



Expression of oncoproteins and the amount of eosinophilic and lymphocytic infiltrates can be used as prognostic factors in gastric cancer

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Summary Preoperative staging of gastric cancer is difficult. Several molecular markers associated with initiation and progression of cancer seem promising for obtaining preoperative prognostic information. To investigate whether these markers are indicative especially for the presence of lymph node metastases in patients with gastric cancer, we have examined primary tumour specimens from 105 patients with primary adenocarcinoma of the stomach entered in a surgical trial. In this trial, lymph node status was determined by strictly quality-controlled lymph node dissection and examination. The selected markers were growth regulators (p53, Rb and *myc*), metastasis-suppressor gene product (nm23), adhesion molecules (Ep-CAM, E-cadherin, CD44v5 and CD44v6) and urokinase-type plasminogen activator (u-PA). Also, the amount of eosinophilic and lymphocytic infiltrates available post-operatively was analysed with respect to its prognostic value for lymph node status. Moreover, the association of these parameters with survival and disease-free period (DFP) was evaluated. Of all molecular markers investigated, only Rb expression had a significant association with the presence of lymph node metastasis in both univariate and multivariate analysis. For curative resectability, a significant association was found with Rb and E-cadherin expression, while in multivariate analysis Rb and *myc* were selected as the combination with additional independent prognostic value, and E-cadherin had no additional independent value. For overall survival in univariate analysis, the amount of both eosinophilic and lymphocytic infiltrates and Rb and *myc* expression were of significant prognostic value. Only the amount of lymphocytic infiltrate had a prognostic significance for DFP. In stepwise multivariate analysis, TNM stage (I+II) and marked lymphocytic infiltrate were associated with better overall survival and longer DFP. We conclude that, if these results are confirmed in a larger series of patients, molecular markers can provide useful prognostic information.

Keywords: gastric cancer; metastasis; prognosis; molecular marker

Despite declining incidence, gastric cancer remains a major clinical management problem with poor prognosis: overall 5 year survival rates vary between 5% and 11%. Only a curative resection (complete tumour removal) offers hope of a cure, but the majority of the patients are diagnosed at an advanced disease stage (Allum *et al.*, 1989). At diagnosis, the first decision to be made is whether or not to attempt a curative resection. This decision is based on stage of disease, which is assessed by radiograph of the stomach, chest and ultrasound of the liver or computerised tomography (CT) scan of the abdomen. Palliative resections offer no survival advantage and are associated with higher operative mortality rates compared with curative resections (Allum *et al.*, 1989; Akoh *et al.*, 1991; Wanebo *et al.*, 1993). Laparoscopy and cytological examination of abdominal washings increase the accuracy of preoperative staging, but are not used routinely (Bonenkamp *et al.*, 1996). Therefore, there is a need for additional, reliable prognostic factors by which tumour aggressiveness can be determined, and the presence of lymph node metastasis and resectability for cure can be predicted, preferably by analysis of biopsy specimens. This would allow better selection of patients suitable for surgery.

A variety of genetic changes are implicated in the process of carcinogenesis and the development of metastasis. These changes are associated with altered expression of proteins, which were reported to be of prognostic significance in different types of malignancies, including gastric cancer. However, not all of these tumour markers have been analysed in the same specimens and it is not known whether some of these markers are superior to others or whether a

combination of several markers has a better predictive value with respect to the presence of lymph node metastasis. As prognostic tumour markers, proteins known to be associated with different steps of carcinogenesis (growth promotion, loss of adhesion, invasion of gastric wall and vessels and distant metastasis) were analysed.

Our first aim was to study the association between the absence or presence of expression of these tumour markers and lymph node status. In addition, the results were also related to powerful clinicopathological prognostic factors, such as infiltration depth (T stage), TNM stage and curative resectability. Staging and curative resectability can be assessed reliably pre- and post-operatively only and are, therefore, of no use for patient selection for surgery.

A second aim was to study the prognostic value of these markers in addition to parameters only determinable post-operatively (eosinophilic and lymphocytic infiltrates) for survival and disease-free period (DFP).

Patients and methods

Patients

Between August 1989 and July 1993 a randomised, controlled multicentre trial was conducted in The Netherlands (Dutch Gastric Cancer Trial, DGCT) to compare the therapeutic efficacy of extended lymph node dissection (N1 and N2 levels, so-called D2) with that of limited lymph node dissection (N1 level, so-called D1) in gastric cancer patients operated on with curative intent. In this trial 1078 patients were entered, 996 of whom met the eligibility criteria. Strict quality control measures were taken to obtain optimum lymph node retrieval and, thus, post-operative staging. In patients undergoing a curative resection, the presence of nodal involvement was assessed histologically and the actual number and location of

the lymph nodes retrieved were recorded by the pathologist. Criteria for curative resectability were published earlier (Bonenkamp *et al.*, 1995). Median follow-up of the patients was 1287 days (range 672–2076 days). In the present study, tumour tissue of all patients ($n=105$) from six randomly selected hospitals out of the 78 participating hospitals was analysed. With respect to nodal involvement, only the presence or absence of nodal involvement was considered, and no attention was paid to the actual number or location of the dissected and involved lymph nodes. In 11 patients, no assessment of the nodal status was performed, because of locally advanced disease with invasion of adjacent organs, in which case a surgical resection was not possible and assessment of nodal status did not have immediate consequences for the patient. The primary tumour in resection specimens (available from 81 patients) was scored by one of the authors (JHJMvK) for the amount of lymphocytic and eosinophilic infiltration (Iwasaki *et al.*, 1986; Pretlow *et al.*, 1983; Jass, 1992).

Tissue specimens and immunohistochemistry

In order to study protein expression in primary adenocarcinomas of the stomach, formalin-fixed, paraffin-embedded tissue blocks of the primary tumour were used. According to the allocated treatment, the specimens were obtained by D1 or D2 resection. If resection was not possible, the specimen was obtained by biopsy of the primary tumour. All cases were reviewed and the tissue block with the largest tumour diameter was used to cut the required number of sections. For practical reasons, this was considered representative for the whole tumour.

Sections (4 μ m thick) were cut, mounted on precoated slides and kept at 37°C overnight. The sections were dewaxed in xylol for 20 min and endogenous peroxidase activity was blocked by methanol/hydrogen peroxide.

The antibodies used were p53 (mAb NCL-p53-DO7, Novocastra Laboratories), Rb (NCL-RB, retinoblastoma gene protein clone 1F8, Novocastra Laboratories), *myc* (NCL-cMYC, human c-myc protein clone 9E11, Novocastra Laboratories), nm23 (CRB-nm23 H1, H2, CRB), Ep-CAM (323/A3), Centocor, Malvern, PA, USA), E-cadherin (anti-E-cadherin clone HECD-1, Zymed Laboratories), CD44v5 and v6 (monoclonal antibodies VFF8 and VFF18 against CD44 variants containing splice variants v5 and v6 respectively, Bender, Vienna, Austria) and urokinase-type plasminogen activator (u-PA), a proteolytic factor (human urokinase monoclonal antibody against B-chain, American Diagnostica).

For p53, Rb, E-cadherin and CD44 splice variants, the sections were first boiled in citrate buffer (pH 6.0) for 25 min. For *myc* and Ep-CAM the sections were pretreated with trypsin (0.1% trypsin with 0.1% calcium chloride) pH 7.4 at 37°C for 20 min. Normal goat serum was applied to reduce non-specific antibody binding. The primary antibody was then applied and incubated overnight in its optimum dilutions in phosphate-buffered saline (PBS)/1% bovine serum albumin (BSA). The optimum dilutions for primary antibodies mentioned are summarised in Table I.

For E-cadherin, a double-step detection system was used: biotinylated rabbit anti-mouse IgG was followed by the streptavidin–biotin complex (each incubation 45 min). For the remaining monoclonal antibodies (Rb, *myc*, Ep-CAM and p53), a secondary antibody rabbit anti-mouse (RAM) was followed by a tertiary antibody swine anti-rabbit (SWAR). Nm23 (polyclonal antibody) was preincubated with normal goat serum for 10 min before applying SWAR. In negative controls the primary antibody was replaced by PBS. The sections were stained with 3-amino-9-ethylcarbazole (AEC) in dimethylformamide with hydrogen peroxide and counterstained with Mayer's haematoxylin.

Assessment and statistical analysis of marker expression

Expression of all proteins, except u-PA, was scored as present ($\geq 50\%$ of the tumour cells positive) or absent ($< 50\%$ of the

tumour cell positive). u-PA expression was scored separately for proportion (1, $< 25\%$; 2, 25–49%; 3, 50–75% or 4, $> 75\%$) and intensity (0, none; 1, weak; 2, moderate or 3, strong). If the proportion was $< 25\%$ and the intensity was zero, expression was considered negative; all other combinations were considered positive. Eosinophilic and lymphocytic infiltration was scored semiquantitatively (1, no or few; 2, moderate or 3, marked) according to the density of stromal infiltrate at the advancing front of the tumour.

All specimens were scored by two independent investigators. In case of disagreement, agreement was obtained by revision of the specimen by both investigators by means of a double-headed microscope. Scoring of protein expression was made without knowledge of the pathological or clinical outcome of the patients.

For statistical analysis the SPSS program was used. The chi-square test (with Yates' correction) was used to evaluate differences in proportions. Step-wise logistic regression analysis was used to study the prognostic value of tumour marker combinations for lymph node metastasis. Survival curves are compared using the log-rank test. Multivariate survival analysis was performed using Cox's regression analysis. For the evaluation of overall survival and disease-free period, post-operative deaths were excluded. Differences are considered statistically significant when $P < 0.05$.

Results

The results are based on the data of 105 patients, 70 males and 35 females. Mean age of the patients was 64.9 years (s.d. 10.8) in men and 68.4 years (s.d. 11.9) in women (NS).

The association between positive protein expression and different tumour characteristics is reported in Tables II and III. Rb was the only marker with a significant association with metastatic/unresectable tumours ($P=0.006$). *Myc*, nm23, E-cadherin and u-PA expression were all associated with T stage ($P < 0.05$). No association was found between protein expression and TNM stage or tumour location (not shown). There was also a statistically significant association between intent of surgery (curative or palliative) and Rb and E-cadherin expression ($P < 0.001$ and $P = 0.03$).

Step-wise logistic regression analysis was used to investigate whether a combination of several markers has a stronger relation with the tumour characteristics than each marker separately. Also, in multivariate analysis, positive Rb expression was the only important predictor (RR 3.66; 95% CI 1.50–8.95; $P=0.004$) for metastatic or unresectable tumours. For the risk of the operation being palliative, Rb (RR 6.47, 95% CI 2.29–18.32, $P=0.0003$) and *myc* (RR 0.24, 95% CI 0.08–0.69, $P=0.007$) were selected as the combination with additional independent prognostic value.

Table I Antibodies with the dilutions used, staining pattern, number of specimen examined and percentage of positive expression in primary adenocarcinomas of the stomach

Markers	Dilution	Staining pattern	Specimen	
			Examined n	Positive (%)
Growth regulation				
p53	1:100	Nuclear	101	39
Rb	1:500	Nuclear	105	53
<i>myc</i>	1:200	Nuclear/ cytoplasmic	105	41
Adhesion				
nm23	1:100	Cytoplasmic	105	50
Ep-CAM	1:100	Membranous	105	58
E-Cadherin	1:1000	Membranous	104	66
CD44v5	1:20 000	Membranous	93	28
CD44v6	1:5000	Membranous	76	90
Plasminogen activation				
u-PA	1:100	Cytoplasmic	84	39

Table IV summarises the results of the comparison (univariate analysis) of marker expression and the amount of eosinophilic and lymphocytic infiltration with overall survival and DFP after curative resection. There was an association between the amount of eosinophilic and lymphocytic infiltration, Rb and *myc* expression and overall survival, whereas only the presence of moderate-marked lymphocytic infiltration was associated with DFP. For overall survival and DFP, multivariate analysis was performed using the step-wise Cox's regression analysis. If only the markers that could be determined preoperatively were used, no other marker had an independent additional value next to Rb expression for survival. For both overall survival and DFP, TNM stage and the amount of lymphocytic infiltrate were selected as the combination with additional independent prognostic value (Table V).

Discussion

Preoperative staging of gastric cancer is suboptimal. While the best prognostic information is provided by the TNM classification, which also determines curative resectability, it

can be obtained post-operatively only. Although at present more accurate staging is possible by means of laparoscopy and cytological examination of abdominal washings, it is not routinely used (Bonenkamp *et al.*, 1996). Therefore, there is still a need for additional, reliable prognostic factors, preferably obtained by means of minimally invasive techniques, such as biopsy specimen. Better preoperative staging or other prognostic factors would be helpful in selecting patients suitable for surgery and preventing laparotomy with associated morbidity and mortality in patients with unresectable (incurable) tumours and would enable us to predict tumour behaviour and patient prognosis.

The presence of moderate to marked eosinophilic infiltration was found in gastric (Iwasaki *et al.*, 1986; Yu *et al.*, 1995) and human colonic cancer (Pretlow *et al.*, 1983) to be associated with prolonged survival. The same association was found for marked lymphocytic infiltration in gastric and rectal cancer (Yu *et al.*, 1995; Jass, 1986). The study of Yu *et al.* (1995) shows clearly that a standardised, quantitative analysis is needed, since there was high interobserver variation. Moreover, because these parameters can only be assessed in resected tumour specimens, they are not useful for preoperative staging or the prediction of resectability. In the present study, only the amount of eosinophilic infiltration was significantly associated with metastatic/unresectable tumours (Table II) and actual surgical procedures (Table III).

We have compared the expression of several proteins, associated with different steps of carcinogenesis and development of metastasis, with lymph node status, T stage, TNM stage and with curative resectability. In univariate analysis, a clear association was found between Rb expression and nodal involvement. Also, in multivariate step-wise logistic regression analysis, positive Rb expression was the only significant prognostic factor for nodal involvement out of all the molecular markers (RR 3.66). Alternatively, the risk of the operation becoming palliative was predicted with an overall accuracy of 77.0% by the combination of Rb and *myc*. The Rb gene is a tumour-suppressor gene, which is frequently mutated in a wide range of human cancer types (Bartek *et al.*, 1992; Harbour *et al.*, 1988; Cryns *et al.*, 1994). Although until now nothing has been known about altered Rb expression in gastric cancer, decreased expression is reported to be associated with invasion in bladder carcinoma (Cordon-Cardo *et al.*, 1992), whereas it had no prognostic value in breast carcinoma (Pietiläinen *et al.*, 1995). Since Rb is a tumour-suppressor gene, mutations leading to decrease or loss of Rb expression are expected to be associated with an unfavourable prognosis. However, not all Rb mutations result in the loss of immunohistochemically detectable expression. Instead, mutations that do not abolish the production of the Rb protein may lead to the production of a not functional, nuclear Rb protein (Mittnacht *et al.*, 1991). It has been demonstrated (Xu *et al.*,

Table II Univariate analysis of the association between lymph node metastasis and positive marker expression (n=105) and lymphocytic and eosinophilic infiltrates (n=81) in primary adenocarcinomas of the stomach

Markers	Node negative		Node positive/ irresectable		P-value
	n	Positive (%)	n	Positive (%)	
p53	34	12 (35)	67	27 (40)	0.79
Rb	34	11 (32)	71	45 (63)	0.006
<i>myc</i>	34	16 (47)	71	27 (38)	0.50
nm23	34	17 (50)	71	35 (49)	1.00
Ep-CAM	34	19 (56)	71	42 (59)	0.92
E-cad	34	19 (56)	70	50 (71)	0.18
CD44v5	33	7 (21)	60	19 (32)	0.40
CD44v6	28	24 (86)	48	44 (92)	0.67
u-PA	27	8 (30)	57	25 (44)	0.31
Infiltrates					
Lymphocytic	32		49		0.14
No or few	7 (22)		16 (33)		
Moderate	12 (38)		23 (47)		
Marked	13 (41)		10 (20)		
Eosinophilic	32		49		0.03
No or few	15 (47)		37 (76)		
Moderate	8 (25)		6 (12)		
Marked	9 (28)		6 (12)		

Table III Univariate analysis of the association between tumour characteristics and positive marker expression and the amount of infiltrates in primary adenocarcinomas of the stomach

Tumour characteristics	Total n	p53 (%)	Rb (%)	myc (%)	Markers (n=105)						Total n	Infiltrates (n=81) ^a			
					nm23 (%)	Ep-CAM (%)	E-Cad (%)	VFF8 (%)	VFF18 (%)	u-PA (%)		Lymphocytic 1-2(%)	Lymphocytic 3(%)	Eosinophilic 1(%)	Eosinophilic 2-3(%)
T stage^b															
T1	17	38	53	41	24	53	53	18	85	15	16	56	44	44	56
T2	51	41	41	49	51	59	66	28	90	45	44	70	30	66	34
T3	17	35	65	47	71	77	53	33	94	21	17	82	18	71	29
T4	16	27	75	6	44	38	100	25	88	71	3	100	0	100	0
TNM stage^c															
I	38	34	40	45	45	50	61	28	87	33	34	59	41	50	50
II	18	47	50	44	50	72	53	25	100	40	16	94	6	69	31
III	16	19	50	56	56	75	63	15	92	42	16	62	38	69	31
IV	28	52	71	29	57	50	79	33	83	46	14	76	14	86	14
Operation															
Curative	69	34	41	48	51	62	59	24	89	36	64	67	33	58	42
Palliative	36	48	78	28	47	50	81	37	90	45	17	88	12	88	12

^aResection specimens were available for 81 cases. Missing cases ^b(n=4), ^c(n=5) are left out of the analysis. (%), row percentages; bold, statistically significant difference (P<0.05); Lymphocytic and eosinophilic infiltration 1, no or few; 2, moderate; 3, marked.

Table IV Univariate analysis of the association between marker expression and median survival (post-operative deaths excluded, *n* = 13) and disease-free period (DFP) after curative resection (post-operative deaths excluded)

Infiltrates		Overall survival			P-value	DFP-(<i>n</i> = 54)			P-value
		1	Amount of infiltrate (<i>n</i> = 68)			1	Amount of infiltrate (<i>n</i> = 54)		
			2	3			2	3	
Eosinophils	(<i>n</i>)	44	12	12	0.02	32	12	10	0.17
Median	days	782	1530 +	1530 +		1350 +	1530 +	1530 +	
Lymphocytes	(<i>n</i>)	20	29	19	0.02	15	22	17	0.01
Median	days	763	1350 +	1530 +		1350 +	1350 +	1530 +	

Markers		Marker expression (<i>n</i> = 92)			P-value	Marker expression (<i>n</i> = 59)			P-value
		Negative	Positive			Negative	Positive		
p53	(<i>n</i>)	55	34		39	19			
Median	(days)	835	764	0.70	1350 +	1530 +	0.09		
Rb	(<i>n</i>)	43	49		35	24			
Median	(days)	1350 +	465	0.02	1530 +	1050 +	0.07		
<i>myc</i>	(<i>n</i>)	55	37		30	29			
Median	(days)	477	1050 +	0.04	1530 +	1530 +	0.74		
nm23	(<i>n</i>)	47	45		29	30			
Median	(days)	825	698	0.93	1530 +	1350 +	0.82		
Ep-CAM	(<i>n</i>)	40	52		22	37			
Median	(days)	720	806	0.20	1530 +	1530 +	0.32		
E-cad	(<i>n</i>)	29	62		23	35			
Median	(days)	1100	674	0.17	1350 +	1530 +	0.88		
VVF8	(<i>n</i>)	59	22		43	11			
Median	(days)	900 +	540	0.27	1530 +	1350 +	0.72		
VFF18	(<i>n</i>)	7	60		6	43			
Median	(days)	1300 +	804	0.51	1530 +	1350 +	0.72		
u-PA	(<i>n</i>)	45	28		29	16			
Median	(days)	896	495	0.13	1530 +	1530 +	0.76		

Eosinophils and lymphocytes 1, no or few; 2, moderate 3, marked.

Table V Results of stepwise Cox's regression analysis on survival (*n* = 64, post-operative deaths excluded) and on disease-free period (*n* = 53, curative resections and post-operative deaths excluded)

		Selected parameters		n	RR	95% CI	P-value
Survival	Dead	TNM stage	I + II	42	1.00		
	Alive		III + IV	22	5.24	2.49 - 11.02	<0.001
			Lymphocytes No or moderate	45	1.00		
			Marked	19	0.21	0.07 - 0.63	0.004
Disease-free period	Relapse free	TNM stage	I + II	40	1.00		
	Relapse		III + IV	13	5.62	2.13 - 14.85	<0.001
			Lymphocytes No or moderate	36	1.00		
			Marked	17	0.12	0.03 - 0.56	0.006

Included parameters are TNM stage, eosinophils, lymphocytes, p53, Rb, *myc*, nm23, Ep-CAM and E-cadherin

1991) that the staining intensity is correlated with the cell cycle (higher in active cells). In our study, we have not analysed whether expression is related to mutations in the Rb gene, but it is conceivable that up-regulation, amplification or even longer half-life of the mutant Rb protein may result in immunohistochemically detectable overexpression of the Rb protein. Expression of *myc* is related to the cell cycle and determines either continuous proliferation or apoptosis (Wyllie, 1993). *Myc* has a high turnover rate, which makes its place as a suitable prognostic marker for routine purposes questionable. This explains the contradictory findings that were reported in studies of *myc* expression (Ninomiya *et al.*, 1991; Auguste *et al.*, 1992; Borg *et al.*, 1992; Fox *et al.*, 1993). Furthermore, the interpretation of the staining pattern is highly variable. For instance, in the study of Ninomiya *et al.* (1991), it is not clear what criteria were used to call a tumour positive. In our study, the expression of CD44 splice variants v5 and v6 was contradictory with an earlier report (Heider *et al.*, 1993), for unknown reasons, whereas the findings with u-PA were similar to earlier reports (Nekarda *et al.*, 1994).

In univariate analysis, we found that moderate and marked eosinophilic infiltration, marked lymphocytic infiltration, Rb and *myc* expression had a prognostic value for

overall survival. Moderate eosinophilic infiltration was just as favourable as marked eosinophilic infiltration, whereas only marked lymphocytic infiltration had a strong association with longer survival. Since there is a correlation between the presence of lymphocytic and eosinophilic infiltrates, these two parameters had no additional value to each other. In multivariate step-wise Cox's regression analysis for both overall survival and DFP, the best prognostic information was provided by TNM stage and lymphocytic infiltrate (Table V). TNM stage is the most important prognostic factor for survival and also for DFP. The presence of marked lymphocytic infiltrate was the only parameter with an additional prognostic value to the TNM stage, while none of the markers had an additional prognostic value.

In conclusion, a prognostic significance of the tumour biological marker Rb was demonstrated in a large series of gastric cancer patients. Adding also post-operatively obtained information, lymphocytic and eosinophilic infiltrates proved to be important prognostic markers for surgical procedure in patients with gastric cancer. The presence and amount of these infiltrates should, therefore, be added to other post-operatively determinable prognostic factors in gastric cancer. To establish reproducibility and the place of tumour markers

as clinical prognostic factors in patients with gastric cancer, we have started further investigations in the tissue specimens of the remaining patients entered in the DGCT.

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