmPsaLT10vs10to20Fon PCA1 (<10 / 10-20)	3,610	,720	18,094	
For cohort ch6983 = normal	3,184	,717	14,145	
For cohort ch6983 = patology	,882	,764	1,018	
N of Valid Cases	88			

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

#### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	2,704	1	,100
Mantel-Haenszel	1,687	1	,194

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	3,610
		In(Estimate)	1,284
		Std. Error of In(Estimate)	,822
		Asymp. Sig. (2-sided)	,119
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,720
Interval		Upper Bound	18,094
	In(Common Odds Ratio)	Lower Bound	-,328
		Upper Bound	2,896

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

#### CROSSTABS

/TABLES=cmPsaLT10vsGT20FonPCA1 BY ch6983
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ RISK CMH(1)
/CELLS=COUNT
/COUNT ROUND CELL.

## **Crosstabs**

Output Created	22-lip-2012 12:12:44
Comments	

Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmPsaLT10vsGT20Fon PCA1 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
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	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Va	lid	Miss	sing	То	tal
	N	Percent	N	Percent	N	Percent
cmPsaLT10vsGT20Fon PCA1 * ch6983	111	27,8%	289	72,3%	400	100,0%

### cmPsaLT10vsGT20FonPCA1 \* ch6983 Crosstabulation

#### Count

Count				
		ch6983		
		normal	patology	Total
cmPsaLT10vsGT20Fon	<10	8	41	49
PCA1	>20	7	55	62
	Total	15	96	111

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	,594 <sup>a</sup>	1	,441

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,62.

### **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>D</sup>	,241	1	,623		
Likelihood Ratio	,590	1	,443		
Fisher's Exact Test				,578	,310
Linear-by-Linear Association	,589	1	,443		
N of Valid Cases	111				

### b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for cmPsaLT10vsGT20Fon PCA1 (<10 / >20)	1,533	,514	4,569	
For cohort ch6983 = normal	1,446	,563	3,712	
For cohort ch6983 = patology	,943	,810	1,098	
N of Valid Cases	111			

#### Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,594	1	,441
Mantel-Haenszel	,239	1	,625

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

Estimate	1,533
In(Estimate)	,427
Std. Error of In(Estimate)	,557
Asymp. Sig. (2-sided)	,443

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

#### Mantel-Haenszel Common Odds Ratio Estimate

Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,514
Interval		Upper Bound	4,569
	In(Common Odds Ratio)	Lower Bound	-,665
		Upper Bound	1,519

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=cmPsa10to20vsGT20FonPCA1 BY ch6983

/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

## **Crosstabs**

#### **Notes**

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	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmPsa10to20vsGT20Fon PCA1 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
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[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# **Case Processing Summary**

	Cases					
	Valid Missing Total				tal	
	N	Percent	N	Percent		
cmPsa10to20vsGT20Fon PCA1 * ch6983	101	25,3%	299	74,8%	400	100,0%

## cmPsa10to20vsGT20FonPCA1 \* ch6983 Crosstabulation

#### Count

			983	
		normal	patology	Total
cmPsa10to20vsGT20Fon	10-20	2	37	39
PCA1	>20	7	55	62
	Total	9	92	101

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	1,120 <sup>a</sup>	1	,290		
Continuity Correction b	,489	1	,484		
Likelihood Ratio	1,203	1	,273		
Fisher's Exact Test				,476	,247
Linear-by-Linear Association	1,109	1	,292		
N of Valid Cases	101				

a. 1 cells (25,0%) have expected count less than 5. The minimum expected count is 3,48.

#### **Risk Estimate**

		95% Confidence Interva		
	Value	Lower	Upper	
Odds Ratio for cmPsa10to20vsGT20Fon PCA1 (10-20 / >20)	,425	,084	2,159	
For cohort ch6983 = normal	,454	,099	2,076	
For cohort ch6983 = patology	1,069	,953	1,200	
N of Valid Cases	101			

# Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

b. Computed only for a 2x2 table

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	1,120	1	,290
Mantel-Haenszel	,485	1	,486

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,425
		In(Estimate)	-,856
		Std. Error of In(Estimate)	,829
		Asymp. Sig. (2-sided)	,302
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,084
Interval		Upper Bound	2,159
	In(Common Odds Ratio)	Lower Bound	-2,482
		Upper Bound	,769

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

## CROSSTABS

/TABLES=cmPsaLT20vsGT20onPCA1 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

## **Crosstabs**

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	Split File	<none></none>
	N of Rows in Working Data File	400

Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmPsaLT20vsGT20on PCA1 BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
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[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Valid Missing			Total		
	N	Percent	N	Percent	N	Percent
cmPsaLT20vsGT20on PCA1 * ch6983	150	37,5%	250	62,5%	400	100,0%

## cmPsaLT20vsGT20onPCA1 \* ch6983 Crosstabulation

### Count

		ch6983		
		normal	patology	Total
cmPsaLT20vsGT20on	,00	10	78	88
PCA1	<10	7	55	62
	Total	17	133	150

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,000 <sup>a</sup>	1	,989		
Continuity Correction b	,000	1	1,000		
Likelihood Ratio	,000	1	,989		
Fisher's Exact Test				1,000	,602
Linear-by-Linear Association	,000	1	,989		
N of Valid Cases	150				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 7,03.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interva	
	Value	Lower	Upper
Odds Ratio for cmPsaLT20vsGT20on PCA1 (,00 / <10)	1,007	,361	2,809
For cohort ch6983 = normal	1,006	,405	2,499
For cohort ch6983 = patology	,999	,890	1,122
N of Valid Cases	150		

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,000	1	,989
Mantel-Haenszel	,061	1	,805

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	1,007
		In(Estimate)	,007
		Std. Error of In(Estimate)	,523
		Asymp. Sig. (2-sided)	,989
Asymp. 95% Confidence Common Oc Interval	Common Odds Ratio	Lower Bound	,361
		Upper Bound	2,809
	In(Common Odds Ratio)	Lower Bound	-1,018
		Upper Bound	1,033

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

## CROSSTABS

/TABLES=cmGgLtvsGt7F BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

## **Crosstabs**

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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmGgLtvsGt7F BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
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	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# **Case Processing Summary**

	Cases					
	Valid Missing Total				tal	
	N Percent N Percent N Per					Percent
cmGgLtvsGt7F * ch6983	93	23,3%	307	76,8%	400	100,0%

# cmGgLtvsGt7F \* ch6983 Crosstabulation

#### Count

000					
		ch69			
		normal	Total		
cmGgLtvsGt7F	<7	7	64	71	
	>7	4 18		22	
	Total	11	82	93	

### **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	1,116 <sup>a</sup>	1	,291

a. 1 cells (25,0%) have expected count less than 5. The minimum expected count is 2,60.

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>D</sup>	,460	1	,498		
Likelihood Ratio	1,025	1	,311		
Fisher's Exact Test				,282	,240
Linear-by-Linear Association	1,104	1	,293		
N of Valid Cases	93				

### b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval  Lower Upper		
	Value			
Odds Ratio for cmGgLtvsGt7F (<7 / >7)	,492	,130	1,871	
For cohort ch6983 = normal	,542	,175	1,681	
For cohort ch6983 = patology	1,102	,892	1,361	
N of Valid Cases	93			

### Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	1,116	1	,291
Mantel-Haenszel	,455	1	,500

Under the conditional independence assumption,
Cochran's statistic is asymptotically distributed as a 1 df
chi-squared distribution, only if the number of strata is
fixed, while the Mantel-Haenszel statistic is always
asymptotically distributed as a 1 df chi-squared
distribution. Note that the continuity correction is removed
from the Mantel-Haenszel statistic when the sum of the
differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,492
		In(Estimate)	-,709
		Std. Error of In(Estimate)	,681
		Asymp. Sig. (2-sided)	,298
	Common Odds Ratio	Lower Bound	,130
Interval		Upper Bound	1,871
	In(Common Odds Ratio)	Lower Bound	-2,044
		Upper Bound	,626

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=cmGgLt7vsEq7F BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmGgLt7vsEq7F BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.017
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Valid Missing Total					tal
	N	Percent	N	Percent	N	Percent
cmGgLt7vsEq7F * ch6983	128	32,0%	272	68,0%	400	100,0%

## cmGgLt7vsEq7F \* ch6983 Crosstabulation

## Count

		ch69	ch6983		
		normal	patology	Total	
cmGgLt7vsEq7F	<7	7	64	71	
	=7	6 51		57	
	Total	13	13 115		

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,015 <sup>a</sup>	1	,901		
Continuity Correction b	,000	1	1,000		
Likelihood Ratio	,015	1	,901		
Fisher's Exact Test				1,000	,564
Linear-by-Linear Association	,015	1	,902		
N of Valid Cases	128				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 5,79.

## **Risk Estimate**

		95% Confidence Interva		
	Value	Lower	Upper	
Odds Ratio for cmGgLt7vsEq7F (<7 / =7)	,930	,294	2,939	
For cohort ch6983 = normal	,937	,333	2,632	
For cohort ch6983 = patology	1,007	,896	1,133	
N of Valid Cases	128			

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

b. Computed only for a 2x2 table

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,015	1	,901
Mantel-Haenszel	,029	1	,865

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,930
		In(Estimate)	-,073
		Std. Error of In(Estimate)	,587
		Asymp. Sig. (2-sided)	,901
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,294
Interval		Upper Bound	2,939
	In(Common Odds Ratio)	Lower Bound	-1,224
		Upper Bound	1,078

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

### CROSSTABS

/TABLES=cmGgEq7vsGt7F BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

## **Crosstabs**

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Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
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	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400

Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=cmGgEq7vsGt7F BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.015
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

## **Case Processing Summary**

	Cases					
	Valid Missing Total					tal
	N	Percent	N	Percent		
cmGgEq7vsGt7F * ch6983	79	19,8%	321	80,3%	400	100,0%

## cmGgEq7vsGt7F \* ch6983 Crosstabulation

#### Count

Count						
		ch69				
		normal	Total			
cmGgEq7vsGt7F	=7	6	51	57		
	>7	4	18	22		
	Total	10	69	79		

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,841 <sup>a</sup>	1	,359		
Continuity Correction b	,291	1	,589		
Likelihood Ratio	,792	1	,374		
Fisher's Exact Test				,452	,285
Linear-by-Linear Association	,831	1	,362		
N of Valid Cases	79				

- a. 1 cells (25,0%) have expected count less than 5. The minimum expected count is 2,78.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower Upper		
Odds Ratio for cmGgEq7vsGt7F (=7 / >7)	,529	,134	2,093	
For cohort ch6983 = normal	,579	,180	1,857	
For cohort ch6983 = patology	1,094	,881	1,357	
N of Valid Cases	79			

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,841	1	,359
Mantel-Haenszel	,288	1	,592

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	,529
		In(Estimate)	-,636
		Std. Error of In(Estimate)	,701
		Asymp. Sig. (2-sided)	,364
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,134
Interval		Upper Bound	2,093
	In(Common Odds Ratio)	Lower Bound	-2,011
		Upper Bound	,739

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=mMeta BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

## **Crosstabs**

	Output Created	22-lip-2012 12:12:47
	Comments	
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	Active Dataset	DataSet1
	Filter	<none></none>
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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mMeta BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.020
	Dimensions Requested	2
	Cells Available	174762

 $[DataSet1] \ \, U:\ \, Data\ \, My \ \, Folders\ \, Science\ \, WorkCurrent\ \, \_rad\_b01\_x\_dsmbdmvf\ \, Lossyman \ \, Lossyman$ 

# **Case Processing Summary**

	Cases					
	Valid Missing Total				tal	
	N Percent N Percent N P				Percent	
mMeta * ch6983	150	37,5%	250	62,5%	400	100,0%

### mMeta \* ch6983 Crosstabulation

#### Count

Count					
		ch6983			
		normal	Total		
mMeta	no	12	83	95	
	yes	5	50	55	
	Total	17	133	150	

### **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	,435 <sup>a</sup>	1	,510

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,23.

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>D</sup>	,154	1	,695		
Likelihood Ratio	,447	1	,504		
Fisher's Exact Test				,601	,354
Linear-by-Linear Association	,432	1	,511		
N of Valid Cases	150				

### b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for mMeta (no / yes)	1,446	,481	4,346	
For cohort ch6983 = normal	1,389	,517	3,736	
For cohort ch6983 = patology	,961	,858,	1,076	
N of Valid Cases	150			

### Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,435	1	,510
Mantel-Haenszel	,153	1	,696

Under the conditional independence assumption,
Cochran's statistic is asymptotically distributed as a 1 df
chi-squared distribution, only if the number of strata is
fixed, while the Mantel-Haenszel statistic is always
asymptotically distributed as a 1 df chi-squared
distribution. Note that the continuity correction is removed
from the Mantel-Haenszel statistic when the sum of the
differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	1,446
		In(Estimate)	,369
		Std. Error of In(Estimate)	,562
		Asymp. Sig. (2-sided)	,512
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,481
Interval		Upper Bound	4,346
	In(Common Odds Ratio)	Lower Bound	-,732
		Upper Bound	1,469

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=mRiskEAU BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:12:47
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mRiskEAU BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
	Elapsed Time	0:00:00.011
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

#### Warnings

The Tests for Homogeneity of the Odds Ratio table and the Mantel-Haenszel Common Odds Ratio Estimate table are not computed for mRiskEAU \* ch6983, because either (1) the group variable does not have exactly two distinct non-missing values or/and (2) the response variable does not have exactly two distinct non-missing values.

### **Case Processing Summary**

		Cases					
	Valid		Missing		Total		
	N Percent		N	Percent	Ν	Percent	
mRiskEAU * ch6983	150	37,5%	250	62,5%	400	100,0%	

### mRiskEAU \* ch6983 Crosstabulation

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Odunt					
		ch6	983		
		normal	patology	Total	
mRiskEAU	low	2	12	14	
	medium	7	48	55	
	high	8	73	81	
	Total	17	133	150	

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	,399 <sup>a</sup>	2	,819
Likelihood Ratio	,394	2	,821
Linear-by-Linear Association	,386	1	,535
N of Valid Cases	150		

a. 1 cells (16,7%) have expected count less than 5. The minimum expected count is 1,59.

#### **Risk Estimate**

	Value
Odds Ratio for mRiskEAU (low / medium)	а

a. Risk Estimate statistics cannot be computed. They are only computed for a 2\*2 table without empty cells.

## CROSSTABS

/TABLES=mRiskEAULowMedium BY ch6983

/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

## Notes

	Output Created	22-lip-2012 12:12:47
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mRiskEAULowMedium BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.016
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# **Case Processing Summary**

		Cases					
	Valid		Missing		Total		
	N	Percent	Ν	Percent	N	Percent	
mRiskEAULowMedium * ch6983	69	17,3%	331	82,8%	400	100,0%	

# mRiskEAULowMedium \* ch6983 Crosstabulation

## Count

Count				
1		ch69		
		normal	patology	Total
mRiskEAULowMedium	low	2	12	14
	medium	7	48	55
	Total	9	60	69

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,024 <sup>a</sup>	1	,877		
Continuity Correction b	,000	1	1,000		
Likelihood Ratio	,023	1	,878		
Fisher's Exact Test				1,000	,587
Linear-by-Linear Association	,024	1	,878		
N of Valid Cases	69				

- a. 1 cells (25,0%) have expected count less than 5. The minimum expected count is 1,83.
- b. Computed only for a 2x2 table

### **Risk Estimate**

		95% Confidence Interva	
	Value	Lower	Upper
Odds Ratio for mRiskEAULowMedium (low / medium)	1,143	,210	6,219
For cohort ch6983 = normal	1,122	,261	4,823
For cohort ch6983 = patology	,982	,775	1,244
N of Valid Cases	69		

# Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

## **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,024	1	,877
Mantel-Haenszel	,083	1	,774

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

Estimate	1,143
In(Estimate)	,134
Std. Error of In(Estimate)	,864
Asymp. Sig. (2-sided)	,877

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

### **Mantel-Haenszel Common Odds Ratio Estimate**

Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,210
Interval		Upper Bound	6,219
	In(Common Odds Ratio)	Lower Bound	-1,560
		Upper Bound	1,828

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS

/TABLES=mRiskEAULowHigh BY ch6983 /FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

	0	00 lin 0040 40:40:47
	Output Created	22-lip-2012 12:12:47
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Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mRiskEAULowHigh BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

Resources	Processor Time	0:00:00.016
	Elapsed Time	0:00:00.014
	Dimensions Requested	2
	Cells Available	174762

 $\label{thm:cond} $$[DataSet1]$ U:\Personal Data\My Folders\Science\WorkCurrent\_rad\_b01_x_dsmbdmvf\rez\SPSS\Stat.sav$ 

# **Case Processing Summary**

	Cases					
	Valid		Miss	sing	То	tal
	N	Percent	Ν	Percent	N	Percent
mRiskEAULowHigh * ch6983	95	23,8%	305	76,3%	400	100,0%

# mRiskEAULowHigh \* ch6983 Crosstabulation

### Count

Count				
		ch6983		
		normal	patology	Total
mRiskEAULowHigh	low	2	12	14
	high	8	73	81
	Total	10	85	95

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,246 <sup>a</sup>	1	,620		
Continuity Correction b	,001	1	,980		
Likelihood Ratio	,228	1	,633		
Fisher's Exact Test				,639	,453
Linear-by-Linear Association	,244	1	,621		
N of Valid Cases	95				

- a. 1 cells (25,0%) have expected count less than 5. The minimum expected count is 1,47.
- b. Computed only for a 2x2 table

### **Risk Estimate**

		95% Confidence Interva	
	Value	Lower	Upper
Odds Ratio for mRiskEAULowHigh (low / high)	1,521	,288	8,042
For cohort ch6983 = normal	1,446	,342	6,117
For cohort ch6983 = patology	,951	,759	1,192
N of Valid Cases	95		

# Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,246	1	,620
Mantel-Haenszel	,001	1	,980

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	1,521
		In(Estimate)	,419
		Std. Error of In(Estimate)	,850
		Asymp. Sig. (2-sided)	,622
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,288
Interval		Upper Bound	8,042
	In(Common Odds Ratio)	Lower Bound	-1,246
		Upper Bound	2,085

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

## CROSSTABS

/TABLES=mRiskEAUMediumHigh BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

Output Created	22-lip-2012 12:12:48
Comments	

Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
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	Filter	<none></none>
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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mRiskEAUMediumHigh BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
	Elapsed Time	0:00:00.014
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# **Case Processing Summary**

	Cases						
	Va	Valid Missing To				Total	
	N	Percent	N	Percent	N	Percent	
mRiskEAUMediumHigh * ch6983	136	34,0%	264	66,0%	400	100,0%	

# mRiskEAUMediumHigh \* ch6983 Crosstabulation

## Count

		ch6983		
		normal		
mRiskEAUMediumHigh	medium	7	48	55
	high	8	73	81
	Total	15	121	136

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	,271 <sup>a</sup>	1	,602

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,07.

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>0</sup>	,059	1	,809		
Likelihood Ratio	,268	1	,605		
Fisher's Exact Test				,592	,400
Linear-by-Linear Association	,269	1	,604		
N of Valid Cases	136				

## b. Computed only for a 2x2 table

### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for mRiskEAUMediumHigh (medium / high)	1,331	,453	3,910	
For cohort ch6983 = normal	1,289	,496	3,348	
For cohort ch6983 = patology	,968	,855	1,096	
N of Valid Cases	136			

## Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,271	1	,602
Mantel-Haenszel	,058	1	,809

Under the conditional independence assumption,
Cochran's statistic is asymptotically distributed as a 1 df
chi-squared distribution, only if the number of strata is
fixed, while the Mantel-Haenszel statistic is always
asymptotically distributed as a 1 df chi-squared
distribution. Note that the continuity correction is removed
from the Mantel-Haenszel statistic when the sum of the
differences between the observed and the expected is 0.

#### Mantel-Haenszel Common Odds Ratio Estimate

Estimate	1,331
In(Estimate)	,286
Std. Error of In(Estimate)	,550
Asymp. Sig. (2-sided)	,603

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

### Mantel-Haenszel Common Odds Ratio Estimate

Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,453
Interval		Upper Bound	3,910
	In(Common Odds Ratio)	Lower Bound	-,792
		Upper Bound	1,364

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

CROSSTABS
/TABLES=mRiskMed BY ch6983
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ RISK CMH(1)
/CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

### **Notes**

	Output Created	22-lip-2012 12:12:48
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
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	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mRiskMed BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.000
	Elapsed Time	0:00:00.013
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# **Case Processing Summary**

	Cases					
	Valid Missing Total					tal
	N	Percent	N Percent		N	Percent
mRiskMed * ch6983	150	37,5%	250	62,5%	400	100,0%

### mRiskMed \* ch6983 Crosstabulation

## Count

		ch6983		
		normal	Total	
mRiskMed	low	8	47	55
	high	9 86		95
	Total	17 133		150

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,892 <sup>a</sup>	1	,345		
Continuity Correction b	,458	1	,498		
Likelihood Ratio	,868	1	,352		
Fisher's Exact Test				,424	,246
Linear-by-Linear Association	,886	1	,347		
N of Valid Cases	150				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,23.

# **Risk Estimate**

		95% Confidence Interva	
	Value	Lower	Upper
Odds Ratio for mRiskMed (low / high)	1,626	,589	4,495
For cohort ch6983 = normal	1,535	,629	3,749
For cohort ch6983 = patology	,944	,831	1,072
N of Valid Cases	150		

# Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

b. Computed only for a 2x2 table

## **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,892	1	,345
Mantel-Haenszel	,455	1	,500

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	1,626
		In(Estimate)	,486
		Std. Error of In(Estimate)	,519
		Asymp. Sig. (2-sided)	,348
Asymp. 95% Confidence Interval	Common Odds Ratio	Lower Bound	,589
		Upper Bound	4,495
	In(Common Odds Ratio)	Lower Bound	-,530
		Upper Bound	1,503

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

### CROSSTABS

/TABLES=mRiskMedLowMedium BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:12:49
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400

Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mRiskMedLowMedium BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.016
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# Warnings

No measures of association are computed for the crosstabulation of mRiskMedLowMedium \* ch6983. At least one variable in each 2-way table upon which measures of association are computed is a constant.

## **Case Processing Summary**

	Cases					
	Va	lid	Missing		Total	
	N	Percent	N	Percent	N	Percent
mRiskMedLowMedium * ch6983	55	13,8%	345	86,3%	400	100,0%

### mRiskMedLowMedium \* ch6983 Crosstabulation

### Count

Count					
		ch6983			
		normal	patology	Total	
mRiskMedLowMedium	low	8	47	55	
	Total	8	47	55	

# **Chi-Square Tests**

	Value
Pearson Chi-Square	а
N of Valid Cases	55

a. No statistics are computed because mRiskMedLowMedium is a constant.

### **Risk Estimate**

	Value
Odds Ratio for mRiskMedLowMedium (low / .)	а

a. No statistics are computed because mRiskMedLowMedium is a constant.

CROSSTABS

/TABLES=mRiskMedLowHigh BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

### **Notes**

	Output Created	22-lip-2012 12:12:49
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400
Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mRiskMedLowHigh BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.031
	Elapsed Time	0:00:00.130
	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# **Case Processing Summary**

	Cases					
	Valid M			Missing		tal
	N	Percent	N	Percent	N	Percent
mRiskMedLowHigh * ch6983	150	37,5%	250	62,5%	400	100,0%

# mRiskMedLowHigh \* ch6983 Crosstabulation

## Count

Count		ch6983		
		normal	patology	Total
mRiskMedLowHigh	low	8	47	55
	high	9	86	95
	Total	17	133	150

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,892 <sup>a</sup>	1	,345		
Continuity Correction b	,458	1	,498		
Likelihood Ratio	,868	1	,352		
Fisher's Exact Test				,424	,246
Linear-by-Linear Association	,886,	1	,347		
N of Valid Cases	150				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,23.

### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for mRiskMedLowHigh (low / high)	1,626	,589	4,495	
For cohort ch6983 = normal	1,535	,629	3,749	
For cohort ch6983 = patology	,944	,831	1,072	
N of Valid Cases	150			

# Tests of Homogeneity of the Odds Ratio

	Chi-Squared	df	Asymp. Sig. (2-sided)
Breslow-Day	,000	0	
Tarone's	,000	0	

b. Computed only for a 2x2 table

### **Tests of Conditional Independence**

	Chi-Squared	df	Asymp. Sig. (2-sided)
Cochran's	,892	1	,345
Mantel-Haenszel	,455	1	,500

Under the conditional independence assumption, Cochran's statistic is asymptotically distributed as a 1 df chi-squared distribution, only if the number of strata is fixed, while the Mantel-Haenszel statistic is always asymptotically distributed as a 1 df chi-squared distribution. Note that the continuity correction is removed from the Mantel-Haenszel statistic when the sum of the differences between the observed and the expected is 0.

### Mantel-Haenszel Common Odds Ratio Estimate

		Estimate	1,626
		In(Estimate)	,486
		Std. Error of In(Estimate)	,519
		Asymp. Sig. (2-sided)	,348
	Common Odds Ratio	Lower Bound	,589
Interval		Upper Bound	4,495
	In(Common Odds Ratio)	Lower Bound	-,530
		Upper Bound	1,503

The Mantel-Haenszel common odds ratio estimate is asymptotically normally distributed under the common odds ratio of 1,000 assumption. So is the natural log of the estimate.

### CROSSTABS

/TABLES=mRiskMedMediumHigh BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:12:49
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
	Active Dataset	DataSet1
	Filter	<none></none>
	Weight	<none></none>
	Split File	<none></none>
	N of Rows in Working Data File	400

Missing Value Handling	Definition of Missing	User-defined missing values are treated as missing.
	Cases Used	Statistics for each table are based on all the cases with valid data in the specified range(s) for all variables in each table.
	Syntax	CROSSTABS /TABLES=mRiskMedMediumHigh BY ch6983 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
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	Dimensions Requested	2
	Cells Available	174762

[DataSet1] U:\Personal Data\My Folders\Science\WorkCurrent\\_rad\_b01\_x\_dsmb dmvf\rez\SPSS\Stat.sav

# Warnings

No measures of association are computed for the crosstabulation of mRiskMedMediumHigh \* ch6983. At least one variable in each 2-way table upon which measures of association are computed is a constant.

## **Case Processing Summary**

	Cases					
	Valid Missing Total					tal
	N Percent N Percent N P					Percent
mRiskMedMediumHigh * ch6983	95	23,8%	305	76,3%	400	100,0%

## mRiskMedMediumHigh \* ch6983 Crosstabulation

### Count

Oddrit				
		ch6983		
		normal	patology	Total
mRiskMedMediumHigh	high	9	86	95
	Total	9	86	95

# **Chi-Square Tests**

	Value
Pearson Chi-Square	a
N of Valid Cases	95

a. No statistics are computed because mRiskMedMediumHigh is a constant.

# **Risk Estimate**

	Value
Odds Ratio for mRiskMedMediumHigh (high / .)	а

a. No statistics are computed because mRiskMedMediumHigh is a constant.