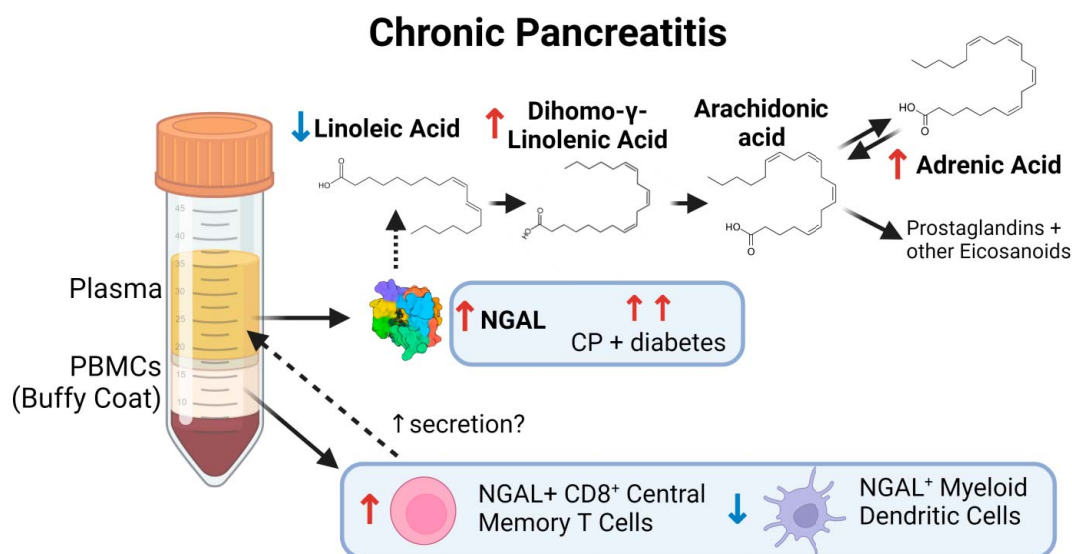


Systemic Neutrophil Gelatinase-Associated Lipocalin Alterations in Chronic Pancreatitis: A Multicenter, Cross-Sectional Study

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NGAL Alterations in Chronic Pancreatitis



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- INTRODUCTION:** Chronic pancreatitis (CP) is a progressive fibroinflammatory disorder lacking therapies and biomarkers. Neutrophil gelatinase-associated lipocalin (NGAL) is a proinflammatory cytokine elevated during inflammation that binds fatty acids (FAs) such as linoleic acid. We hypothesized that systemic NGAL could serve as a biomarker for CP and, with FAs, provide insights into inflammatory and metabolic alterations.
- METHODS:** NGAL was measured by immunoassay, and FA composition was measured by gas chromatography in plasma (n = 171) from a multicenter study, including controls (n = 50), acute and recurrent acute pancreatitis (AP/RAP) (n = 71), and CP (n = 50). Peripheral blood mononuclear cells (PBMCs) from controls (n = 16), AP/RAP (n = 17), and CP (n = 15) were measured by cytometry by time-of-flight.
- RESULTS:** Plasma NGAL was elevated in subjects with CP compared with controls (area under the curve [AUC] = 0.777) or AP/RAP (AUC = 0.754) in univariate and multivariate analyses with sex, age, body mass index, and smoking (control AUC = 0.874; AP/RAP AUC = 0.819). NGAL was elevated in CP and diabetes compared with CP without diabetes ($P < 0.001$). NGAL⁺ PBMC populations distinguished CP from controls (AUC = 0.950) or AP/RAP (AUC = 0.941). Linoleic acid was lower, whereas dihomo- γ -linolenic and adrenic acids were elevated in CP ($P < 0.05$). Linoleic acid was elevated in CP with diabetes compared with CP subjects without diabetes ($P = 0.0471$).
- DISCUSSION:** Elevated plasma NGAL and differences in NGAL⁺ PBMCs indicate an immune response shift that may serve as biomarkers of CP. The potential interaction of FAs and NGAL levels provide insights into the metabolic pathophysiology and improve diagnostic classification of CP.

KEYWORDS: lipocalin 2; body mass index; smoking; mass CyTOF; peripheral blood mononuclear cells; linoleic acid

SUPPLEMENTARY MATERIAL accompanies this paper at <http://links.lww.com/CTG/B85>.

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INTRODUCTION

Pancreatitis is an inflammatory disease that ranges from acute (AP) to recurrent acute (RAP) to chronic pancreatitis (CP), and cases have increased by 13.3% in the past 2 decades (1). Patients with CP are at risk of long-term complications, including chronic abdominal pain, diabetes, exocrine pancreatic dysfunction (EPD), osteopathy, and pancreatic cancer (2). Our understanding of CP pathophysiology remains limited, leading to extensive testing, delayed diagnosis, and a lack of biomarkers and effective therapies (3–5).

Our systematic review investigating potential CP biomarkers indicated that neutrophil gelatinase-associated lipocalin (NGAL/lipocalin 2) was elevated in pancreatic juice of CP patients, warranting further evaluation as a potential biomarker of CP (5). NGAL is a secreted protein in the lipocalin family that binds hydrophobic ligands, including fatty acids (FAs) such as linoleic acid, and siderophores, as part of an acute-phase inflammatory response (6–11). Increased NGAL expression promotes lipolysis and FA oxidation, releasing FAs from intracellular storage during inflammation (12). NGAL contributes to pancreatic cancer pathogenesis by regulating inflammation in pancreatic tumors and is increased in the blood of subjects with pancreatic cancer compared with controls (13). Increased NGAL expression in immune cell subtypes within the peripheral blood mononuclear cells (PBMCs), such as B cells in chronic lymphocytic leukemia (14) and CD4⁺ T cells in acute kidney injury (15), is associated with apoptosis inhibition and enhanced proinflammatory or profibrotic effects (14,16). Although studies have assessed NGAL as a potential biomarker for CP in various biofluids, the studies were small, lacked well-controlled sample acquisition and clinical criteria, and focused on a comparison between CP and pancreatic cancer instead of a control group (13,17–19).

We evaluated whether NGAL levels or NGAL⁺ PBMCs could serve as diagnostic biomarkers of CP and gain insights into inflammatory alterations associated with CP. Furthermore, we investigated whether FAs known to interact with NGAL were altered in CP to better understand the metabolic alterations in this disease.

METHODS

Study design

This study used well-defined human samples from the prospective multicenter study Prospective Evaluation of Chronic Pancreatitis for Epidemiologic and Translational Studies (PROCEED) (NCT03099850), a study of the Chronic Pancreatitis, Diabetes, and Pancreatic Cancer consortium (20,21). Subjects selected for the current analysis were enrolled into PROCEED, from June 2017 to September 2020 and included a random sampling of healthy controls (n = 50), AP/RAP (n = 71), and CP (n = 50). Sample size was determined by a small pilot study using a separate set of samples from the Ohio State University. PBMCs from controls (n = 16), AP/RAP (n = 17; 1 subject with AP), and CP (n = 15) subjects were selected from the PROCEED biorepository based on sample availability. AP and RAP were diagnosed using the revised Atlanta criteria (22), and CP was diagnosed as previously described (20) based on pancreatic calcifications and/or advanced Cambridge stage. Inclusion and exclusion criteria for the PROCEED cohort included in this study are previously published and summarized in Supplementary Table 1 (see Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>) (20,21). Samples (plasma, urine, and PBMCs) were collected 1 month after an acute attack to prevent bias from acute inflammation. Data for sex, age, body mass index (BMI), etiology, presence of EPD, diabetes status, and smoking history were retrieved from PROCEED (20).

Enzyme-linked immunosorbent assay

Plasma and urine NGAL concentrations were measured using an enzyme-linked immunosorbent assay (DLCN20; R&D Systems, Minneapolis, MN) following the manufacturer's recommended protocol. All samples were measured in duplicate and reported in nanograms per milliliter determined from a standard curve. The personnel and study team were blinded to group assignment.

Mass cytometry by time-of-flight

PBMCs were thawed and processed as previously described (21). PBMCs were stained using the Maxpar Direct Immune Profiling Assay kit (PN 400286 B1; Standard BioTools, San Francisco, CA) with the addition of CD11b (Standard BioTools) and CD33 (BioLegend, San Diego, CA) antibodies (see Supplementary Table 2, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>) (23). After surface staining, the cells were fixed, permeabilized, and intracellularly stained for NGAL (ab224264; Abcam, Cambridge, UK), which was conjugated to metal isotope ¹⁵⁹Tb, using the Maxpar X8 Antibody Labeling Kit, ¹⁵⁹Tb-4 RXN (Standard BioTools, see Supplementary Table 2, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>). The stained PBMCs were analyzed on a Fluidigm Helios Mass Cytometer. Data concatenation and normalization were performed using the Cytobank software v7.0 (Beckman Coulter). Samples with more than 100,000 events were used for analysis. PBMC populations and subsets were gated based on phenotypic markers of each population (see Supplementary Table 3, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>) (23). Differences in NGAL⁺ of PBMC populations were calculated as the number of cells positive for NGAL within an immune population divided by the number of live PBMCs positive for NGAL. Differences in NGAL⁺ PBMC subsets were calculated as the number of cells positive for NGAL within an immune subpopulation divided by the number of cells positive for NGAL within the respective parent population (see Supplementary Table 3, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>).

FA composition analysis

Total lipids were extracted from plasma samples using the 2:1 chloroform:methanol method and methylated (24,25) as we previously described (26–29), blinded to disease group. The methyl ester form of the FAs was analyzed using gas chromatography run on a 30-m Omegawax TM 320 fused silica capillary column (Supelco, Bellefonte, PA), with the oven temperature and carrier gas flow rate settings as previously described (29,30). Results were compared with purchased standards for each FA methyl ester (Matreya, LLC, Pleasant Gap, PA; Nu-Check Prep, Elysian, MN). The data are presented as the relative composition of all identified FAs (% area), as recommended by experts in the field for clinical studies (30,31).

Statistical analysis

Statistical analyses were performed using GraphPad Prism 9 (GraphPad Software, San Diego, CA), JMP 16 (SAS Institute, Cary, NC), and R version 4.1.2 (R Foundation for Statistical Computing, Vienna, Austria). Relative frequencies were reported and compared for categorical variables using Fisher exact test or χ^2 tests of independence, whereas mean values and SDs were reported and compared by study group for the continuous age variable using 1-way ANOVA. Because EPD was an exclusion

criterion for the control group in the PROCEED study and none of our control subjects had diabetes, we did not include controls when comparing the difference in these clinical factors using χ^2 tests between AP/RAP and CP. Propensity score matching on the covariates of sex, age, BMI, and smoking was used to calculate the average treatment effect for NGAL when comparing control with CP, control with AP/RAP, and AP/RAP with CP. Receiver operator characteristic curves were generated, the area under the curve (AUC) was calculated to assess the ability of NGAL concentration to distinguish between study groups, and a multiple logistic regression was used to control for clinical characteristics. A 2-way ANOVA with Tukey's testing for pairwise comparisons on log-transformed NGAL concentrations compared NGAL across groups by categorical subject characteristics. Because age is a continuous variable, we performed a correlation analysis between age and NGAL and assessed differences in the correlation slopes and intercepts. Nonparametric Kruskal-Wallis tests were performed on the percentages of NGAL⁺ PBMC populations and subsets and on overall PBMC populations and subsets to compare the differences between controls, AP/RAP, and CP. FA composition was compared among the disease groups using 1-way ANOVA parametric test, or Kruskal-Wallis nonparametric test, with Holm-Šidák correction for multiple testing. T-test or Mann-Whitney nonparametric test was used to assess differences in FA composition in CP based on diabetes.

RESULTS

Study group characteristics

All groups had similar sex distributions. On average, the CP group was older, leaner, more likely to smoke, and more likely to have EPD compared with the AP/RAP group and, where applicable, the control group. Idiopathic and hypertriglyceridemia etiologies were more common in the AP/RAP group, whereas alcoholic and genetic etiologies were more common in the CP group (Table 1, top). In the subset of subjects with available PBMCs, age, etiology, EPD, and smoking were all different between the controls and disease groups (Table 1, bottom).

Plasma NGAL is elevated in CP

Plasma NGAL levels were higher, on average, in CP compared with control and AP/RAP (Figure 1a). Two-way ANOVAs assessing the interaction between study group and the clinical characteristics of sex, BMI, pancreatitis etiology, EPD, and smoking history and a correlation analysis between NGAL and age demonstrated that NGAL remained higher, on average, in CP compared with control and AP/RAP within the subgroups of each of the clinical features, indicating the differences in NGAL are attributable to disease group rather than these clinical features (see Supplementary Figure 1A–F, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>). NGAL was significantly elevated in subjects with both CP and diabetes, compared with CP subjects without diabetes ($P = 0.0009$) (Figure 1b). We did not observe any significant difference in the urine NGAL levels between any of the groups (see Supplementary Figure 2, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>). We generated receiver operator characteristic curves to assess NGAL's potential as a biomarker and found that NGAL distinguished CP from control (AUC = 0.777; $P < 0.0001$) (Figure 1c) and CP from AP/RAP (AUC = 0.754; $P < 0.0001$) (Figure 1d). Propensity score matching to mitigate the possible effect of clinical features (age, sex, BMI, and smoking) on the difference in

Table 1. Comparison of patient characteristics by PROCEED study group

Plasma/urine	Control n = 50	AP/RAP n = 71	CP n = 50	P value
Age, yr, mean (SD)	51.4 (14.1)	48.7 (13.8) ^a	55.5 (11.9) ^a	0.028
Sex, n (%)				0.915
Male	29 (58.0)	42 (59.2)	31 (62.0)	
Female	21 (42.0)	29 (40.8)	19 (38.0)	
BMI, kg/m ² , n (%)				<0.001
<24.9	14 (28.0)	27 (38.0)	34 (68.0)	
25–29.9	26 (52.0)	26 (36.6)	9 (18.0)	
>30	10 (20.0)	18 (25.4)	7 (14.0)	
Etiology, n (%)				0.006
Idiopathic	N/A	41 (57.7)	17 (34.0)	
Alcoholic		18 (25.4)	26 (52.0)	
Genetic		1 (1.4)	3 (6.0)	
Obstructive		2 (2.8)	1 (2.0)	
Hypertriglyceridemia		9 (12.7)	1 (2.0)	
Postnecrotic		0 (0.0)	1 (2.0)	
Other		0 (0.0)	1 (2.0)	
EPD present, n/total n (%)	N/A	10/49 (20.4) ^b	24/40 (60.0) ^b	<0.001
Diabetes present, n/total n (%)	N/A	18/71 (25.4)	19/49 (38.8) ^b	0.118
Smoking, n (%)				<0.001
Current	2 (4.0)	13 (18.3)	25 (50.0)	
Past	10 (20.0)	23 (32.4)	13 (26.0)	
Never	38 (76.0)	35 (49.3)	12 (24.0)	
PBMCs	Control n = 16	AP/RAP n = 17	CP n = 15	P value
Age, yr, mean (SD)	45.7 (11.9)	40.2 (11.2)	54.5 (10.4)	0.006
Sex, n (%)				0.608
Male	9 (56.3)	11 (64.7)	8 (53.3)	
Female	7 (53.7)	8 (35.3)	7 (46.7)	
BMI, kg/m ² , n (%)				0.069
<24.9	6 (37.5)	4 (23.5)	8 (53.3)	
25–29.9	7 (43.8)	3 (17.7)	4 (26.7)	
>30	3 (18.8)	10 (58.8)	3 (20.0)	
Etiology, n (%)				0.030
Idiopathic	N/A	9 (52.9)	7 (46.7)	
Alcoholic		4 (23.5)	8 (53.3)	
Genetic		0 (0.0)	0 (0.0)	
Obstructive		0 (0.0)	0 (0.0)	
Hypertriglyceridemia		4 (23.5)	0 (0.0)	
Postnecrotic		0 (0.0)	0 (0.0)	
Other		0 (0.0)	0 (0.0)	
EPD present, n (%)	N/A	3/11 (27.3) ^b	10/13 (76.9) ^b	0.038
Diabetes present, n (%)	N/A	5 (29.4)	9 (60.0)	0.080

Table 1. (continued)

PBMCs	Control n = 16	AP/RAP n = 17	CP n = 15	P value
Smoking, n (%)				0.001
Current	0 (0.0)	5 (29.4)	9 (60.0)	
Past	3 (18.8)	5 (29.4)	4 (26.7)	
Never	13 (81.3)	7 (41.2)	2 (13.3)	

Results for subjects with available plasma and urine are shown at the top, whereas results for subjects with available PBMCs are shown at the bottom. Subject age was analyzed using 1-way ANOVA with Tukey's HSD for multiple comparisons. Subject sex, BMI, pancreatitis etiology, EPD, diabetes status, and cigarette use were analyzed using χ^2 analysis.

AP, acute pancreatitis; BMI, body mass index; CP, chronic pancreatitis; EPD, exocrine pancreatic dysfunction; HSD, honestly significant difference; PBMC, peripheral blood mononuclear cell; RAP, recurrent acute pancreatitis.

^a $P < 0.05$ between AP/RAP and CP.

^bMissing data from some of the cases.

NGAL levels between groups continued to provide evidence of significantly higher NGAL levels in subjects with CP compared with controls ($P = 0.011$) and AP/RAP ($P = 0.007$) (Table 2). The matching resulted in similar AUCs for NGAL differentiating CP vs control or AP/RAP (Table 2).

To evaluate the predictive performance of NGAL in combination with clinical features, multivariate models to predict CP

(vs control and vs AP/RAP) were fit using NGAL, age, sex, BMI, and smoking as predictors. Diabetes and EPD were excluded because these are complications of CP that may cause an over-estimation of NGAL's discriminative ability. The regression results (Table 3) showed that a model that includes clinical features in addition to NGAL improved AUC over NGAL alone for CP compared with control ($AUC = 0.874$; $P < 0.0001$) and for CP

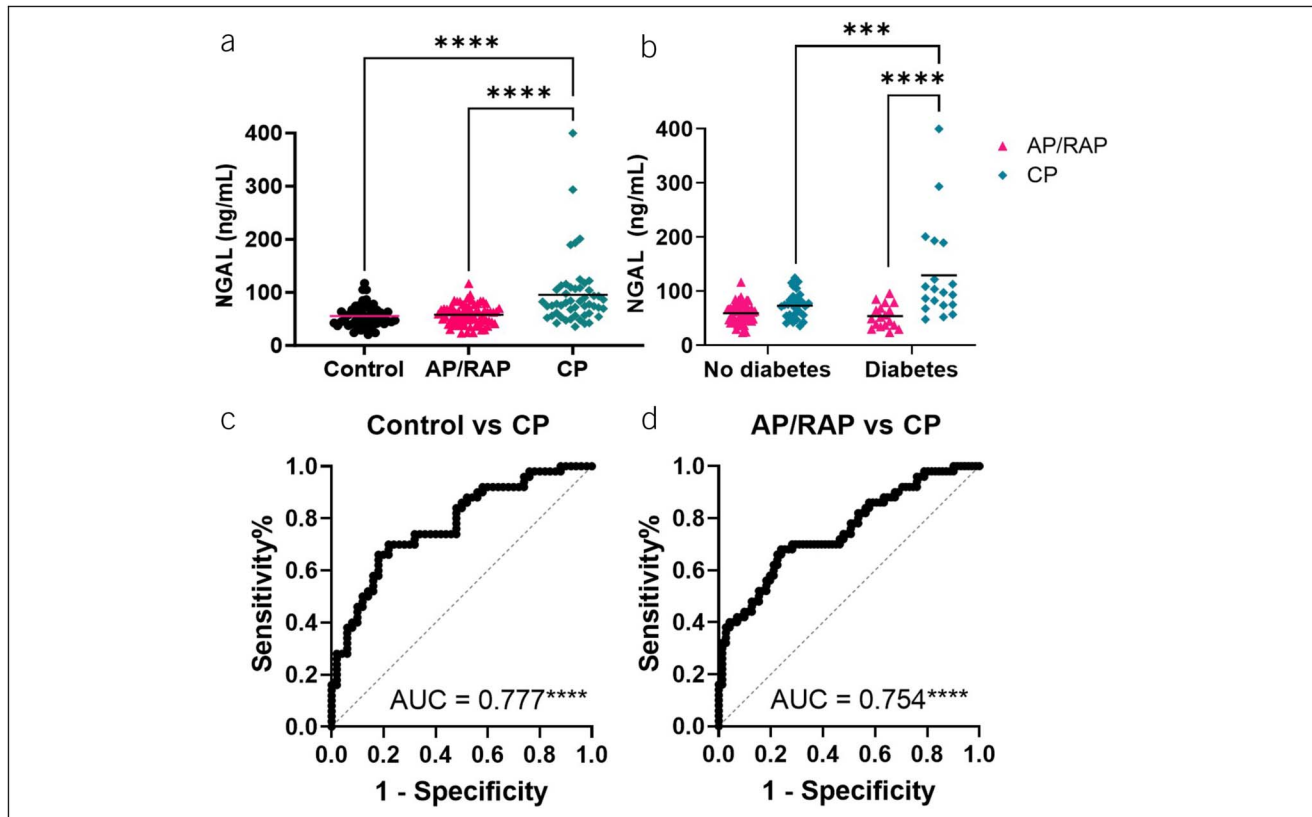


Figure 1. NGAL expression increases in the plasma of subjects with CP. (a) NGAL measured by ELISA from the multicenter (PROCEED) study discovery set. Statistical significance was determined by a non-parametric 1-way ANOVA with Tukey's multiple comparisons correction. (b) AP/RAP and CP samples were subdivided by diabetes status and compared for plasma NGAL concentration. Statistical significance was determined by 2-way ANOVA with Tukey's multiple testing correction. (c) ROCs with AUCs of plasma NGAL comparing control vs CP and (d) AP/RAP vs CP. *** $P < 0.001$, **** $P < 0.0001$. AP, acute pancreatitis; CP, chronic pancreatitis; ELISA, enzyme-linked immunosorbent assay; NGAL, neutrophil gelatinase-associated lipocalin; RAP, recurrent acute pancreatitis.

Table 2. Propensity score matching for plasma NGAL in CP

Disease	ATT (SE)	f statistic	ROC AUC
CP (n = 50) vs control (n = 21)	42.06 (16.56)	2.54*	0.785***
CP (n = 50) vs AP/RAP (n = 27)	35.83 (13.33)	2.69**	0.739***
AP/RAP (n = 71) vs control (n = 34)	6.27 (4.83)	1.30	

ATT, average treatment effect on the disease group; AP, acute pancreatitis; AUC, area under the curve; CP, chronic pancreatitis; RAP, recurrent acute pancreatitis; ROC, receiver operator characteristic.
* $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$.

compared with AP/RAP (AUC = 0.819; $P < 0.0001$). In the multivariable models, NGAL significantly contributed to the differentiation of CP and controls ($P < 0.001$) and CP and AP/RAP ($P < 0.001$) after controlling for these clinical features. History of smoking and low BMI also made a significant contribution to the differentiation of CP vs controls or AP/RAP.

Profiling PBMCs identified altered NGAL⁺ immune cell populations in CP

Immune cells are one of several sources of systemic NGAL, particularly in inflammatory conditions (15). Therefore, we performed mass cytometry by time-of-flight (CyTOF) to determine whether there were differences in NGAL⁺ PBMCs for patients with CP. When assessing NGAL⁺ cells as a percentage of live PBMCs, there were more NGAL⁺ B cells in CP compared with controls ($P = 0.021$) (Figure 2a), with similar results in the overall PBMC populations (see Supplementary Figure 3A, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>). We observed differences in NGAL⁺ PBMC subpopulations as a percentage of the NGAL⁺ parent population between controls, AP/RAP, and CP (Figure 2b). Compared with controls, CP subjects had less NGAL⁺ naive CD4⁺ T cells ($P = 0.037$) and memory B cells ($P = 0.019$) and more NGAL⁺ CD8⁺ central memory T cells ($P = 0.009$). Compared with AP/RAP, the CP group had fewer NGAL⁺ memory resting B cells ($P = 0.027$) and myeloid dendritic cells ($P = 0.042$) and more NGAL⁺ CD8⁺ central memory T cells ($P = 0.001$) and CD8⁺ effector T cells ($P = 0.041$). There were fewer NGAL⁺ CD16⁻ and more NGAL⁺ CD16⁺ natural killer cells ($P = 0.034$) in subjects with AP/RAP compared with control subjects (Figure 2b). Some of these shifts occurred in the overall PBMCs immune cell subpopulations (see Supplementary Figure 3B, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>). These differences in immune cells suggest a shift away from an innate immune response and toward an adaptive immune response during CP that is enhanced in NGAL⁺ cells.

Multiple logistic regressions of the immune cell populations that were significantly different between CP and either the control or AP/RAP groups were fit to determine the utility of this combination of NGAL⁺ PBMC subpopulations for distinguishing CP from either controls or AP/RAP. A panel of NGAL⁺ CD8⁺ central memory T cells, CD8⁺ effector T cells, naive CD4⁺ T cells, memory resting B cells, and myeloid dendritic cells differentiated control from CP (AUC = 0.950; $P < 0.0001$) and AP/RAP from CP (AUC = 0.941; $P < 0.0001$) (Table 3). By contrast, the same overall PBMC subpopulations resulted in lower AUCs comparing control with CP (AUC = 0.883; $P < 0.0001$) and AP/RAP with CP (AUC = 0.890; $P < 0.0001$) (see Supplementary Table 6, Supplementary Digital Content 1, <http://links.lww.com/CTG/B85>). This

suggests that a subset of circulating NGAL⁺ immune cells could be used as a potential biomarker panel for CP.

FA composition is altered in CP

Because NGAL binds FAs such as linoleic acid, we measured FA composition in the plasma of the same subjects for whom we measured NGAL levels. Three polyunsaturated FAs (PUFAs) were differentially affected in CP compared with controls or AP/RAP. Linoleic acid was lower, whereas dihomo- γ -linolenic and adrenic acids were higher in CP compared with AP/RAP (Figure 3a). Because several FAs interact with NGAL and NGAL is modulated by both CP and diabetes, we assessed whether FA composition was also different in subjects with CP and diabetes compared with those without diabetes. Interestingly, like NGAL, plasma linoleic acid was also significantly higher in subjects with diabetes and CP compared with those with CP without diabetes ($P = 0.0471$) (Figure 3b). Overall, the differences observed suggest a dysregulation of PUFAs in CP.

DISCUSSION

One of the key distinctions between AP/RAP and CP is the constant proinflammatory state in CP, leading to complications such as fibrosis and calcification. However, these complications are typically observable by imaging once CP is severe as the spectrum of pancreatitis lacks biomarkers to discern between disease stages. In this study, we demonstrated elevated plasma NGAL levels in patients with CP, especially those with diabetes. NGAL emerged as a promising biomarker for CP, particularly when coupled with easily accessible clinical characteristics. Furthermore, our investigation revealed alterations in NGAL⁺ PBMCs, suggesting a transition from a naive state to memory and effector cell types in CP. These differences could potentially form a panel for distinguishing CP from controls or AP/RAP. Furthermore, we identified differences in plasma linoleic acid composition in CP and CP combined with diabetes. The changes observed in NGAL and PUFAs known to interact with NGAL in CP reflect metabolic changes that may explain some of the pathophysiologic features of CP.

NGAL regulates inflammation, particularly in cancers such as pancreatic cancer, and has been suggested to have clinical value in diagnosing the spectrum of pancreatic diseases, including AP, CP, and pancreatic cancer (7). However, previous studies assessing NGAL as a biomarker of pancreatic diseases mainly compared CP with pancreatic cancer or AP to controls (5,7). These studies had low sample sizes, different criteria to define CP, and did not account for the impact the timing of sample collection, and clinical characteristics could have had on plasma NGAL concentrations. The multicenter PROCEED discovery sample set used in this

Table 3. Multiple logistic regressions using plasma NGAL (top) or NGAL⁺ PBMC subpopulations (bottom) to predict CP

	Odds ratio	95% CI	P value	AUC
Plasma NGAL				
Control vs CP				0.874****
Intercept	5.52	0.20–170.4	0.316	
NGAL (1 ng/mL)	1.05	1.02–1.08	<0.001	
Sex (male)	0.63	0.20–1.85	0.405	
Age (yr)	1.01	0.97–1.05	0.809	
BMI (kg/m ²)	0.87	0.77–0.96	0.012	
Smoking (never)	0.14	0.04–0.38	<0.001	
AP/RAP vs CP				0.819****
Intercept	0.56	0.03–10.5	0.703	
NGAL (1 ng/mL)	1.04	1.02–1.07	<0.001	
Sex (male)	0.89	0.35–2.27	0.809	
Age (yr)	1.03	1.0–1.07	0.096	
BMI (kg/m ²)	0.87	0.78–0.95	0.003	
Smoking (never)	0.42	0.16–1.05	0.069	
NGAL⁺ PBMCs				
Control vs CP				0.950****
Intercept	10.39	0.002–93833	0.585	
CD8 ⁺ central memory T cells	1.59	1.13–2.82	0.033	
CD8 ⁺ effector T cells	0.98	0.89–1.08	0.750	
Naive CD4 ⁺ T cells	0.95	0.85–1.04	0.273	
Memory resting B cells	0.83	0.60–0.96	0.137	
Myeloid dendritic cells	0.94	0.82–1.05	0.298	
AP/RAP vs CP				0.941****
Intercept	10.64	0.001–645744	0.631	
CD8 ⁺ central memory T cells	1.63	1.14–3.15	0.051	
CD8 ⁺ effector T cells	1.06	0.94–1.22	0.392	
Naive CD4 ⁺ T cells	1.04	0.92–1.19	0.565	
Memory resting B cells	0.90	0.65–1.19	0.447	
Myeloid dendritic cells	0.88	0.75–0.98	0.044	

PROCEED study NGAL⁺ PBMC subpopulations that were significantly different between CP and either control or AP/RAP were considered. AP, acute pancreatitis; BMI, body mass index; CP, chronic pancreatitis; NGAL, neutrophil gelatinase-associated lipocalin; PBMC, peripheral blood mononuclear cell; RAP, recurrent acute pancreatitis. ****ROC AUC $P < 0.0001$.

study enabled us to compare larger numbers of well-defined pancreatitis subjects to healthy controls, allowing us to better adhere to the phase 1 of the prospective-specimen-collection, retrospective-blinded-evaluation study design for biomarker discovery (20,32,33). A recent study using the Olink technology in a similar sample set from the PROCEED study identified IL-17 as another potential biomarker of CP (NGAL was not a part of that cytokine panel) (34). Interestingly, IL-17 can promote expression of NGAL (35). Combining NGAL with other potential mechanistic markers such as IL-17 may improve accuracy for CP compared with controls or AP/RAP (36,37). Urine NGAL is currently being investigated as a biomarker for acute kidney injury (38,39), diabetic nephropathy, or diabetic kidney disease

(40,41), and others are working on ways to test plasma NGAL using automated assays (42). Because we did not see differences in NGAL levels in the urine, urine NGAL may be specific to kidney injury and not CP. It is important to note that we do not expect plasma NGAL to be a differential diagnostic marker between pancreatic and kidney disease or even other inflammatory disorders. Rather, persistent elevated plasma NGAL with other indicators of pancreatic disease may be markers of CP inflammation in patients suspected of having a pancreatic disease, but for whom the diagnosis is unclear.

Diabetes is a common complication of CP and often represents a progression in severity (2). Although many patients presenting with CP already have diabetes, those who do not are likely

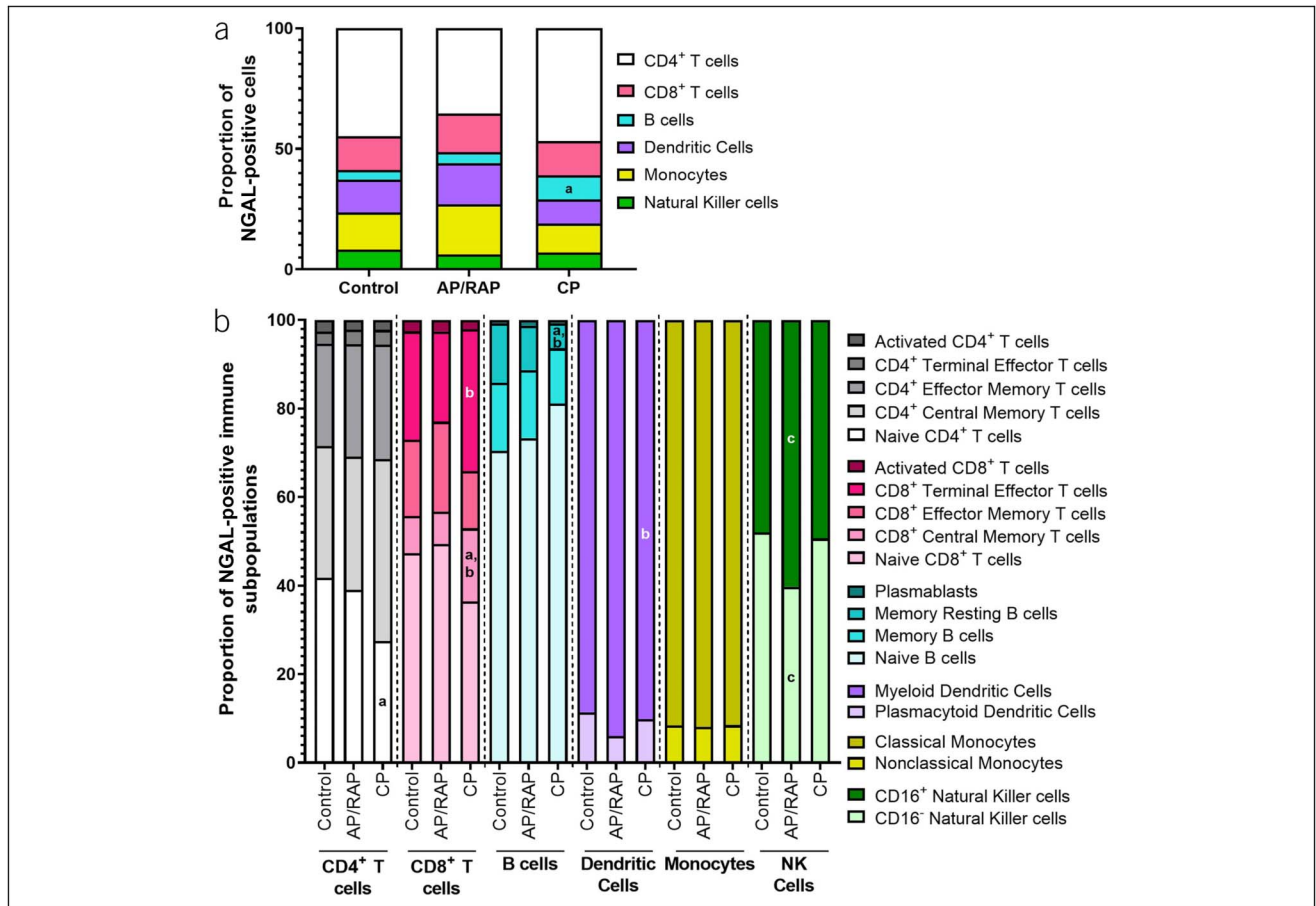


Figure 2. NGAL⁺ immune cells within PBMC populations and subsets tends to increase in the adaptive immune cells of subjects with CP. PBMCs from multicenter (PROCEED) discovery set (healthy controls, AP/RAP, and subjects with CP) collected during enrollment visit were analyzed by CyTOF. (a) Proportion of NGAL⁺ immune populations (b) and subpopulations. ^a $P < 0.05$ between control and CP; ^b $P < 0.05$ between AP/RAP and CP; ^c $P < 0.05$ between control and AP/RAP; **** $P < 0.001$. AP, acute pancreatitis; CP, chronic pancreatitis; CyTOF, mass cytometry by time-of-flight; NGAL, neutrophil gelatinase-associated lipocalin; PBMC, peripheral blood mononuclear cell; RAP, recurrent acute pancreatitis.

to develop type 3c diabetes as pancreatitis progresses (43). There is conflicting evidence on whether NGAL promotes insulin resistance or counteracts diabetes (44,45). However, several studies suggest NGAL is elevated in diabetes associated with microvascular complications (neuropathy and nephropathy) (40,41,46). Here, NGAL levels were higher in subjects with both CP and diabetes compared with CP alone, although we were unable to compare with a control group with diabetes because of a limitation in the PROCEED study design. Our analysis does not provide definitive certainty regarding whether NGAL concentrations in the CP and diabetes participants reflect the influence of the diabetic state or if it represents a more advanced clinical phenotype of disease (i.e., CP with endocrine insufficiency). Based on these data, future studies would include additional controls with diabetes to better understand how exocrine and endocrine diseases influence plasma NGAL levels.

NGAL is essential for innate and adaptive immune functions, with roles in iron homeostasis (10,47), T-cell and B-cell development and proliferation (48), antibody production (49), and dendritic cell antigen presentation (50). Although the processing and storage methods used in PROCEED (21) caused neutrophil degradation, an established source of NGAL (9,51,52), we detected a shift in NGAL⁺ PBMCs of subjects with CP away from

the naive state and toward memory and effector cell types. Unlike other studies, we did not observe a difference in circulating monocytes (53) or lymphocytes (54), and we saw an increase in CD8⁺ central memory T cells that conflicts with earlier studies (55). The differences are likely due to variations in detection method (e.g., flow cytometry (54,56,57), barcoding cells for CyTOF (53), and metal-labeled antibodies for CyTOF like the one used in this study), immune cell source (tissue (53,57) vs circulating (54,58)), and time after an acute attack (53,54,56,58). The increase observed in NGAL⁺ B cells and NGAL⁺ B-cell subpopulations detected in subjects with CP in our study results contrasts with a study that found B cells to only be increased in AP (and further increased in severe AP), compared with healthy subjects (54). However, since the samples in the aforementioned study were collected on admission rather than 1 month after an AP attack (as is the case for PROCEED), differences in these findings may be due to acute rather than chronic disease response. Overall, the differences in NGAL⁺ PBMCs point toward a rise in a cell-mediated immune response through cytotoxic mechanisms (14,59) that could lead to increased tissue immune infiltration and inflammation of subjects with CP. Furthermore, our data suggest that a small panel of NGAL⁺ immune cell populations may be developed as another effective diagnostic biomarker tool

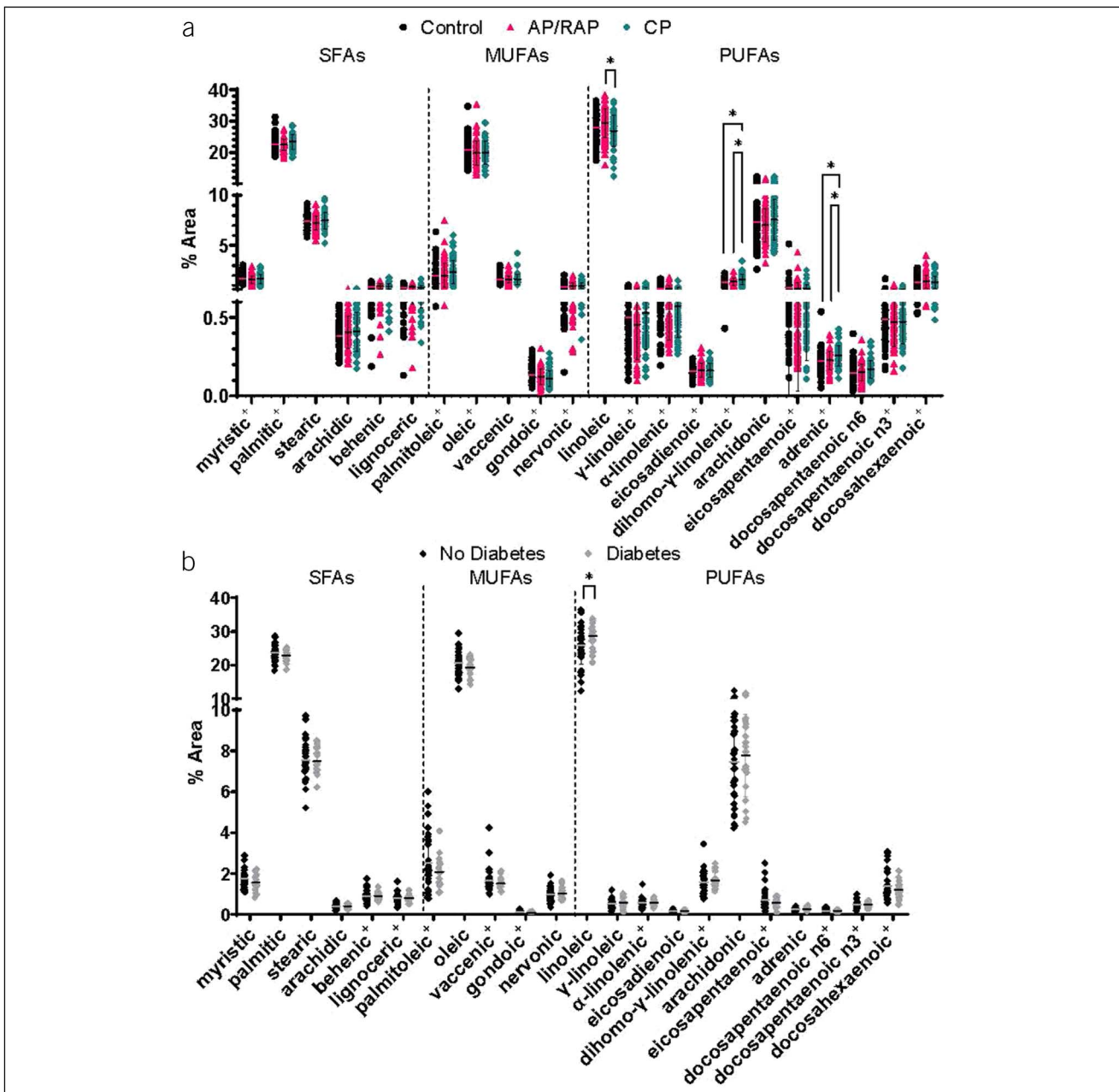


Figure 3. PUFA composition is altered in the plasma of subjects with CP. (a) Plasma FAs ordered by saturation and carbon chain number compared by 1-way ANOVA with multiple testing for significance. (b) Plasma FAs from subjects with CP subdivided by diabetes compared by *t* tests. †Nonparametric Kruskal-Wallis or Mann-Whitney **P* < 0.05. AP, acute pancreatitis; CP, chronic pancreatitis; FA, fatty acid; MUFA, monounsaturated FA; PUFA, polyunsaturated FA; RAP, recurrent acute pancreatitis; SFA, saturated FA.

for CP compared with the expensive and less widely available CyTOF technology; however, this will need to be validated.

NGAL interacts with several hydrophobic ligands such as linoleic, palmitic, and oleic acids, and these interactions have been associated with endothelial dysfunction and promoting inflammation (11). Arachidonic acid, a downstream metabolite of linoleic and dihomo- γ -linolenic acids, is further metabolized into eicosanoids, which are elevated in the pancreatic juice of subjects with CP (60) and can mediate pain sensation, inflammation, and promote fibrosis, all of which are present in CP (61,62). It is possible that excessive NGAL in CP may be binding and

trafficking linoleic acid into cells for use and metabolism; however, because we do not have dietary information, we cannot rule out the impact of an altered diet in subjects with CP. In subjects with both CP and diabetes, we observed an increase in linoleic acid, which was surprising because higher levels of linoleic acid have been correlated with a decreased incidence of type 2 diabetes mellitus (63). Further mechanistic studies will be required to tease out this complex interaction between CP and diabetes in modulating NGAL and linoleic acid to determine whether this interaction is targetable for future therapeutic development for pancreatitis.

This study on NGAL is strengthened over previous studies (5,7) from the use of an extensive set of blood and urine samples with associated clinical data gathered from multiple clinical centers using standardized procedures (21). We also conducted a comprehensive assessment of NGAL⁺ immune populations and subpopulations from PBMCs. Our study enabled a direct comparison of circulating NGAL in subjects with CP to subjects with AP/RAP and control subjects. Finally, we identified a potential association between alterations in NGAL levels and its hydrophobic ligands, particularly in CP and diabetes. In subsequent studies, we will explore whether binding and trafficking of linoleic acid by NGAL in CP contributes to pathophysiologic processes such as inflammation, fibrosis, and pain in the progression of pancreatic disease from acute pancreatitis to pancreatic cancer and validate NGAL as a biomarker of CP. Overall, this study presents a meticulous analysis of the diagnostic utility of NGAL in CP and identifies a potential biological relevance of elevated NGAL and its associated FAs in pancreatitis.

CONFLICTS OF INTEREST

Guarantor of the article: Zobeida Cruz-Monserrate, PhD.

Specific author contributions: K.G.-F. and K.C.: study concept and design; acquisition of data; analysis and interpretation of data; drafting of the manuscript; critical revision of the manuscript for important intellectual content; and statistical analysis. V.P.-G.:

acquisition of data; analysis and interpretation of data;

administrative, technical, or material support; and critical revision of

the manuscript for important intellectual content. M.T. and T.P.:

acquisition of data; administrative, technical, or material support.

T.A.M.: acquisition of data; analysis and interpretation of data; and

critical revision of the manuscript for important intellectual content.

R.M.C. and M.A.B.: acquisition of data; analysis and interpretation of

data; administrative, technical, or material support; and critical

revision of the manuscript for important intellectual content. S.C.:

analysis and interpretation of data; administrative, technical, or

material support; critical revision of the manuscript for important

intellectual content; and statistical analysis. P.A.H.: analysis and

interpretation of data; administrative, technical, or material support;

drafting of the manuscript; and critical revision of the manuscript for

important intellectual content. S.G.K., L.F.L., M.L.R., W.F., E.L.F.,

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administrative, technical, or material support; critical revision of the

manuscript for important intellectual content. D.L.C.: analysis and

interpretation of data; administrative, technical, or material support;

drafting of the manuscript; and critical revision of the manuscript for

important intellectual content. Z.C.-M.: study concept and design;

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Study Highlights

WHAT IS KNOWN

- ✓ Chronic pancreatitis (CP) is an inflammatory disease of the pancreas with significant morbidity and mortality.
- ✓ There are no diagnostic biomarkers for CP.
- ✓ The pathophysiology of CP remains poorly understood.

WHAT IS NEW HERE

- ✓ Plasma neutrophil gelatinase-associated lipocalin (NGAL) in combination with clinical characteristics is a potential biomarker for CP.
- ✓ NGAL⁺ immune cells associated with CP that suggest a shift in the adaptive immune response in CP.
- ✓ NGAL and its binding partner linoleic acid are altered in subjects with CP and diabetes.

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