CROSSTABS
/TABLES=diagPca BY ch3760
/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

#### Notes

	0	00 1:- 0040 40:40:47
	Output Created	22-lip-2012 12:18:47
	Comments	
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	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=diagPca BY ch3760 /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					
	Valid Missing Total				tal	
	N	N Percent		Percent	Ν	Percent
diagPca * ch3760	300	75,0%	100	25,0%	400	100,0%

## diagPca \* ch3760 Crosstabulation

Count

Count					
		ch3			
		normal	Total		
diagPca	no	65	85	150	
	yes	70	80	150	
	Total	135	165	300	

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,337 <sup>a</sup>	1	,562		
Continuity Correction b	,215	1	,642		
Likelihood Ratio	,337	1	,562		
Fisher's Exact Test				,643	,321
Linear-by-Linear Association	,336	1	,562		
N of Valid Cases	300				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 67,50.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for diagPca (no / yes)	,874	,554	1,378	
For cohort ch3760 = normal	,929	,723	1,193	
For cohort ch3760 = patology	1,063	,866	1,304	
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	,337	1	,562
Mantel-Haenszel	,215	1	,643

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	,874
		In(Estimate)	-,135
		Std. Error of In(Estimate)	,232
		Asymp. Sig. (2-sided)	,562
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,554
Interval		Upper Bound	1,378
	In(Common Odds Ratio)	Lower Bound	-,590
		Upper Bound	,320

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:18:47
	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=cmDiagPca0Kont BY ch3760 /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					
	Valid Missing Total					tal
	N	Percent	N	Percent	N	Percent
cmDiagPca0Kont * ch3760	250	62,5%	150	37,5%	400	100,0%

# cmDiagPca0Kont \* ch3760 Crosstabulation

# Count

000					
		ch3			
		normal	patology	Total	
cmDiagPca0Kont	no	65	85	150	
	control	51	49	100	
	Total	116	134	250	

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	1,418 <sup>a</sup>	1	,234		
Continuity Correction b	1,127	1	,289		
Likelihood Ratio	1,418	1	,234		
Fisher's Exact Test				,246	,144
Linear-by-Linear Association	1,412	1	,235		
N of Valid Cases	250				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 46,40.

# **Risk Estimate**

		95% Confide	nce Interval
	Value	Lower	Upper
Odds Ratio for cmDiagPca0Kont (no / control)	,735	,442	1,221
For cohort ch3760 = normal	,850	,652	1,108
For cohort ch3760 = patology	1,156	,906	1,476
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	1,418	1	,234
Mantel-Haenszel	1,122	1	,289

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	,735
		In(Estimate)	-,308
		Std. Error of In(Estimate)	,259
		Asymp. Sig. (2-sided)	,234
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,442
Interval		Upper Bound	1,221
	In(Common Odds Ratio)	Lower Bound	-,816
		Upper Bound	,200

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=cmDiagPcalKont BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:18:48
	Comments	
Input	Data	U:\Personal Data\My Folders\Science\WorkCurrent\_rad_ b01_x_dsmbdmvf\rez\SPSS\Stat.sav
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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=cmDiagPca1Kont BY ch3760 /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	Ν	Percent	N	Percent
cmDiagPca1Kont * ch3760	250	62,5%	150	37,5%	400	100,0%

# cmDiagPca1Kont \* ch3760 Crosstabulation

#### Count

Count					
		ch3760			
		normal patology		Total	
cmDiagPca1Kont	yes	70	80	150	
	control	51	49	100	
	Total	121	129	250	

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,451 <sup>a</sup>	1	,502		
Continuity Correction b	,294	1	,587		
Likelihood Ratio	,451	1	,502		
Fisher's Exact Test				,521	,294
Linear-by-Linear Association	,449	1	,503		
N of Valid Cases	250				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 48,40.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interv	
	Value	Lower	Upper
Odds Ratio for cmDiagPca1Kont (yes / control)	,841	,507	1,395
For cohort ch3760 = normal	,915	,707	1,183
For cohort ch3760 = patology	1,088	,848	1,397
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	,451	1	,502
Mantel-Haenszel	,293	1	,588

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	,841
		In(Estimate)	-,174
		Std. Error of In(Estimate)	,258
		Asymp. Sig. (2-sided)	,502
Asymp. 95% Confidence Interval	Common Odds Ratio In(Common Odds Ratio)	Lower Bound	,507
		Upper Bound	1,395
		Lower Bound	-,680
		Upper Bound	,333

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 ch3760

/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

## **Crosstabs**

	Output Created	22-lip-2012 12:18:48
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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=kontrol BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.047
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	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 * ch3760	400						

## kontrol \* ch3760 Crosstabulation

# Count

		ch3				
			normal patology			
kontrol	no control	135	165	300		
	control	51	49	100		
	Total	186	214	400		

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	1,085 <sup>a</sup>	1	,298

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 46,50.

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>0</sup>	,858	1	,354		
Likelihood Ratio	1,083	1	,298		
Fisher's Exact Test				,301	,177
Linear-by-Linear Association	1,083	1	,298		
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)	,786	,500	1,237	
For cohort ch3760 = normal	,882	,702	1,110	
For cohort ch3760 = patology	1,122	,897	1,405	
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	1,085	1	,298
Mantel-Haenszel	,855	1	,355

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	,786
		In(Estimate)	-,241
		Std. Error of In(Estimate)	,231
		Asymp. Sig. (2-sided)	,298
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,500
Interval		Upper Bound	1,237
	In(Common Odds Ratio)	Lower Bound	-,694
		Upper Bound	,213

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:18:48
	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 ch3760 /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					
	Va	Valid Missing Total				tal
	N Percent N Percent N Percei					Percent
cmTStadOnly12 * ch3760	100	25,0%	300	75,0%	400	100,0%

# cmTStadOnly12 \* ch3760 Crosstabulation

#### Count

000					
		ch3			
		normal	patology	Total	
cmTStadOnly12	T1	13	15	28	
	T2	37 35		72	
	Total	50	50	100	

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,198 <sup>a</sup>	1	,656		
Continuity Correction b	,050	1	,824		
Likelihood Ratio	,199	1	,656		
Fisher's Exact Test				,824	,412
Linear-by-Linear Association	,196	1	,658		
N of Valid Cases	100				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 14,00.

## **Risk Estimate**

		95% Confidence Interval		
	Value	Lower Upper		
Odds Ratio for cmTStadOnly12 (T1 / T2)	,820	,342	1,966	
For cohort ch3760 = normal	,903	,572	1,427	
For cohort ch3760 = patology	1,102	,725	1,675	
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	,198	1	,656
Mantel-Haenszel	,049	1	,825

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	,820
		In(Estimate)	-,199
		Std. Error of In(Estimate)	,446
		Asymp. Sig. (2-sided)	,656
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,342
Interval		Upper Bound	1,966
	In(Common Odds Ratio)	Lower Bound	-1,073
		Upper Bound	,676

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:18:48
	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=cmTStadOnly13 BY ch3760 /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	Percent	
cmTStadOnly13 * ch3760	78						

# cmTStadOnly13 \* ch3760 Crosstabulation

#### Count

Count						
		ch3760				
		normal	patology	Total		
cmTStadOnly13	T1	13	15	28		
	T3,T4	20	30	50		
	Total	33	45	78		

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,304 <sup>a</sup>	1	,581		
Continuity Correction b	,098	1	,755		
Likelihood Ratio	,303	1	,582		
Fisher's Exact Test				,637	,376
Linear-by-Linear Association	,300	1	,584		
N of Valid Cases	78				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 11,85.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for cmTStadOnly13 (T1 / T3, T4)	1,300	,511	3,307	
For cohort ch3760 = normal	1,161	,688,	1,958	
For cohort ch3760 = patology	,893	,591	1,349	
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	,304	1	,581
Mantel-Haenszel	,096	1	,756

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	1,300
		In(Estimate)	,262
		Std. Error of In(Estimate)	,476
		Asymp. Sig. (2-sided)	,582
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,511
Interval		Upper Bound	3,307
	In(Common Odds Ratio)	Lower Bound	-,671
		Upper Bound	1,196

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 ch3760
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ RISK CMH(1)
/CELLS=COUNT
/COUNT ROUND CELL.

## **Crosstabs**

	Output Created	22-lip-2012 12:18:49
	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=cmTStadOnly23 BY ch3760 /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					
	Va	Valid Missir			To	tal
	N Percent N Percent			Ν	Percent	
cmTStadOnly23 * ch3760	122	30,5%	278	69,5%	400	100,0%

# cmTStadOnly23 \* ch3760 Crosstabulation

Count

Count		ch3760		
		normal	patology	Total
cmTStadOnly23	T2	37	35	72
	T3,T4	20	30	50
	Total	57	65	122

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	1,538 <sup>a</sup>	1	,215		
Continuity Correction b	1,114	1	,291		
Likelihood Ratio	1,544	1	,214		
Fisher's Exact Test				,269	,146
Linear-by-Linear Association	1,525	1	,217		
N of Valid Cases	122				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 23,36.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for cmTStadOnly23 (T2 / T3, T4)	1,586	,764	3,292	
For cohort ch3760 = normal	1,285	,855	1,930	
For cohort ch3760 = patology	,810	,584	1,125	
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	1,538	1	,215
Mantel-Haenszel	1,105	1	,293

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	1,586
In(Estimate)	,461
Std. Error of In(Estimate)	,373
Asymp. Sig. (2-sided)	,216

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	,764
Interval		Upper Bound	3,292
	In(Common Odds Ratio)	Lower Bound	-,270
		Upper Bound	1,192

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 ch3760

/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

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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=cmPsaLT10vs10to20Fon PCA1 BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

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 $\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	То	tal
	N	Percent	N	Percent	N	Percent
cmPsaLT10vs10to20Fon PCA1 * ch3760	88	22,0%	312	78,0%	400	100,0%

## cmPsaLT10vs10to20FonPCA1 \* ch3760 Crosstabulation

#### Count

			ch3760		
		normal	patology	Total	
cmPsaLT10vs10to20Fon	<10	23	26	49	
PCA1	10-20	19	20	39	
	Total	42	46	88	

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,028 <sup>a</sup>	1	,868		
Continuity Correction b	,000	1	1,000		
Likelihood Ratio	,028	1	,868		
Fisher's Exact Test				1,000	,519
Linear-by-Linear Association	,027	1	,869		
N of Valid Cases	88				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 18,61.
- b. Computed only for a 2x2 table

### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower	Upper	
Odds Ratio for cmPsaLT10vs10to20Fon PCA1 (<10 / 10-20)	,931	,401	2,161	
For cohort ch3760 = normal	,963	,621	1,494	
For cohort ch3760 = patology	1,035	,691	1,549	
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	,028	1	,868
Mantel-Haenszel	,002	1	,961

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	,931
		In(Estimate)	-,071
		Std. Error of In(Estimate)	,430
		Asymp. Sig. (2-sided)	,868
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,401
Interval		Upper Bound	2,161
	In(Common Odds Ratio)	Lower Bound	-,913
		Upper Bound	,771

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	_
Output Created	22-lip-2012 12:18:49
Comments	

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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=cmPsaLT10vsGT20Fon PCA1 BY ch3760 /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					
	Va	Valid Missing Total				
	N	Percent	Ν	Percent	N	Percent
cmPsaLT10vsGT20Fon PCA1 * ch3760	111	27,8%	289	72,3%	400	100,0%

### cmPsaLT10vsGT20FonPCA1 \* ch3760 Crosstabulation

#### Count

Count				
		ch3760		
		normal	patology	Total
cmPsaLT10vsGT20Fon	<10	23	26	49
PCA1	>20	28	34	62
	Total	51	60	111

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	,035 <sup>a</sup>	1	,852

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 22,51.

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>D</sup>	,000	1	1,000		
Likelihood Ratio	,035	1	,852		
Fisher's Exact Test				1,000	,502
Linear-by-Linear Association	,035	1	,853		
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)	1,074	,507	2,278	
For cohort ch3760 = normal	1,039	,693	1,558	
For cohort ch3760 = patology	,968	,684	1,369	
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	,035	1	,852
Mantel-Haenszel	,000	1	,996

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	1,074
In(Estimate)	,072
Std. Error of In(Estimate)	,383
Asymp. Sig. (2-sided)	,852

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	,507
Interval		Upper Bound	2,278
	In(Common Odds Ratio)	Lower Bound	-,680
		Upper Bound	,823

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 ch3760 /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 ch3760 /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	N	Percent
cmPsa10to20vsGT20Fon PCA1 * ch3760	101	25,3%	299	74,8%	400	100,0%

# cmPsa10to20vsGT20FonPCA1 \* ch3760 Crosstabulation

## Count

		ch3760		
		normal	patology	Total
cmPsa10to20vsGT20Fon	10-20	19	20	39
PCA1	>20	28	34	62
	Total	47	54	101

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,122 <sup>a</sup>	1	,727		
Continuity Correction b	,021	1	,885		
Likelihood Ratio	,122	1	,727		
Fisher's Exact Test				,838	,442
Linear-by-Linear Association	,121	1	,728		
N of Valid Cases	101				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 18,15.

#### **Risk Estimate**

		95% Confidence Interv	
	Value	Lower	Upper
Odds Ratio for cmPsa10to20vsGT20Fon PCA1 (10-20 / >20)	1,154	,517	2,574
For cohort ch3760 = normal	1,079	,707	1,647
For cohort ch3760 = patology	,935	,639	1,368
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	,122	1	,727
Mantel-Haenszel	,021	1	,886

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	1,154
		In(Estimate)	,143
		Std. Error of In(Estimate)	,410
		Asymp. Sig. (2-sided)	,727
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,517
Interval		Upper Bound	2,574
	In(Common Odds Ratio)	Lower Bound	-,660
		Upper Bound	,946

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

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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=cmPsaLT20vsGT20on PCA1 BY ch3760 /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		
	N	N Percent N Percent		N	Percent	
cmPsaLT20vsGT20on PCA1 * ch3760	150	37,5%	250	62,5%	400	100,0%

# cmPsaLT20vsGT20onPCA1 \* ch3760 Crosstabulation

#### Count

Count					
			ch3760		
		normal	patology	Total	
cmPsaLT20vsGT20on	,00	42	46	88	
PCA1	<10	28	34	62	
	Total	70	80	150	

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,096 <sup>a</sup>	1	,756		
Continuity Correction b	,021	1	,885		
Likelihood Ratio	,096	1	,756		
Fisher's Exact Test				,868	,443
Linear-by-Linear Association	,096	1	,757		
N of Valid Cases	150				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 28,93.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interva	
	Value	Lower	Upper
Odds Ratio for cmPsaLT20vsGT20on PCA1 (,00 / <10)	1,109	,578	2,128
For cohort ch3760 = normal	1,057	,744	1,501
For cohort ch3760 = patology	,953	,705	1,289
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	,096	1	,756
Mantel-Haenszel	,021	1	,886

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	1,109
		In(Estimate)	,103
		Std. Error of In(Estimate)	,333
		Asymp. Sig. (2-sided)	,756
Asymp. 95% Confidence	Common Odds Ratio In(Common Odds Ratio)	Lower Bound	,578
Interval		Upper Bound	2,128
		Lower Bound	-,549
		Upper Bound	,755

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

## **Crosstabs**

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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=cmGgLtvsGt7F BY ch3760 /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 N Perc					Percent
cmGgLtvsGt7F * ch3760	93	23,3%	307	76,8%	400	100,0%

# cmGgLtvsGt7F \* ch3760 Crosstabulation

#### Count

		ch3			
		normal	patology	Total	
cmGgLtvsGt7F	<7	33	38	71	
	>7	10 12		22	
	Total	43	50	93	

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	,007 <sup>a</sup>	1	,933

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 10,17.

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>D</sup>	,000	1	1,000		
Likelihood Ratio	,007	1	,933		
Fisher's Exact Test				1,000	,565
Linear-by-Linear Association	,007	1	,933		
N of Valid Cases	93				

## b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value			
Odds Ratio for cmGgLtvsGt7F (<7 / >7)	1,042	,399	2,722	
For cohort ch3760 = normal	1,023	,607	1,722	
For cohort ch3760 = patology	,981	,633	1,522	
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	,007	1	,933
Mantel-Haenszel	,025	1	,873

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	1,042
		In(Estimate)	,041
		Std. Error of In(Estimate)	,490
		Asymp. Sig. (2-sided)	,933
- /	Common Odds Ratio	Lower Bound	,399
Interval		Upper Bound	2,722
	In(Common Odds Ratio)	Lower Bound	-,919
		Upper Bound	1,001

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

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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=cmGgLt7vsEq7F BY ch3760 /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		Total		
	N	Percent	N	Percent	N	Percent
cmGgLt7vsEq7F * ch3760	128	32,0%	272	68,0%	400	100,0%

# cmGgLt7vsEq7F \* ch3760 Crosstabulation

# Count

		ch3	ch3760		
		normal	Total		
cmGgLt7vsEq7F	<7	33	38	71	
	=7	27 30		57	
	Total	60	68	128	

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,010 <sup>a</sup>	1	,920		
Continuity Correction b	,000	1	1,000		
Likelihood Ratio	,010	1	,920		
Fisher's Exact Test				1,000	,531
Linear-by-Linear Association	,010	1	,920		
N of Valid Cases	128				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 26,72.

## **Risk Estimate**

		95% Confidence Interval		
	Value	Lower Upper		
Odds Ratio for cmGgLt7vsEq7F (<7 / =7)	,965	,480	1,940	
For cohort ch3760 = normal	,981	,678	1,421	
For cohort ch3760 = patology	1,017	,732	1,412	
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	,010	1	,920
Mantel-Haenszel	,006	1	,938

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	,965
		In(Estimate)	-,036
		Std. Error of In(Estimate)	,356
		Asymp. Sig. (2-sided)	,920
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,480
Interval		Upper Bound	1,940
	In(Common Odds Ratio)	Lower Bound	-,734
		Upper Bound	,663

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

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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=cmGgEq7vsGt7F BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.031
	Elapsed Time	0:00:00.019
	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 Percei				N	Percent
cmGgEq7vsGt7F * ch3760	79	19,8%	321	80,3%	400	100,0%

# cmGgEq7vsGt7F \* ch3760 Crosstabulation

### Count

Count						
		ch3				
		normal	Total			
cmGgEq7vsGt7F	=7	27	30	57		
	>7	10	12	22		
	Total	37	42	79		

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,023 <sup>a</sup>	1	,879		
Continuity Correction b	,000	1	1,000		
Likelihood Ratio	,023	1	,878		
Fisher's Exact Test				1,000	,540
Linear-by-Linear Association	,023	1	,879		
N of Valid Cases	79				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 10,30.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interval		
	Value	Lower Upper		
Odds Ratio for cmGgEq7vsGt7F (=7 / >7)	1,080	,402	2,898	
For cohort ch3760 = normal	1,042	,611	1,776	
For cohort ch3760 = patology	,965	,613	1,519	
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	,023	1	,879
Mantel-Haenszel	,010	1	,922

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	1,080
		In(Estimate)	,077
		Std. Error of In(Estimate)	,504
		Asymp. Sig. (2-sided)	,879
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,402
Interval		Upper Bound	2,898
	In(Common Odds Ratio)	Lower Bound	-,910
		Upper Bound	1,064

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	0	00 5- 0040 40:40:50
	Output Created	22-lip-2012 12:18:52
	Comments	
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	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=mMeta BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
	Elapsed Time	0:00:00.016
	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				Percent		
mMeta * ch3760	150	N Percent N Percent					

#### mMeta \* ch3760 Crosstabulation

#### Count

Count					
		ch3760			
		normal	Total		
mMeta	no	48	47	95	
	yes	22	33	55	
	Total	70	80	150	

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	1,551 <sup>a</sup>	1	,213

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 25,67.

## **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>D</sup>	1,157	1	,282		
Likelihood Ratio	1,558	1	,212		
Fisher's Exact Test				,238	,141
Linear-by-Linear Association	1,540	1	,215		
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)	1,532	,782	3,002	
For cohort ch3760 = normal	1,263	,864	1,847	
For cohort ch3760 = patology	,825	,613	1,109	
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	1,551	1	,213
Mantel-Haenszel	1,149	1	,284

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	1,532
		In(Estimate)	,427
		Std. Error of In(Estimate)	,343
		Asymp. Sig. (2-sided)	,214
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,782
Interval		Upper Bound	3,002
	In(Common Odds Ratio)	Lower Bound	-,246
		Upper Bound	1,099

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=mRiskEAU BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:18:52
	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=mRiskEAU BY ch3760 /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

The Tests for Homogeneity of the Odds Ratio table and the Mantel-Haenszel Common Odds Ratio Estimate table are not computed for mRiskEAU \* ch3760, 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					
	Va	Valid Missing Total			tal		
	N	N Percent N Percent		N	Percent		
mRiskEAU * ch3760	150	37,5%	250	62,5%	400	100,0%	

#### mRiskEAU \* ch3760 Crosstabulation

#### Count

Count						
		ch3				
		normal	Total			
mRiskEAU	low	6	8	14		
	medium	28	27	55		
	high	36	45	81		
	Total	70	80	150		

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	,640 <sup>a</sup>	2	,726
Likelihood Ratio	,640	2	,726
Linear-by-Linear Association	,098	1	,754
N of Valid Cases	150		

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,53.

#### **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 ch3760

/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ RISK CMH(1)

/CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

### Notes

	Output Created	22-lip-2012 12:18:52
	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=mRiskEAULowMedium BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.032
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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	Ν	Percent	
mRiskEAULowMedium * ch3760	69	17,3%	331	82,8%	400	100,0%	

# mRiskEAULowMedium \* ch3760 Crosstabulation

#### Count

		ch3760		
		normal	patology	Total
mRiskEAULowMedium	low	6	8	14
	medium	28	27	55
	Total	34	35	69

### **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,289 <sup>a</sup>	1	,591		
Continuity Correction b	,057	1	,811		
Likelihood Ratio	,290	1	,590		
Fisher's Exact Test				,766	,406
Linear-by-Linear Association	,285	1	,593		
N of Valid Cases	69				

- a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,90.
- b. Computed only for a 2x2 table

#### **Risk Estimate**

		95% Confidence Interva		
	Value	Lower	Upper	
Odds Ratio for mRiskEAULowMedium (low / medium)	,723	,222	2,361	
For cohort ch3760 = normal	,842	,436	1,626	
For cohort ch3760 = patology	1,164	,687	1,973	
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	,289	1	,591
Mantel-Haenszel	,056	1	,813

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	,723
In(Estimate)	-,324
Std. Error of In(Estimate)	,604
Asymp. Sig. (2-sided)	,591

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	,222
Interval		Upper Bound	2,361
	In(Common Odds Ratio)	Lower Bound	-1,507
		Upper Bound	,859

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT

/COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:18:53
	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=mRiskEAULowHigh BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

Resources	Processor Time	0:00:00.016	T
	Elapsed Time	0:00:00.098	ı
	Dimensions Requested	2	ı
	Cells Available	174762	ı

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

# **Case Processing Summary**

		Cases				
	Valid		Missing		Total	
	N	Percent	Ν	Percent	N	Percent
mRiskEAULowHigh * ch3760	95	23,8%	305	76,3%	400	100,0%

# mRiskEAULowHigh \* ch3760 Crosstabulation

#### Count

Count		ch3760		
		normal	patology	Total
mRiskEAULowHigh	low	6	8	14
	high	36	45	81
	Total	42	53	95

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	,012 <sup>a</sup>	1	,912		
Continuity Correction b	,000	1	1,000		
Likelihood Ratio	,012	1	,912		
Fisher's Exact Test				1,000	,575
Linear-by-Linear Association	,012	1	,913		
N of Valid Cases	95				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 6,19.

### **Risk Estimate**

		95% Confidence Interva	
	Value	Lower	Upper
Odds Ratio for mRiskEAULowHigh (low / high)	,938	,298	2,948
For cohort ch3760 = normal	,964	,502	1,851
For cohort ch3760 = patology	1,029	,628	1,685
N of Valid Cases	95		

b. Computed only for a 2x2 table

### 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	,012	1	,912
Mantel-Haenszel	,032	1	,857

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	,938
		In(Estimate)	-,065
		Std. Error of In(Estimate)	,585
		Asymp. Sig. (2-sided)	,912
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,298
Interval		Upper Bound	2,948
	In(Common Odds Ratio)	Lower Bound	-1,210
		Upper Bound	1,081

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

Output Created	22-lip-2012 12:18:53
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=mRiskEAUMediumHigh BY ch3760 /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					
	Va	lid	Miss	Missing		tal
	N	Percent	N	Percent	Ν	Percent
mRiskEAUMediumHigh * ch3760	136	34,0%	264	66,0%	400	100,0%

# mRiskEAUMediumHigh \* ch3760 Crosstabulation

# Count

		ch3760		
		normal	patology	Total
mRiskEAUMediumHigh	medium	28	27	55
	high	36	45	81
	Total	64	72	136

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	,549 <sup>a</sup>	1	,459

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 25,88.

### **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Continuity Correction <sup>0</sup>	,321	1	,571		
Likelihood Ratio	,549	1	,459		
Fisher's Exact Test				,488	,286
Linear-by-Linear Association	,545	1	,460		
N of Valid Cases	136				

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

#### **Risk Estimate**

		95% Confide	nce Interval
	Value	Lower	Upper
Odds Ratio for mRiskEAUMediumHigh (medium / high)	1,296	,652	2,576
For cohort ch3760 = normal	1,145	,802	1,635
For cohort ch3760 = patology	,884	,634	1,232
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	,549	1	,459
Mantel-Haenszel	,318	1	,573

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	1,296
In(Estimate)	,260
Std. Error of In(Estimate)	,350
Asymp. Sig. (2-sided)	,459

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	,652
Interval		Upper Bound	2,576
	In(Common Odds Ratio)	Lower Bound	-,427
		Upper Bound	,946

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

#### **Notes**

	Output Created	22-lip-2012 12:18:53
	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=mRiskMed BY ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.011
	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	N Percent N Percent		N	Percent	
mRiskMed * ch3760	150	37,5%	250	62,5%	400	100,0%

#### mRiskMed \* ch3760 Crosstabulation

### Count

		ch3760		
		normal	patology	Total
mRiskMed	low	30	25	55
	high	40	55	95
	Total	70	80	150

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	2,166 <sup>a</sup>	1	,141		
Continuity Correction b	1,695	1	,193		
Likelihood Ratio	2,166	1	,141		
Fisher's Exact Test				,175	,096
Linear-by-Linear Association	2,151	1	,142		
N of Valid Cases	150				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 25,67.

# **Risk Estimate**

		95% Confidence Interv	
	Value	Lower	Upper
Odds Ratio for mRiskMed (low / high)	1,650	,845	3,221
For cohort ch3760 = normal	1,295	,925	1,815
For cohort ch3760 = patology	,785	,561	1,099
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	2,166	1	,141
Mantel-Haenszel	1,684	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	1,650
		In(Estimate)	,501
		Std. Error of In(Estimate)	,341
		Asymp. Sig. (2-sided)	,142
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,845
Interval		Upper Bound	3,221
	In(Common Odds Ratio)	Lower Bound	-,168
		Upper Bound	1,170

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:18:54
	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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.015
	Elapsed Time	0:00:00.013
	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 \* ch3760. 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	
	N	Percent	N	Percent	N	Percent
mRiskMedLowMedium * ch3760	55	13,8%	345	86,3%	400	100,0%

### mRiskMedLowMedium \* ch3760 Crosstabulation

#### Count

Oddit				
		ch3760		
		normal	patology	Total
mRiskMedLowMedium	low	30	25	55
	Total	30	25	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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

#### **Notes**

	Output Created	22-lip-2012 12:18:54
	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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
	Elapsed Time	0:00:00.238
	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**

			Cas	ses		
	Va	lid	Missing		Total	
	N	Percent	N	Percent	Ν	Percent
mRiskMedLowHigh * ch3760	150	37,5%	250	62,5%	400	100,0%

# mRiskMedLowHigh \* ch3760 Crosstabulation

#### Count

Obdite				
		ch3	ch3760	
		normal	patology	Total
mRiskMedLowHigh	low	30	25	55
	high	40	55	95
	Total	70	80	150

# **Chi-Square Tests**

	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	2,166 <sup>a</sup>	1	,141		
Continuity Correction b	1,695	1	,193		
Likelihood Ratio	2,166	1	,141		
Fisher's Exact Test				,175	,096
Linear-by-Linear Association	2,151	1	,142		
N of Valid Cases	150				

a. 0 cells (,0%) have expected count less than 5. The minimum expected count is 25,67.

#### **Risk Estimate**

		95% Confide	nce Interval
	Value	Lower	Upper
Odds Ratio for mRiskMedLowHigh (low / high)	1,650	,845	3,221
For cohort ch3760 = normal	1,295	,925	1,815
For cohort ch3760 = patology	,785	,561	1,099
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	2,166	1	,141
Mantel-Haenszel	1,684	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	1,650
		In(Estimate)	,501
		Std. Error of In(Estimate)	,341
		Asymp. Sig. (2-sided)	,142
Asymp. 95% Confidence	Common Odds Ratio	Lower Bound	,845
Interval		Upper Bound	3,221
	In(Common Odds Ratio)	Lower Bound	-,168
		Upper Bound	1,170

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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.

# **Crosstabs**

	Output Created	22-lip-2012 12:18:54
	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 ch3760 /FORMAT=AVALUE TABLES /STATISTICS=CHISQ RISK CMH(1) /CELLS=COUNT /COUNT ROUND CELL.
Resources	Processor Time	0:00:00.016
	Elapsed Time	0:00:00.094
	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 \* ch3760. 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	Percent	
mRiskMedMediumHigh * ch3760	95	23,8%	305	76,3%	400	100,0%

### mRiskMedMediumHigh \* ch3760 Crosstabulation

#### Count

Count				
		ch3	760	
		normal	patology	Total
mRiskMedMediumHigh	high	40	55	95
	Total	40	55	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.