RESEARCH ARTICLE

Neutrophil gelatinase-associated lipocalin (NGAL) and matrix metalloproteinase-9 (MMP-9) prognostic value in lung adenocarcinoma

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Abstract Prognosis in patients with lung cancer is poor. Neutrophil gelatinase-associated lipocalin (NGAL) and matrix metalloproteinase-9 (MMP-9) are proteins involved in the invasion and metastases of cancer. The objective of this study is to determine if there is a relationship between tumor expression of NGAL and MMP-9 in lung adenocarcinoma patients with prognosis and overall survival. Retrospective analysis was made of patients with lung adenocarcinoma treated at Medica Sur Hospital between 2005 and 2013. Tumor tissue was analyzed for NGAL and MMP-9 expression by immunohistochemistry. We identified 41 patients. Mean overexpression in tumoral tissue of NGAL was 70 % and 30 % for MMP-9. Univariate analysis revealed that prognostic factors associated with overall survival (OS) were NGAL expression and stage at diagnosis. Median OS for NGAL expression <70 % was 45.7 months (95 % CI; 15.2-76.2) and for patients with \geq 70 % 4.6 months (95 % CI; 0.5–18.8; *P*<0.0001), and for stage at diagnosis (stages I and II mean not reached), stage III mean OS 15.57 months (95 % CI; 9.8-21.2) and stage IV 9.6 months (95 % CI; 0.8-18.4. P=0.002). No differences in OS were found for expression of MMP-9. Multivariate analysis revealed significance for OS in NGAL expression (HR 5.01 [95 % CI; 1.68–14.93] P=0.004) and stage at diagnosis

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D. Motola-Kuba Gastroenterology Clinic, Medica Sur Hospital, Mexico City, Mexico (HR 2.05 [95 % CI 1.30–3.22] P=0.002). Tumoral tissue expression of NGAL \geq 70 % confers a worse prognosis compared to those who did not. NGAL is an independent prognostic factor of stage at diagnosis.

 $\label{eq:constraint} \begin{array}{l} \textbf{Keywords} \ \ Lipocalin \cdot Lepidic \ \cdot \ Adenocarcinoma \ \cdot \ Matrix \\ metalloproteinase \end{array}$

Abbreviations

NGAL Neutrophil gelatinase-associated lipocalin MMP-9 Matrix metalloproteinase–9

Introduction

Lung cancer is an aggressive disease. In 2012, it was estimated that 1,825,000 patients around the world were diagnosed with lung carcinoma, and mortality rates were as high as 1, 590,000 deaths that year. In México, in that same year it had an incidence of 8,439 new cases (5,471 for men and 2,968 for women) and 7,608 casualties in both sexes representing the first cause of death from all malignancies [1]. As today, the most important factor for the prognosis of patients is the stage at diagnosis, with a worse outcome for those who present with a more advanced disease [2].

Despite new treatments, survival has been low even in the earliest stages (IA with a median overall survival of 60 months) and even lower in advanced stages (median survival of 6 months in stage IV) [2]. Due the impact of AJCC staging system for patient staging, it is also used to decide the most appropriate treatment. Generally, stages I and II benefit from local treatment with surgical resection and in some specific

cases the use of adjuvant chemotherapy. Stage III patients need more aggressive treatment, even combining three modes of therapeutics: surgery, chemotherapy, and/or radiotherapy. Stage IV benefits from the palliative use of chemotherapy or molecular targeted therapy [3]. This staging system and its use for treatment only reflect the anatomical extent of the disease without a correlation with progression and/or overall survival of patients. Even in early stages, the prognosis can be more aggressive than expected and reflected with a rapid progression of regional disease or distant metastases [4], hence the importance for identifying the subset of individuals that are within this risk group.

Neutrophil gelatinase-associated lipocalin (NGAL) is a 25kDa protein which is stored in the granules of human neutrophils. Overexpression of NGAL in human tumors and its impact in the prognosis and clinical outcomes has led to contradictory results. It has been described a worst stage at diagnosis and a short progression-free and overall survival in breast, esophagus, stomach, pancreas, and colon cancer [5–7] and a better prognosis in anaplastic thyroid and ovarian cancer [8].

One of the possible explanations of the negative prognosis of patients with malignant tumors that overexpress NGAL is the ability of this protein to positively modulate the activity of the matrix metalloproteinase-9 (MMP-9) enzyme. MMP-9 degrades the basement membranes and extracellular matrix, thus enabling angiogenesis, invasion, and metastasis of malignant cells [9]. There is a lack of prognostic information and clinical outcomes of NGAL and MMP-9 overexpression in lung adenocarcinoma. In this study, we investigated the NGAL and MMP-9 prognostic value in lung adenocarcinoma.

Material and methods

Patients

This retrospective study was approved by the institutional review board of our hospital. Tumoral tissue samples were obtained from patients with a diagnosis of non-small cell lung carcinoma (NSCLC) (stages I to IV according to TNM Staging System, 7th edition), with histological confirmation of adenocarcinoma, who attended the outpatient Oncology Clinic from Médica Sur Hospital from year 2005 to 2013. We excluded patients with non-adenocarcinoma histology and with previous chemotherapy or radiation therapy. Clinical data of all patients were obtained retrospectively, and it included clinical history, histopathology diagnosis, and smoking history. Never smoker was defined as having a lifetime exposure of <100 cigarettes [10]; the tobacco smoking index was calculated by multiplying the number of cigarette packs consumed per day by the number of years spent smoking [11].

Tissue procurement and expression of NGAL or MMP-9 by immunohistochemistry

Selecting the procedure for obtaining a biopsy from all patients depended on patient and tumor characteristics. Tumor specimens were collected at the time of diagnosis using computed tomography (CT)-guided tru-cut, lobectomy, or by bronchoscopy and were analyzed by the pathology department for their histologic diagnosis, who were blinded to the clinical outcomes. Immunostains were performed using the standard avidin-biotin-peroxidase method. Samples were later embedded in paraffin until processing. Antibodies were used with appropriate controls (Lipocalin-2 antibodies [NGAL-GTX63306] Clone: EPR5084 Brand: GeneTex® Dilution 1:100. Anti-metalloproteinase-9 [MMP-9-GTX100458] Clone: NC1 Brand: GeneTex® Dilution 1:200) then permanently mounted in resin for microscopy, and coverslips were placed for evaluation by an expert pathologist. To register the positivity of the immunoreactions, a qualitative and quantitative system determined by direct light microscopy observation was used. Tissue used as control to evaluate the positivity of NGAL-2 was the intense staining (3+) in the cytoplasm of the glandular cells of the endocervix. The tissue used as control to evaluate the reactivity of MMP-9 was a strongly positive reaction in alveolar macrophage cytoplasm (3+). A scale of 1+ (for weak staining) and for up to a 3+ (intense staining) for both antibodies in the cytoplasm of tumor cells was used. Likewise, in each case, the percentage of neoplastic cells that showed weak to strong positive staining was determined. Immunohistochemical staining results were delivered to researchers in a single-blinded way by the pathology department.

Regimen of treatment

Patients were treated according to the international guidelines for the treatment of lung cancer [3]. If they were suitable for adjuvant or palliative chemotherapy (CT), all patients received platinum-based CT as the first line of treatment. CT regimens included paclitaxel plus cisplatin or carboplatin and pemetrexed plus cisplatin or carboplatin. Second-line treatment choice was determined by the oncologist's criteria.

Outcome measurement

All patients underwent CT or positron emission tomography (PET-CT) scans and were evaluated according to RECIST 1.1 at the baseline (before each line of treatment) and afterwards, every 2 to 3 cycles of the treatment in case of palliative treatment. Progression-free survival (PFS) was defined as the time from the start of the treatment until disease progression or

last visit. Overall survival (OS) was defined as the time from histologic diagnosis until death or last follow-up visit.

Statistical analysis

Continuous variables were summarized as arithmetic means, medians, and standard deviations (SDs) for descriptive purposes, and categorical variables consisted of percentages and their respective 95 % confidence intervals (95 % CI). Inferential comparisons were made using the Student's t test or the Mann-Whitney U test, according to the data distribution (normal or abnormal) determined by the Kolmogorov-Smirnov test. The χ^2 test or Fisher's exact test were used for assessing the statistical significance of categorical variables, determined as being P < 0.05 when using a two-tailed test. Statistically significant and borderline significant variables (P < 0.1) were included in a multivariate logistic regression analysis. OS was analyzed with the Kaplan-Meier method, whereas comparisons among the subgroups were analyzed using the log rank or Breslow tests, if the two survival curves crossed. For survival curve analysis, all the variables were dichotomized (for age, we utilized the median). Adjustment for potential confounders was performed by using a multivariate Cox regression model, and hazard ratios (HR) were calculated along with their corresponding 95 % CIs as a measure of association. Statistical significance was determined as P < 0.05 using a two-tailed test. SPSS software version 19 (IBM) was used for all statistical analyses.

Results

Patient population

It was possible to identify a total of 41 patients who met the inclusion criteria within the time frame 2005–2013. The median age was 61.4 ± 10.1 years, 53.7 % were women and 46.3 % men; 75.6 % of patients had a smoking history.

The most common types of invasive adenocarcinoma were as follows: the predominant type was the mixed type (70.7 %), followed by lepidic (14.6 %). The grade of differentiation was moderately differentiated in the majority of patients (53.7 %), followed by poorly differentiated (39.1 %) and well differentiated (7.3 %). Vascular permeation was present in 68.3 % and perineural invasion in 9.8 % of all patients.

The stages at diagnosis were stage I (19.5 %), II (12.2 %), III (19.5 %), and IV (48.8 %). All patients stage II through IV received platinum-based cytotoxic chemotherapy treatment (adjuvant or palliative). The median follow-up of all patients was 25.6 ± 32.2 months. For most patients, the EGFR mutation status was unknown (n=36, 87.80 %). Patient general characteristics are listed in Table 1.

Table 1 Basal characteristics of the population

Characteristic	Total	Percentage	
Gender			
Female	22	53.7	
Male	19	46.3	
Age			
Mean	61.41±10.10		
Range	39 - 80 years		
Smoking status			
No	10	24.4	
Yes	31	75.6	
Tobacco index (mean)	28.4±31	.2	
Range	0-100		
Hypertension			
No	27	65.9	
Yes	14	34.1	
Diabetic			
No	39	95.1	
Yes	2	4.9	
Adenocarcinoma type			
Lepidic	6	14.6	
Papillary	1	2.4	
Solid	1	2.4	
Others	4	9.8	
Mixed	29	70.7	
Grade			
Well differentiated	3	7.3	
Moderately differentiated	22	53.7	
Poorly differentiated	16	39.1	
Vascular permeation			
Yes	28	68.3	
No	13	31.7	
Perineural invasion			
Yes	4	98	
No	37	90.2	
Stage at diagnosis	5,	, oi z	
I	8	19.5	
I	5	12.2	
Ш	8	19.5	
IV	20	48.8	
Carcinoembryonic antigen (CEA) mean	20 74+2	93	
Rango	20.77 ± 27.5		
Follow-up months (median)	25 65+32 20		
ronow up, months (moduli)	0_156.0	2.20 a	
EGER mutation status	0 100.0	,	
Ves	3	7 31	
No	2	4.87	
Unknown	- 36	87.80	
Chemotherany regimen	50	07.00	
Paclitaxel + carbonlatin	18	60	
Pemetrexed + carbonlatin	5	16.6	
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Table 1 (continued)

Characteristic	Total	Percentage
Gemcitabine + cisplatin	2	6.6
Pemetrexed + cisplatin	2	6.6
Docetaxel + cisplatin	1	3.3
Vinorelbine + cisplatin	1	3.3
Gemcitabine + carboplatin	1	3.3

Expression and intensity of NGAL and MMP-9 by immunohistochemistry

Of the 41 patients, 39 expressed NGAL in tumor tissue (95.12 %) (Fig. 1) and 31 patients (75.60 %) expressed MMP-9 (Fig. 2). All patients negative for NGAL immunohistochemistry were also negative for MMP-9. Most patients had a stain intensity of 3+ for NGAL (32 patients, 78.04 %) and 2+ for MMP-9 (15 patients, 35.68 %). Mean NGAL expression was 70 and 30 % for MMP-9. (Table 2) We found a relationship between the percentage expression of NGAL related to percentage of expression of MMP-9. For patients with NGAL <70 % (n=19), the mean expression of MMP-9 was 22.3 %±27. For those with NGAL expression \geq 70 % (n=22), the mean expression of MMP-9 was 52.1 %± 39 (P=0.007, R=0.48) (Fig. 3).

Progression-free survival

The median PFS for stages I–II was not reached. Median PFS for stages III–IV was 10.8 months (5.9–15.7). In univariate and multivariate analyses, no variables directly influenced PFS. A suggestive trend for better PFS with NGAL expression <70 % was 13.1 months (95 % CI 8.3–18) and for patients with NGAL expression \geq 70 % was 7.8 months



Fig. 1 NGAL expression in pulmonary adenocarcinoma. 3+ Immunostain, 95 % for NGAL in lung adenocarcinoma (40×)



Fig. 2 MMP-9 expression in pulmonary a denocarcinoma. 3+ Immunostain, 95 % for MMP-9 in lung a denocarcinoma (40×)

(95 % CI, 5.3–10.2, P=0.062). For MMP-9, non-significant difference in PFS was shown: MMP-9 expression <30 % was 13.5 months (95 % CI 12.4–14.7) and for those with expression \geq 30 % was 7.8 months (95 % CI, 1.7–13.9) (Table 3).

Response to chemotherapy

Only 30 patients were evaluated for RECIST with an objective response of 40 % (CR + PR 12 patients). There were no statistically significant differences regarding the objective responses in patients treated with CT related to sex,

Table 2	Immunohistochemistry	results for	r NGAL	and MMP-9
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Characteristic	Total (<i>n</i> =41)	Percentage
NGAL expression		
Yes	39	95.12
No	2	4.82
NGAL expression %, mean	70 %	
NGAL intensity		
0	2	4.87
1+	1	2.43
2+	6	14.63
3+	32	78.04
MMP-9 expression		
Yes	31	75.60
No	10	24.40
MMP-9 expression %, mean	30 %	
MMP-9, intensity		
0	10	24.3
1+	7	17.07
2+	15	35.68
3+	9	21.95

Fig. 3 Association between NGAL and MMP-9 expression in tumoral tissue



differentiation, vascular permeation, perineural invasion, and percentage of expression of MMP-9. We found a suggestive trend for worse response to chemotherapy with NGAL expression \geq 70 % (no overall response in 11/15 (73.3 %) *P*= 0.065) (Table 4). The most common schemes used were paclitaxel and carboplatin (CBP) (55.4 %) and Pemetrexed + CBP (15.2 %) (Table 1).

Overall survival

The median OS was 17.0 months (range 4.7–29.2 months). In univariate analysis, only two variables directly influenced OS, stage at diagnosis and NGAL expression. Stages I and II (median OS not reached), stage III (median 15.57 months [95 % CI 9.8–21.2]), and stage IV (median 9.6 months [95 % 0.8–18.4 months], P=0.002). OS with NGAL expression <70 % was 45.7 months (95 % CI 15.2–76.2) and for patients with NGAL expression \geq 70 % was 4.6 months (95 % CI, 0.5–18.8; P<0.0001) (Fig. 4). We found no relationship between OS and NGAL stain intensity in tumoral cells.

Median OS for patients with MMP-9 expression <30 % was 36 months (95 % CI 1.92–70.1) and for those with expression ≥ 30 % was 15 months (CI 95 %, 7.3–22.7). No statistical difference between groups was found (*P*=0.568) (Fig. 5). No differences in OS were either found for MMP-9 stain intensity in tumoral cells, gender, smoking status, vascular permeation, or degree of differentiation.

In the multivariate analysis, stage at diagnosis (HR 2.05 [CI 95 %; 1.30–3.22] P<0.002) and NGAL expression \geq 70 % (HR 5.01 [CI 95 %; 1.68–14.93] P<0.004) were statistically significant for OS (Table 5).

Discussion

For lung cancer, there are conflicting data regarding the prognosis of histopathological variables (lymphovascular invasion, adenocarcinoma vs. epidermoid histology), with controversial results, and they are only described for resectable disease, with no information for metastatic patients [12–15]. Another variable with no definite results is the degree of tumor differentiation in patients with operable disease (worst prognosis in undifferentiated tumors) [14, 15]. Vascular invasion was described as a negative prognostic factor for patients with T1-2 N0 tumors. In a series of 746 patients, microscopic vascular permeation was identified in 257 individuals (34 %) and conferring them a worse 5-year OS compared to those who did not (65 vs. 55 %, respectively) [16], hence the importance of identifying new histological prognostic markers in order to use them in conjunction with the TNM system and allow a better selection of treatment for each patient.

NGAL has been known for many years for its role in the innate immune system. It also receives the names of lipocalin-2, siderocalin, uterocalin, and 24p3 [17]. With more than 50 members, it belongs to the lipocalin superfamily. They all have similar three-dimensional structure in a single eight-stranded anti-parallel b-barrel surrounding a central pocket [18]. NGAL has an ability to bind and transport small hydrophobic substances such as fatty acids, prostaglandins, arachidonic acid, retinoids, and hormones [19].

NGAL's principal mechanism of action is to capture the extracellular iron particles (also known as siderophores) and transport them to the inner cell after interacting with the specific membrane receptor 24p3R. This causes an increase in the iron cytoplasmic levels [20]. By activating an iron

Table 3Progression-freesurvival (PFS) in patients withlung adenocarcinoma

PFS	Mean (months) (95 % CI)	Univariate analysis <i>P</i>	Multivariate analysis		
			HR	95 % CI	Р
Gender					
Female Male	13.5 (9.6–17.4) 7.8 (5.3–10.2)	0.092	2.29	0.78–6.76	0.13
Age					
≤60 >60	10.1 (2.7–17.5) 11.4 (4.6–18.1)	0.942	0.97	0.93-1.02	0.33
Smoking status					
Yes No	8.1 (3.6–12.8) 15.2 (11.6–18.9)	0.056	1.42	0.39–5.17	0.58
Vascular permeation					
Yes No	10.8 (4.3–17.4) 8.1 (0–18.8)	0.698	1.88	0.58-6.1	0.28
Grade					
Moderately differentiated	13.1 (7.9–18.2)	0.061	1.83	0.49-6.80	0.36
Poorly differentiated	7.8 (4.3–11.2)				
NGAL %, expression					
<70 % ≥70 %	13.1 (8.3–18) 7.8 (5.3–10.2)	0.062	2.30	0.60-8.85	0.21
MMP-9 %, expression					
<30 % ≥30 %	13.5 (12.4–14.7) 7.8 (1.7–13.9)	0.1	1.19	0.38–3.69	0.75

CI confidence interval, HR hazard ratio

depletion strategy, this protein participates in the irondependent enzymatic defensive systems [21]. Through the

Table 4Overall response

Variable	Overall response (Overall response (OR)		
	Yes	No		
Gender				
Female Male	7/17 (41.2 %) 6/13 (46.2 %)	10/17 (58.8 %) 7/13 (53.8 %)	0.785	
Differentiation				
Well Moderately	0/2 (0 %) 9/7 (47.1 %)	2/2 (100 %) 9/17 (52.9 %)	0.439	
Poor	5/11 (45.5 %)	6/11 (54.5 %)		
Vascular permeati	ion			
No Yes	3/9 (33 %) 10/21 (47.6 %)	6/9 (66.7 %) 11/21 (52.4 %)	0.469	
Perineural invasio	m	· · · · ·		
No Yes	13/29 (44.8 %) 0/1 (0 %)	16/29 (55.2 %) 1/1 (100 %)	1.00	
NGAL				
<70 % ≥70 %	9/15 (60.0 %) 4/15 (26.7 %)	6/12 (40 %) 11/15 (73.3 %)	0.065	
MMP-9				
<30 % ≥30 %	6/15 (40 %) 7/15 (46.7 %)	9/15 (60 %) 8/15 (53.3 %)	0.713	

regulation of iron-responsive genes, which are important in the differentiation of primordial cells, NGAL seems to participate in the proliferation, differentiation, and development of human tumors, by favoring iron uptake from extracellular space within the malignant cells [9]. Various in vitro experimental models demonstrated that factors known to promote proliferation of malignant cells such as hepatocyte growth factor, the neu-transforming factor, the phorbol ester PMA, retinoic acid, poliomavirus infection, and even glucocorticoids can induce the synthesis and release of NGAL. This has suggested that its overexpression in tumoral tissue is a negative prognostic factor and associated with shorter overall and disease-free survival in several neoplasms [22, 23].

NGAL was originally identified as a protein covalently associated with 92-kDa gelatinase/MMP-9 from humanactivated neutrophils (for hence its name, being the MMP-9 a gelatinase). Indeed, by forming the NGAL/MMP-9 dimeric complex, NGAL can protect MMP-9 from its auto and proteolytic degradation and consequently results in a higher gelatinolytic action of MMP-9 on extracellular matrix [24]. This would trigger an enhancement of the enzymatic activity of MMP-9 and explain the tumoral invasiveness and diffusion associated with NGAL overexpression [25, 26]. The overexpression in tumoral tissue of NGAL/MMP-9 complex as a worse prognostic factor has been identified in bladder, brain, breast, esophageal, gastric, and hematological malignancies [24].

Fig. 4 Mean OS in patients according to NGAL percentage of expression in tumoral tissue



In our study, the expression of NGAL was prognostic for overall survival in all patients and not only limited to those with operable disease. We are aware that one limitation of our study is the vast heterogeneity of the population and that we included local, locally advanced, and metastatic disease. Most of the patients had advanced disease at diagnosis: stage III (n= 8; 19.5 %) and stage IV (n=20; 48.8 %), and tissue for those patients was obtained from core needle biopsies. On the other hand, stages I and II were present in only 13 patients (31.7 %), and tissue was obtained from therapeutic lobectomies performed at our institution that influenced in the prognosis of the local disease. However, in the multivariate analysis, a ≥ 70 % expression of NGAL in tumoral tissue correlated with

a less favorable prognosis, and it was independent of the stage at diagnosis. To our knowledge, only one study has evaluated the role of NGAL expression as a marker of resistance to EGFR-TKIs in cell lines of lung cancer [27], but there are no reports that have assessed the prognostic role of NGAL in patients with NSCLC and limited to the subset of lung adenocarcinoma and neither with the expression of both NGAL and MMP-9.

The prognostic value of NGAL, independently of MMP-9 expression, has been previously reported in patients with stage I colon adenocarcinoma [7]. NGAL expression and its poor prognosis have been evaluated in breast, esophagus, and stomach cancer [5–7, 22, 28]. There is inconclusive or





Table 5 OS analysis in patientswith lung adenocarcinoma

		Univariate analysis	Multivariate analysis		
OS	Mean (months) (95 % CI)	Р	HR	95 % CI	Р
Gender					
Female Male	36.76 (2.5–70.9) 12.38 (3.5–21.2)	0.47	0.73	0.32–1.66	0.45
Age					
≤60 >60	17.4 (7.4–26.6) 21.1 (6.6–35.4)	0.846	0.99	0.95–1.03	0.71
Smoking status					
Yes No	15.5 (8.1–23.0) 36.7 (8.8–64.7)	0.69	1.20	0.42-3.43	0.72
Vascular permeation					
Yes No	17.4 (3.8–31.0) 22.7 (6.–39.2)	0.68	1.14	0.47–2.73	0.76
Grade					
Well differentiated	45.7 (0.4–91.0)	0.78	0.6	0.22-1.58	0.30
Moderately differentiated	22.7 (11.0-34.9)				
Poorly differentiated	11.3 (9.6–12.9)				
Stage at diagnosis					
I and II III	NA 15.57 (9.8-21-2)	0.002	2.05	1.30-3.22	0.002
IV	9.6 (0.8–18.4)				
NGAL %, expression					
<70 % ≥70 %	45.7 (15.2–76.2) 4.6 (0.5–18.8)	<0.0001	5.01	1.68–14.93	0.004
MMP-9 %, expression					
<30 % ≥30 %	36.0 (1.92–70.1) 15.0 (7.3–22.7)	0.568	0.72	0.28–1.84	0.49

CI confidence interval, *HR* hazaro ratio

conflicting data in patients with renal [29] and pancreatic carcinoma [30] and a less unfavorable prognosis in anaplastic thyroid and ovarian cancer [8]. Our results may suggest that identifying patients with a worse prognosis (overexpression of NGAL \geq 70 % in tumoral tissue) regardless of stage at diagnosis could help the physician selecting an aggressive treatment approach and a stricter follow-up. On the other hand, patients who do not could avoid overtreatment and have a loose surveillance.

Furthermore, it has been found that NGAL results as a physiological regulator of cell proliferation, and it is active in the process of growth and differentiation of tissues since the embryonic stage. In fact, it was found that binding to the 24p3R receptor, NGAL regulates cell proliferation by a mechanism of depletion and increase of the intracellular iron, which induces cell to apoptosis or survival, respectively [31]. In our series, we assume that NGAL overexpression of 95.12 % is secondary to an increased iron need for NSCLC cancerogenesis [32].

MMP-9 and its expression as an unfavorable prognostic factor have been extensively studied in NSCLC [33–41] besides being a possible therapeutic target in this group of patients [42–44]. The lack of correlation between MMP-9 and overall survival in our study might reflect the relative insensitivity of Northern blot analyses used in the previously mentioned studies that, compared to immunohistochemistry, cannot precisely discriminate between stromal and neoplastic cell protein expression.

NGAL also has been described in pre-clinical models as a novel mechanism of NSCLC resistance to the small molecule tyrosine-kinase inhibitor erlotinib. In a recent study of mice NSCLC cell lines, NGAL overexpression in erlotinibsensitive cells augmented apoptosis resistance. This was mediated by NGAL-dependent modulation of the pro-apoptotic protein Bcl-2-like protein 11 (BIM). This might contribute to the TKI resistance in some adenocarcinoma lung cancer patients by overcoming the apoptosis pathway in malignant cells [27]. In an Asian population study, BIM deletion polymorphism predicted shorter PFS to EGFR-TKIs and OS in advanced NSCLC [45].

Our results indicate that NGAL expression is associated with a worse OS prognosis compared to patients who did not, independently of stage at diagnosis. On the other hand, we found no prognostic significance for the expression of MMP-9 and OS or any correlation with the expression of NGAL. This suggests that NGAL may be independent of the regulation of MMP-9. This has not been documented in NSCLC, but observed in cell lines of colon adenocarcinoma. In this study, in contrast to the positive relationship between the two markers in colon cancer previously published [46], it was observed that the expression of NGAL promotes invasion and the metastatic potential of neoplastic cells by a mechanism that is iron dependent, thus decreasing the inter-cellular-mediated E-cadherin adhesion [47]. This suggests a novel therapeutic strategy with iron-chelating drugs for the inhibition of NGAL, used as anti-neoplastic drugs. By reducing iron stores and transporters such as NGAL protein, these drugs inhibit cell proliferation. This has already been investigated in some animal models with promising results [48, 49].

Conclusion

In this group of patients with lung adenocarcinoma, overexpression of NGAL in tumor tissue greater than or equal to 70 % confers a worse prognosis compared to those who did not, and it is independent of the stage at diagnosis. Expression of MMP-9 did not confer any significant prognostic value nor correlation of this marker with NGAL, suggesting an independent mechanism of NGAL in lung adenocarcinoma for invasion and metastases. The prognostic value of NGAL is currently being validated in a larger series of patients and in a multi-institutional form.

Conflicts of interest None

Ethical approval All procedures performed were in accordance with the ethical standards of the institutional committee. For this type of study, formal consent is not required.

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