

Overexpression of neutrophil gelatinase-associated lipocalin and its receptor in colorectal carcinoma: Significant correlation with cell differentiation and tumour invasion

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Abstract. Neutrophil gelatinase-associated lipocalin (NGAL), a member of the lipocalin family, is related to inflammation and tumour. Recently, a specific cell-surface receptor (24p3R/NGALR) for lipocalin 24p3 was reported. However, the characteristics of NGALR expression in colorectal carcinoma (CRC) are not known. The objectives of this study were to investigate the expression of NGAL and NGALR in CRC specimens, and determine any relationship between the expression of these proteins and tumour progression. In the present study, CRC specimens of 102 patients were obtained, and the expression of NGAL, NGALR, ferritin and Ki67 was analyzed in paraffin sections by immunohistochemistry. Statistical analyses of the data collected were performed with SPSS software. We found that the cytoplasmic staining of NGAL, NGALR and ferritin, as well as the nuclear staining of Ki67 were significantly up-regulated in CRC tissues compared with normal colorectal tissues. Expression of NGAL was related to the deeper invasion of CRC ($P=0.026$), while NGALR was significantly associated with a deeper invasion ($P=0.018$) and a high degree of Tumor, Node and Metastasis stages ($P=0.042$) in CRC. The NGAL/NGALR co-expression was associated with poor cellular differentiation ($P=0.004$). Positive correlations between NGAL and NGALR ($r=0.432$, $P<0.01$), NGAL and ferritin ($r=0.374$, $P<0.001$), NGALR and Ki67 ($r=0.228$, $P<0.05$), NGAL/NGALR co-expression and ferritin ($r=0.349$, $P<0.001$), as well as NGAL/NGALR co-expression and Ki67 ($r=0.205$,

$P<0.05$) were observed. However, the expression of NGAL or NGALR was not significantly associated with patient survival. These findings detected an elevated expression of NGAL and NGALR resulting in poor cellular differentiation and a deeper invasion of CRC. Thus, NGALR may be a novel target for the treatment of CRC.

Introduction

Colorectal carcinoma (CRC) is one of the most common malignancies worldwide. Each year in the United States and Canada, CRC is diagnosed in >160,000 people and ~65,000 succumb to the disease, accounting for at least 10% of all cancer deaths. The lifetime risk of developing CRC is 1 in 17, affecting men and women alike, with 90% of cases occurring after the age of 50 years. Of note is that the incidence of CRC has increased substantially in Asia during the past few decades (1).

Neutrophil gelatinase-associated lipocalin (NGAL) is a 25-kDa secreted protein of the lipocalin superfamily (2). An elevated NGAL expression is observed in human cancers such as breast, pancreatic, ovarian and cholesteatoma as well as esophageal squamous cell carcinoma (3-7). Our previous studies demonstrated that the over-expression of NGAL plays an important role in the malignant transformation of human immortalized esophageal epithelial cells and it is involved in the invasion of esophageal squamous cell carcinoma cells (8-10). Others have reported very high expression levels of NGAL in colonic epithelium in areas of inflammation, with a weak expression occasionally seen in some epithelial cells of normal colon (11). However, Lee *et al* found that the ectopic expression of NGAL in colon cancer cells had little effect on their growth and viability (12). Conflicting results have been reported from various laboratories. Consequently, the clinical importance of NGAL expression in CRC remains unsettled.

Lipocalins are characterized by multiple molecular recognition properties, including the ability to bind to cell surface receptors (13). Several studies found that NGAL binds iron and delivers the latter to cells (14-16). Devireddy *et al* isolated a specific cell-surface receptor (24p3R/NGALR) for lipocalin 24p3, a highly conserved murine homolog of NGAL, and demonstrated that the expression of this receptor conferred

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on cells the ability to take up iron or undergo apoptosis depending on the state of 24p3, independent of the cell type (17). However, the expression patterns and specific features of NGALR in CRC are still unknown. Therefore, this study aimed to investigate the expression of NGAL and NGALR in CRC specimens, and determine any relationship between the expression of these proteins and tumour progression.

Materials and methods

Cases and clinical parameters. This study was approved by the ethics committee of the Medical College of Shantou University. Written informed consent to use resected samples for research was obtained from patients undergoing surgery.

For this retrospective study, archival formalin-fixed, paraffin-embedded specimens from 102 primary CRC patients were obtained from the Pathology Department of the Medical College of Shantou University, collected between 1992 and 2006.

The patients were 53 males and 49 females (median age 57 years, range 20–81). All patients except one were deceased at the end of follow-up.

Immunohistochemical staining and scoring. Briefly, each tissue section was de-paraffinised, rehydrated and then incubated with fresh 3% hydrogen peroxide for 10 min. After PBS rinse, antigen retrieval from the tissue was carried out by autoclaving in 0.01 M citrate buffer (pH 6.0) at 120°C for 3 min. Two drops (100 µl) or enough to completely cover tissue of the primary antibody were applied to each section and incubated in a moist chamber for 30 min. After the PBS rinse, the tissue sections were incubated for 10 min at room temperature with HRP polymer conjugate. Subsequently, they were stained with 0.003% 3,3-diaminobenzidine tetrahydrochloride and 0.005% hydrogen peroxide in 0.05 M Tris•HCl (pH 7.2), counterstained with Mayer's hematoxylin, dehydrated and mounted.

Negative controls were prepared by substituting PBS for primary antibody. The polyclonal rabbit NGALR antibody was raised using a C-terminal NGALR peptide (CDHVPLLATPNAL) as the immunogen and then affinity-purified on a peptide-coupled Sepharose column (Beijing Biosynthesis Biotechnology Co., Ltd). This antibody has been selected for its ability to recognize human NGALR. Rat anti-human NGAL antibody (1:50 dilution, R&D Systems, USA), rabbit anti-human ferritin antibody (1:2000 dilution, Sigma, USA), mouse anti-Ki-67 nuclear antigen (1:100 dilution, Golden Bridge International) and NGALR (1:10 dilution) were used.

The SuperPicTure Polymer Detection and Liquid DAB Substrate kits (Zymed, Carlsbad, CA, USA) were used to carry out immunohistochemical staining. Positive samples were defined as those showing brown signals in the cell cytoplasm or nucleus. When >5% of cells in a given specimen were positively stained, it was defined as a positive case.

Statistical analysis

The association between NGAL or NGALR expression and clinicopathological features of patients was analysed by the

Chi-square or Fisher's exact probability test. The concordance of NGAL and NGALR, NGAL and ferritin, or NGALR and Ki67 expression was determined by the κ test. The significance of NGAL or NGALR expression levels to patient survival was examined using the Kaplan-Meier method and log-rank test. Statistical analyses were performed using SPSS for Windows (Version 13.0). The accepted level of significance was $P < 0.05$.

Results

Overexpression of NGAL is significantly associated with tumour invasion in CRC. Representative results of NGAL immunostaining are shown in Fig. 1. Cytoplasm positivity for NGAL staining was observed in tumour tissues (Fig. 1B), and a weak expression was observed in adjacent normal colon tissues (Fig. 1A). A total of 102 CRC cases were included in the final analysis. NGAL expression was significantly increased in CRC (29/102, 28.4%) compared with adjacent normal tissues (0/81, 0.0%) ($P < 0.001$). The percentage of NGAL positivity was significantly associated with CRC invasion ($P < 0.01$), and NGAL expression appeared to intensify with the development of the cancer invasion. No significant association was observed between NGAL expression and cell differentiation, although the P-value was borderline ($P = 0.057$). NGAL expression was not significantly associated with age, gender, cell differentiation, lymph node metastasis or Tumor, Node and Metastasis (TNM) stages ($P > 0.05$; Table I) in patients with CRC. No significant association between NGAL expression and patient survival was noted ($P = 0.290$) (Fig. 3).

Overexpression of NGALR is significantly associated with tumour invasion and TNM stages in CRC. NGALR-positive staining was observed in the cytoplasm of CRC tumour cells (Fig. 1D), whereas weak positive staining was observed in a restricted intracellular area close to the cell membrane in adjacent normal tissues (Fig. 1C). The number of CRC specimens staining positive for NGALR (37/102, 36.6%) was significantly higher than that for normal glandular organ (8/93, 8.6%) ($P < 0.01$). NGALR expression was strongly associated with deep cancer invasion and with later TNM stages ($P < 0.05$ and $P < 0.05$). No significant association was found between NGALR expression and gender, age or lymph node metastasis. A significant association was found between tumour TNM stages and patient survival ($P = 0.043$; Fig. 3). However, there was no significant association between NGALR expression levels and patient survival (Fig. 3).

NGAL and NGALR co-expression is correlated with deeper invasion and tumour progression in patients with CRC. A positive correlation between NGAL and NGALR protein expression was observed in the 102 CRC cases ($r = 0.432$; $P < 0.01$). We therefore analyzed the relationship between NGAL and NGALR expression in these cases and the clinicopathological factors. The cases in which NGAL and NGALR were both positive (NGAL/NGALR co-expression) were stratified into group I ($n = 19$, 18.6%), the positive group, while the rest of the cases were put into group II ($n = 83$, 71.4%), the negative group. Statistical analysis showed that the two groups exhibited differences in invasion ($P < 0.05$), cell differentiation

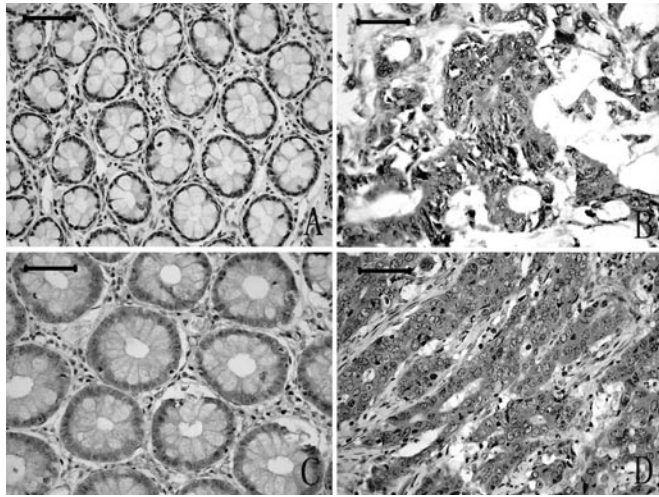


Figure 1. NGAL and NGALR protein expression in CRC (B and D) and adjacent normal glandular organ tissues (A and C). Scale bars, 50 μ m.

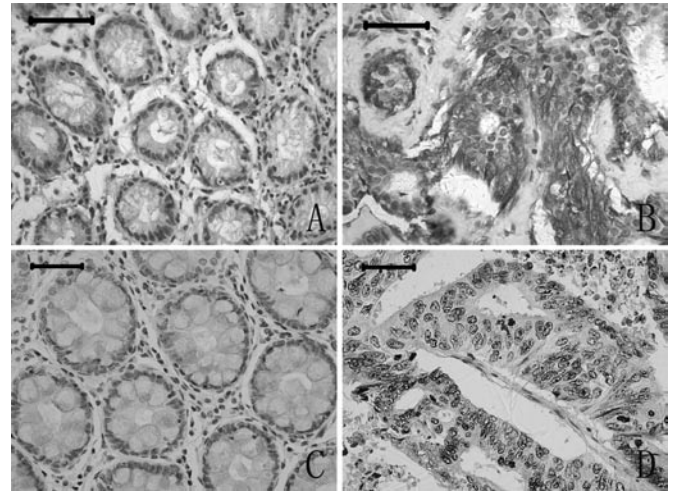


Figure 2. Ferritin and Ki67 protein expression in CRC (B and D) and adjacent normal glandular organ tissues (A and C). Scale bars, 50 μ m.

Table I. Relationship between clinicopathological features and NGALR and/or NGAL positivity.

Parameters	NGALR		NGAL		NGAL/NGALR	
	% positive rate	P-value	% positive rate	P-value	% positive rate ^a	P-value
Mean age, years (range)						
<57 (46)	39.1 (18/46)	0.587	30.4 (14/46)	0.684	19.6 (9/46)	0.825
≥57 (56)	33.9 (19/56)		26.8 (15/56)		17.9 (10/56)	
Gender						
Male	30.2 (16/53)	0.184	22.4 (11/49)	0.198	17.0 (9/53)	0.675
Female	42.9 (21/49)		34.0 (18/53)		20.4 (10/49)	
Regional lymph node						
N0	28.8 (15/52)	0.112	25.0 (13/52)	0.593	13.5 (7/52)	0.172
N1	44.0 (22/50)		32.0 (16/50)		24.0 (12/50)	
Primary tumour						
T1/T2	10.5 (2/19)	0.018	10.5 (2/19)	0.026	5.3 (1/19)	0.049
T3	38.3 (23/60)		26.6 (16/60)		16.7 (10/60)	
T4	52.2 (12/23)		47.8 (11/23)		34.8 (8/23)	
Histopathology						
Well-differentiated	22.7 (5/22)	0.193	9.1 (2/22)	0.057	0 (0/22)	0.004
Moderately differentiated	38.0 (27/71)		32.4 (23/71)		21.3 (15/71)	
Poorly differentiated	55.6 (5/9)		44.4 (4/9)		44.4 (4/9)	
TNM stages						
I/IIa	23.7 (9/38)	0.042	23.7 (9/38)	0.413	7.9 (3/38)	0.032
IIb/III/IV	43.6 (28/64)		31.3 (20/64)		25.0 (16/64)	
Colorectal cancer	36.3 (37/102)	0.000	28.4 (29/102)	0.000		
Normal mucosa	8.6 (8/93)		0 (0/81)			

^aPositive, both NGAL and NGALR were positive in one case. Positive rate, positive cases in the proportion of the overall.

($P<0.01$) and TNM stages ($P<0.05$). No significant associations with age, gender and patient survival were found for either group (Table I; Fig. 3).

NGAL and NGALR expression is associated with iron transport in cancer cells and increased cell proliferation. To explore whether a high expression of NGAL and NGALR

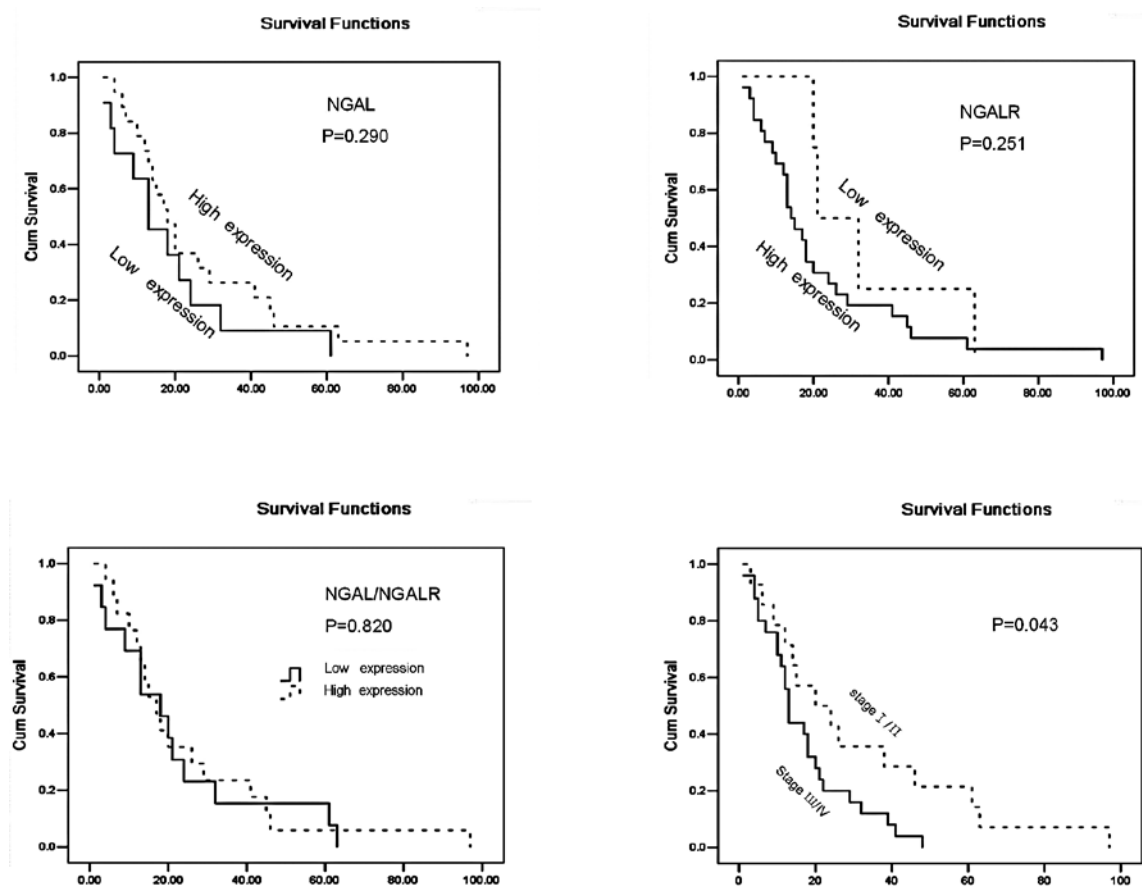


Figure 3. Kaplan-Meier estimates of the overall survival by NGAL, NGALR and NGAL/NGALR and TNM stage status.

is associated with iron transport and cell proliferation, we measured the expression level of ferritin and Ki67 in CRC cells by immunohistochemistry. Ferritin-positive staining was observed in the cytoplasm of tumour and normal cells (Figs. 2A and B). However, of the 102 tumor cases examined, ferritin expression was positive in 40 (39.2%), whereas of the 57 normal specimens examined, ferritin expression was positive in only 2 cases (3.51%); a statistically significant difference ($P < 0.01$). Expression of Ki67 was observed in the nucleus of CRC tumour and normal cells (Figs. 2C and D). Of the 102 tumour specimens examined, Ki67 expression was positive in 54 cases (52.9%), compared to 7 positive specimens in 44 normal cases (15.9%); again a significant difference in expression levels ($P < 0.01$).

As for the relationships among the proteins themselves, there were significant positive associations between the expression of NGAL and ferritin ($r = 0.374$, $P < 0.001$), as well as NGALR and Ki67 ($r = 0.228$, $P < 0.05$). An analysis of the relationship between NGAL/NGALR co-expression and the expression of ferritin or Ki67 revealed significantly positive correlations for NGAL/NGALR co-expression with ferritin ($r = 0.349$, $P < 0.001$) and Ki67 ($r = 0.205$; $P < 0.05$) in CRC.

Discussion

NGAL, a member of the lipocalin family, was originally discovered as a protein stored in specific granules of the human neutrophil (2). Our group reported that the over-expression of

NGAL is associated with poor differentiation of oesophageal squamous cell carcinoma (3). In the present study, we found that the expression of NGAL is significantly increased in CRC tissues, and that NGAL/NGALR co-expression is associated with poor cellular differentiation.

A series of studies suggested that NGAL is a novel iron transporter with functions distinct from those of transferrin (16-18). Moreover, a specific cell-surface receptor of 24p3/NGAL (24p3R/NGALR) was cloned (17). Over-expression of 24p3R/NGALR in cells induces binding and the uptake of 24p3/NGAL, which results in specific biological responses: iron-loaded 24p3/NGAL increases the intracellular iron concentration without promoting apoptosis, while iron-lacking 24p3/NGAL decreases intracellular iron levels, which induces expression of the proapoptotic protein Bim, leading to apoptosis (17). In the present study, NGALR was observed to be significantly up-regulated in CRC tissues. Furthermore, the co-expression of NGAL and NGALR is associated with CRC development, as is the expression of ferritin and Ki67. A correlation analysis revealed significant positive correlations for NGAL and NGALR, NGAL and ferritin, as well as NGALR and Ki67. These data suggest that NGAL and NGALR participate in intracellular iron transport/accumulation and contribute to the poor differentiation of tumour cells. Results of a large prospective epidemiological study suggest that iron results in an increased risk for CRC (19). High dietary iron promotes the production of reactive oxygen species, which activates the activator protein 1 and nuclear factor κ B signal

transduction pathways, leading to transcription of the genes involved in cell growth regulation (20-24). We assume that NGAL and NGALR play an important role in CRC through an increase of the iron content of cells, possibly activating some iron-sensitive gene(s) involved in tumour infiltration.

Our previous work showed that NGAL expression is significantly correlated with the depth of tumour invasion in oesophageal squamous cell carcinoma, a pathological process accompanied by over-expression of the NGAL/MMP-9 (matrix metalloproteinase 9) complex (4). MMP-9 is a proteolytic enzyme that degrades the extracellular matrix leading to connective tissue remodelling during normal biological processes and tumour invasion. NGAL directly associates with MMP-9, protecting it from degradation and resulting in increased MMP-9 activity (14,25,26). Interestingly, we found that the levels of NGAL, NGALR and NGAL/NGALR are associated with CRC invasion. A role for NGAL in breast cancer invasion has been suggested following the observation that the NGAL-overexpressing human breast cancer cell line MCF-7 exhibits an elevated tumour cell proliferative fraction (27).

Several examples of ligand-receptor interactions involved in tumour progression are known. Relaxin binds to LGR7 (relaxin receptor) and activates signalling cascades, leading to changes in tumour cell proliferation and altered motility (28). Receptor activator of nuclear factor κ B is expressed on prostate cancer cells and promotes invasion in a receptor activator of nuclear factor κ B ligand-dependent manner (29). Koshiba *et al* reported that the SDF-1/CXCR4 receptor-ligand system may be involved in the progression of pancreatic cancer, participating in tumour cell migration and angiogenesis (30). In light of these discoveries, we speculate that the NGAL/NGALR interaction is a key regulator of tumour growth and invasion in two distinct pathways: one regulating genes which promote cancer invasion, and the other activating iron-related genes also involved in cancer invasion. However, whether the aberrant expression of NGAL and NGALR in these pathways leads to an enhanced invasion of CRC needs to be investigated.

This study is the first to investigate expression of the NGALR protein in a large series of CRC, and to reveal a role of NGAL and NGALR in tumour invasion and cell differentiation in CRC. The most important findings are: i) expression of NGAL, NGALR, ferritin and Ki67 is elevated in CRC; and ii) over-expression of NGAL and NGALR is associated with cell differentiation and tumour invasion in CRC. Thus, our findings suggest that NGAL and NGALR are involved in the transformation and progression of CRC. Therefore, NGALR may be a novel target for the treatment of CRC.

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