

## **Supplementary Materials**

### **New prognostic frontiers for lung neuroendocrine tumors**

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**Short title:** New prognostic frontiers for lung N

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**Supplementary results. Associations with age, peripheral versus central location of the primary tumor and necrosis**

**1. AGE**

**Case Processing Summary**

	Cases					
	Valid		Missing		Total	
	N	Percent	N	Percent	N	Percent
Age * Smoke yes 1 no 0	165	55,0%	135	45,0%	300	100,0%
Age* Central 1 vs Periferic 0 primary lesion	200	66,7%	100	33,3%	300	100,0%
Age * 18 FDG PET scan Not done Positive 1 Negative 0	103	34,3%	197	65,7%	300	100,0%
Age* Octreoscan/68GalliumPET Not done NA Positive 1 Negative 0	63	21,0%	237	79,0%	300	100,0%
Age * Tcat	192	64,0%	108	36,0%	300	100,0%
Age * Npos1neg0	190	63,3%	110	36,7%	300	100,0%
Age * Stage	196	65,3%	104	34,7%	300	100,0%
Age * Ki67	165	55,0%	135	45,0%	300	100,0%
Age* Necrosis Not done NA Positive 1 Negative 0	199	66,3%	101	33,7%	300	100,0%
Age* Diagnosis Typical carcinoid 0 Atypical carcinoid 1	200	66,7%	100	33,3%	300	100,0%
Age * Typesurgery	178	59,3%	122	40,7%	300	100,0%

**Count**

		Smoke yes 1 no 0		Total
		0	1	
Age	<= MED	47	37	84
	>MED	55	26	81
Total		102	63	165

### Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	2,494 <sup>a</sup>	1	0,114		
Continuity Correction <sup>b</sup>	2,014	1	0,156		
Likelihood Ratio	2,504	1	0,114		
Fisher's Exact Test				0,149	0,078
Linear-by-Linear Association	2,479	1	0,115		
N of Valid Cases	165				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 30,93.

b. Computed only for a 2x2 table

### Crosstab

#### Count

		Central 1 vs Periferic 0 primary lesion		
		0	1	Total
Age	< = MED	27	76	103
	>MED	44	53	97
Total		71	129	200

### Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	7,998 <sup>a</sup>	1	0,005		
Continuity Correction <sup>b</sup>	7,184	1	0,007		
Likelihood Ratio	8,053	1	0,005		
Fisher's Exact Test				0,005	0,004
Linear-by-Linear Association	7,958	1	0,005		
N of Valid Cases	200				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 34,43.

b. Computed only for a 2x2 table

Crosstab

Count

		18 FDG PET scan Not done Positive 1 Negative 0		
		0	1	Total
Age	< = MED	12	36	48
	>MED	14	41	55
Total		26	77	103

Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	0,003 <sup>a</sup>	1	0,958		
Continuity Correction <sup>b</sup>	0,000	1	1,000		
Likelihood Ratio	0,003	1	0,958		
Fisher's Exact Test				1,000	0,570
Linear-by-Linear Association	0,003	1	0,958		
N of Valid Cases	103				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 12,12.

b. Computed only for a 2x2 table

Crosstab

Count

		Octreoscan/68GalliumPET Not done NA Positive 1 Negative 0		
		0	1	Total
Age	< = MED	14	24	38
	>MED	8	17	25
Total		22	41	63

### Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	0,156 <sup>a</sup>	1	0,693		
Continuity Correction <sup>b</sup>	0,015	1	0,901		
Likelihood Ratio	0,156	1	0,693		
Fisher's Exact Test				0,790	0,453
Linear-by-Linear Association	0,153	1	0,696		
N of Valid Cases	63				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 8,73.

b. Computed only for a 2x2 table

### Crosstab

#### Count

		Tcat				
		1	2	3	4	Total
Age	< = MED	53	33	10	3	99
	>MED	63	23	4	3	93
Total		116	56	14	6	192

### Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)
Pearson Chi-Square	5,037 <sup>a</sup>	3	0,169
Likelihood Ratio	5,128	3	0,163
Linear-by-Linear Association	3,152	1	0,076
N of Valid Cases	192		

a. 2 cells (25,0%) have expected count less than 5. The minimum expected count is 2,91.

**Crosstab**

**Count**

		NposIneg0		
		0	1	Total
Age	< = MED	68	31	99
	>MED	72	19	91
Total		140	50	190

**Chi-Square Tests**

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	2,662 <sup>a</sup>	1	0,103		
Continuity Correction <sup>b</sup>	2,151	1	0,142		
Likelihood Ratio	2,686	1	0,101		
Fisher's Exact Test				0,137	0,071
Linear-by-Linear Association	2,648	1	0,104		
N of Valid Cases	190				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 23,95.

b. Computed only for a 2x2 table

**Crosstab**

**Count**

		Stage				
		1	2	3	4	Total
Age	< = MED	51	33	8	7	99
	>MED	60	23	9	5	97
Total		111	56	17	12	196

### Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	2,887 <sup>a</sup>	3	0,409
Likelihood Ratio	2,899	3	0,407
Linear-by-Linear Association	1,065	1	0,302
N of Valid Cases	196		

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 5,94.

### Crosstab

#### Count

		Ki67			Total
		1	2	3	
Age	< = MED	49	35	4	88
	>MED	34	39	4	77
Total		83	74	8	165

### Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	2,204 <sup>a</sup>	2	0,332
Likelihood Ratio	2,208	2	0,332
Linear-by-Linear Association	1,754	1	0,185
N of Valid Cases	165		

a. 2 cells (33,3%) have expected count less than 5. The minimum expected count is 3,73.

Crosstab

Count

		Necrosis Not done NA Positive 1 Negative 0		Total
		0	1	
Age	< = MED	83	20	103
	>MED	82	14	96
Total		165	34	199

Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	0,820 <sup>a</sup>	1	0,365		
Continuity Correction <sup>b</sup>	0,514	1	0,473		
Likelihood Ratio	0,824	1	0,364		
Fisher's Exact Test				0,452	0,237
Linear-by-Linear Association	0,816	1	0,366		
N of Valid Cases	199				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 16,40.

b. Computed only for a 2x2 table

Crosstab

Count

		Diagnosis Typical carcinoid 0 Atypical carcinoid 1		Total
		0	1	
Age	< = MED	67	36	103
	>MED	72	25	97
Total		139	61	200



### Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	1,985 <sup>a</sup>	1	0,159		
Continuity Correction <sup>b</sup>	1,576	1	0,209		
Likelihood Ratio	1,994	1	0,158		
Fisher's Exact Test				0,170	0,105
Linear-by-Linear Association	1,975	1	0,160		
N of Valid Cases	200				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 29,59.

b. Computed only for a 2x2 table

### Count

		Typesurgery								
		pneumone ctomy	bilobecto my	lobecto my	sleeve resection	segmental resection	wedge resection	extended resection	other	Total
Age	<= MED	7	8	62	2	3	5	4	1	92
	>MED	4	2	67	0	2	9	2	0	86
Total		11	10	129	2	5	14	6	1	178

### Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)
Pearson Chi-Square	9,430 <sup>a</sup>	7	0,223
Likelihood Ratio	10,874	7	0,144
Linear-by-Linear Association	0,245	1	0,620
N of Valid Cases	178		

a. 9 cells (56,3%) have expected count less than 5. The minimum expected count is 0,48.

## 2. TUMOR LOCATION: PERIPHERAL VERSUS CENTRAL

### Case Processing Summary

	Cases				Total	
	Valid		Missing		N	Percent
	N	Percent	N	Percent		
Central 1 vs Periferic 0 primary lesion * Smoke yes 1 no 0	246	82,0 %	54	18,0 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * 18 FDG PET scan Not done Positive 1 Negative 0	200	66,7 %	100	33,3 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Octreoscan/68Gallium PET Not done NA Positive 1 Negative 0	64	21,3 %	236	78,7 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Tcat	276	92,0 %	24	8,0 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Npos1neg0	265	88,3 %	35	11,7 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Stage	274	91,3 %	26	8,7 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Ki67	231	77,0 %	69	23,0 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Necrosis Not done NA Positive 1 Negative 0	293	97,7 %	7	2,3 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Diagnosis Typical carcinoid 0 Atypical carcinoid 1	298	99,3 %	2	0,7 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Typesurgery	276	92,0 %	24	8,0 %	300	100,0 %
Central 1 vs Periferic 0 primary lesion * Age	200	66,7 %	100	33,3 %	300	100,0 %

Crosstab

Count

		Smoke yes 1 no 0		Total
		0	1	
Central 1 vs Periferic 0 primary lesion	0	65	37	102
	1	77	67	144
Total		142	104	246

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	2,572 <sup>a</sup>	1	0,109		
Continuity Correction <sup>b</sup>	2,169	1	0,141		
Likelihood Ratio	2,587	1	0,108		
Fisher's Exact Test				0,118	0,070
Linear-by-Linear Association	2,562	1	0,109		
N of Valid Cases	246				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 43,12.

b. Computed only for a 2x2 table

Crosstab

Count

		18 FDG PET scan Not done		Total
		Positive 1	Negative 0	
Central 1 vs Periferic 0 primary lesion	0	21	62	83
	1	51	66	117
Total		72	128	200

**Chi-Square Tests**

	<b>Value</b>	<b>df</b>	<b>Asymptotic Significance (2- sided)</b>	<b>Exact Sig. (2-sided)</b>	<b>Exact Sig. (1-sided)</b>
<b>Pearson Chi-Square</b>	<b>7,049<sup>a</sup></b>	<b>1</b>	<b>0,008</b>		
<b>Continuity Correction<sup>b</sup></b>	<b>6,277</b>	<b>1</b>	<b>0,012</b>		
<b>Likelihood Ratio</b>	<b>7,206</b>	<b>1</b>	<b>0,007</b>		
<b>Fisher's Exact Test</b>				<b>0,011</b>	<b>0,006</b>
<b>Linear-by-Linear Association</b>	<b>7,013</b>	<b>1</b>	<b>0,008</b>		
<b>N of Valid Cases</b>	<b>200</b>				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 29,88.

b. Computed only for a 2x2 table

**Count**

		<b>Octreoscan/68GalliumPET Not done NA Positive 1 Negative 0</b>		<b>Total</b>
		<b>0</b>	<b>1</b>	
<b>Central 1 vs Periferic 0 primary lesion</b>	<b>0</b>	<b>9</b>	<b>13</b>	<b>22</b>
	<b>1</b>	<b>13</b>	<b>29</b>	<b>42</b>
<b>Total</b>		<b>22</b>	<b>42</b>	<b>64</b>

### Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)	Exact Sig. (2-sided)	Exact Sig. (1- sided)
<b>Pearson Chi-Square</b>	<b>0,634<sup>a</sup></b>	<b>1</b>	<b>0,426</b>		
<b>Continuity Correction<sup>b</sup></b>	<b>0,270</b>	<b>1</b>	<b>0,603</b>		
<b>Likelihood Ratio</b>	<b>0,627</b>	<b>1</b>	<b>0,428</b>		
<b>Fisher's Exact Test</b>				<b>0,580</b>	<b>0,300</b>
<b>Linear-by-Linear Association</b>	<b>0,625</b>	<b>1</b>	<b>0,429</b>		
<b>N of Valid Cases</b>	<b>64</b>				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 7,56.

b. Computed only for a 2x2 table

### Crosstab

#### Count

		Tcat				Total
		1	2	3	4	
<b>Central 1 vs Periferic 0 primary lesion</b>	<b>0</b>	<b>72</b>	<b>22</b>	<b>7</b>	<b>2</b>	<b>103</b>
	<b>1</b>	<b>103</b>	<b>56</b>	<b>9</b>	<b>5</b>	<b>173</b>
<b>Total</b>		<b>175</b>	<b>78</b>	<b>16</b>	<b>7</b>	<b>276</b>

**Chi-Square Tests**

	<b>Value</b>	<b>df</b>	<b>Asymptotic Significance (2- sided)</b>
<b>Pearson Chi-Square</b>	<b>4,375<sup>a</sup></b>	<b>3</b>	<b>0,224</b>
<b>Likelihood Ratio</b>	<b>4,480</b>	<b>3</b>	<b>0,214</b>
<b>Linear-by-Linear Association</b>	<b>1,415</b>	<b>1</b>	<b>0,234</b>
<b>N of Valid Cases</b>	<b>276</b>		

a. 2 cells (25,0%) have expected count less than 5. The minimum expected count is 2,61.

**Crosstab**

**Count**

		<b>Npos1neg0</b>		
		<b>0</b>	<b>1</b>	<b>Total</b>
<b>Central 1 vs Periferic 0 primary lesion</b>	<b>0</b>	<b>70</b>	<b>27</b>	<b>97</b>
	<b>1</b>	<b>130</b>	<b>38</b>	<b>168</b>
<b>Total</b>		<b>200</b>	<b>65</b>	<b>265</b>

**Chi-Square Tests**

	<b>Value</b>	<b>df</b>	<b>Asymptotic Significance (2- sided)</b>	<b>Exact Sig. (2-sided)</b>	<b>Exact Sig. (1-sided)</b>
<b>Pearson Chi-Square</b>	<b>0,904<sup>a</sup></b>	<b>1</b>	<b>0,342</b>		
<b>Continuity Correction<sup>b</sup></b>	<b>0,644</b>	<b>1</b>	<b>0,422</b>		
<b>Likelihood Ratio</b>	<b>0,894</b>	<b>1</b>	<b>0,344</b>		
<b>Fisher's Exact Test</b>				<b>0,375</b>	<b>0,210</b>
<b>Linear-by-Linear Association</b>	<b>0,900</b>	<b>1</b>	<b>0,343</b>		
<b>N of Valid Cases</b>	<b>265</b>				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 23,79.

b. Computed only for a 2x2 table

**Crosstab**

**Count**

		<b>Stage</b>				<b>Total</b>
		<b>1</b>	<b>2</b>	<b>3</b>	<b>4</b>	
<b>Central 1 vs Periferic 0 primary lesion</b>	<b>0</b>	<b>62</b>	<b>21</b>	<b>10</b>	<b>6</b>	<b>99</b>
	<b>1</b>	<b>108</b>	<b>49</b>	<b>12</b>	<b>6</b>	<b>175</b>
<b>Total</b>		<b>170</b>	<b>70</b>	<b>22</b>	<b>12</b>	<b>274</b>

### Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	2,978 <sup>a</sup>	3	0,395
Likelihood Ratio	2,946	3	0,400
Linear-by-Linear Association	0,542	1	0,462
N of Valid Cases	274		

a. 1 cells (12,5%) have expected count less than 5. The minimum expected count is 4,34.

### Crosstab

#### Count

		Ki67			Total
		1	2	3	
Central 1 vs Periferic 0 primary lesion	0	59	23	2	84
	1	83	57	7	147
Total		142	80	9	231

### Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	4,432 <sup>a</sup>	2	0,109
Likelihood Ratio	4,536	2	0,104
Linear-by-Linear Association	4,312	1	0,038
N of Valid Cases	231		

a. 1 cells (16,7%) have expected count less than 5. The minimum expected count is 3,27.



Crosstab

Count

		Necrosis Not done NA Positive 1		Total
		Negative 0	1	
Central 1 vs Periferic 0 primary lesion	0	97	12	109
	1	157	27	184
Total		254	39	293

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	0,797 <sup>a</sup>	1	0,372		
Continuity Correction <sup>b</sup>	0,511	1	0,475		
Likelihood Ratio	0,815	1	0,367		
Fisher's Exact Test				0,477	0,239
Linear-by-Linear Association	0,794	1	0,373		
N of Valid Cases	293				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 14,51.

b. Computed only for a 2x2 table

Crosstab

Count

		Diagnosis Typical carcinoid 0		Total
		Atypical carcinoid 1	1	
Central 1 vs Periferic 0 primary lesion	0	87	25	112
	1	136	50	186
Total		223	75	298

**Chi-Square Tests**

	<b>Value</b>	<b>df</b>	<b>Asymptotic Significance (2- sided)</b>	<b>Exact Sig. (2-sided)</b>	<b>Exact Sig. (1-sided)</b>
<b>Pearson Chi-Square</b>	<b>0,772<sup>a</sup></b>	<b>1</b>	<b>0,380</b>		
<b>Continuity Correction<sup>b</sup></b>	<b>0,549</b>	<b>1</b>	<b>0,459</b>		
<b>Likelihood Ratio</b>	<b>0,781</b>	<b>1</b>	<b>0,377</b>		
<b>Fisher's Exact Test</b>				<b>0,411</b>	<b>0,230</b>
<b>Linear-by-Linear Association</b>	<b>0,769</b>	<b>1</b>	<b>0,380</b>		
<b>N of Valid Cases</b>	<b>298</b>				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 28,19.

b. Computed only for a 2x2 table

### 3. NECROSIS

#### Case Processing Summary

	Cases				Total	
	Valid		Missing		N	Percent
	N	Percent	N	Percent		
Necrosis Not done NA Positive 1 Negative 0 * Smoke yes 1 no 0	245	81,7%	55	18,3%	300	100,0%
Necrosis Not done NA Positive 1 Negative 0 * 18 FDG PET scan Not done Positive 1 Negative 0	198	66,0%	102	34,0%	300	100,0%
Necrosis Not done NA Positive 1 Negative 0 * Octreoscan/68Ga IliumPET Not done NA Positive 1 Negative 0	64	21,3%	236	78,7%	300	100,0%
Necrosis Not done NA Positive 1 Negative 0 * Tcat	274	91,3%	26	8,7%	300	100,0%
Necrosis Not done NA Positive 1 Negative 0 * NposIneg0	264	88,0%	36	12,0%	300	100,0%
Necrosis Not done NA Positive 1 Negative 0 * Stage at diagnosis	295	98,3%	5	1,7%	300	100,0%
Necrosis Not done NA Positive 1 Negative 0 * Gruppoki67	229	76,3%	71	23,7%	300	100,0%

**Crosstab**

**Count**

		Smoke yes 1 no 0		Total
		0	1	
Necrosis Not done NA	0	126	81	207
Positive 1 Negative 0	1	16	22	38
<b>Total</b>		<b>142</b>	<b>103</b>	<b>245</b>

**Chi-Square Tests**

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	4,639 <sup>a</sup>	1	0,031		
Continuity Correction <sup>b</sup>	3,901	1	0,048		
Likelihood Ratio	4,578	1	0,032		
Fisher's Exact Test				0,048	0,025
Linear-by-Linear Association	4,620	1	0,032		
N of Valid Cases	245				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 15,98.

b. Computed only for a 2x2 table

**Crosstab**

**Count**

		18 FDG PET scan Not done		Total
		Positive 1 Negative 0	0	
Necrosis Not done NA	0	56	116	172
Positive 1 Negative 0	1	12	14	26
<b>Total</b>		<b>68</b>	<b>130</b>	<b>198</b>

**Chi-Square Tests**

	<b>Value</b>	<b>df</b>	<b>Asymptotic Significance (2-sided)</b>	<b>Exact Sig. (2- sided)</b>	<b>Exact Sig. (1- sided)</b>
<b>Pearson Chi-Square</b>	<b>1,851<sup>a</sup></b>	<b>1</b>	<b>0,174</b>		
<b>Continuity Correction<sup>b</sup></b>	<b>1,298</b>	<b>1</b>	<b>0,255</b>		
<b>Likelihood Ratio</b>	<b>1,786</b>	<b>1</b>	<b>0,181</b>		
<b>Fisher's Exact Test</b>				<b>0,189</b>	<b>0,128</b>
<b>Linear-by-Linear Association</b>	<b>1,842</b>	<b>1</b>	<b>0,175</b>		
<b>N of Valid Cases</b>	<b>198</b>				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 8,93.

b. Computed only for a 2x2 table

**Crosstab**

**Count**

		<b>Octreoscan/68GalliumPET Not done NA</b>		<b>Total</b>
		<b>Positive 0</b>	<b>Negative 1</b>	
<b>Necrosis Not done NA</b>	<b>0</b>	<b>18</b>	<b>32</b>	<b>50</b>
<b>Positive 1 Negative 0</b>	<b>1</b>	<b>4</b>	<b>10</b>	<b>14</b>
<b>Total</b>		<b>22</b>	<b>42</b>	<b>64</b>

### Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	0,268 <sup>a</sup>	1	0,605		
Continuity Correction <sup>b</sup>	0,040	1	0,842		
Likelihood Ratio	0,274	1	0,601		
Fisher's Exact Test				0,755	0,429
Linear-by-Linear Association	0,263	1	0,608		
N of Valid Cases	64				

a. 1 cells (25,0%) have expected count less than 5. The minimum expected count is 4,81.

b. Computed only for a 2x2 table

### Crosstab

#### Count

		Tcat				Total
		1	2	3	4	
Necrosis Not done NA	0	168	56	9	3	236
Positive 1 Negative 0	1	7	20	7	4	38
<b>Total</b>		<b>175</b>	<b>76</b>	<b>16</b>	<b>7</b>	<b>274</b>

### Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	47,059 <sup>a</sup>	3	0,000
Likelihood Ratio	42,735	3	0,000
Linear-by-Linear Association	46,326	1	0,000
N of Valid Cases	274		

a. 2 cells (25,0%) have expected count less than 5. The minimum expected count is ,97.

**Crosstab**

**Count**

		Npos1neg0		Total
		0	1	
Necrosis Not done NA	0	180	47	227
Positive 1 Negative 0	1	19	18	37
<b>Total</b>		<b>199</b>	<b>65</b>	<b>264</b>

**Chi-Square Tests**

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	13,386 <sup>a</sup>	1	,000		
Continuity Correction <sup>b</sup>	11,922	1	,001		
Likelihood Ratio	11,881	1	,001		
Fisher's Exact Test				,001	,001
Linear-by-Linear Association	13,335	1	,000		
N of Valid Cases	264				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 9,11.

b. Computed only for a 2x2 table

**Crosstab**

**Count**

		Gruppoki67			Total
		1	2	3	
Necrosis Not done NA	0	135	54	5	194
Positive 1 Negative 0	1	7	25	3	35
<b>Total</b>		<b>142</b>	<b>79</b>	<b>8</b>	<b>229</b>

Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)
Pearson Chi-Square	31,141 <sup>a</sup>	2	,000
Likelihood Ratio	30,850	2	,000
Linear-by-Linear Association	29,173	1	,000
N of Valid Cases	229		

a. 1 cells (16,7%) have expected count less than 5. The minimum expected count is 1,22.

Crosstab

Count

		Diagnosis Typical carcinoid 0 Atypical carcinoid 1		Total
		0	1	
Necrosis Not done NA Positive 1 Negative 0	0	225	31	256
	1	0	39	39
Total		225	70	295

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2- sided)	Exact Sig. (1- sided)
Pearson Chi-Square	144,455 <sup>a</sup>	1	0,000		
Continuity Correction <sup>b</sup>	139,639	1	0,000		
Likelihood Ratio	134,302	1	0,000		
Fisher's Exact Test				0,000	0,000
Linear-by- Linear Association	143,965	1	0,000		
N of Valid Cases	295				

a. 0 cells (0,0%) have expected count less than 5. The minimum expected count is 9,25.

b. Computed only for a 2x2 table



**Kaplan-Meier survival curves of: high mitotic count (> 10 per 10 HPF) versus mitotic count ≤ than 10 per 10 HPF in terms of PFS and OS (upper part of the figure, left and right side, respectively); and of high Ki67 value (> 20%) versus Ki67 lower than 20% in terms of both PFS and OS (lower part of the figure, left and right side, respectively). P-values have been considered as significant if < 0.05 and are reported according to the corresponding univariant COX regression analysis.**

