



# Glycosylation alterations in acute pancreatitis and pancreatic cancer: CA19-9 expression is involved in pathogenesis and maybe targeted by therapy

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## Introduction

Dear editor, we have read with great interest the basic research paper entitled ‘*The glycan CA19-9 promotes pancreatitis and pancreatic cancer in mice*’, published recently by Engle *et al.* in the *Science* journal (1). This paper underlines that CA 19-9 is not only the most used and well-known tumor biomarker for pancreatic cancer (2), but an active player in the acute pancreatitis and pancreatic cancer pathogenesis, and a potential target for future therapies (1).

For the present editorial we have scrutinized the PubMed/Medline and Web of Science databases, using the search strategy detailed in *Table 1*. The semantic analysis of the abstracts retrieved in Web of Science may be observed in *Figure 1*, with inter-relations between ‘CA 19-9’ ‘expression’, primary ‘tumor’, ‘borderline’ and ‘metastatic’ tumors.

Pancreatic cancer is associated with an increasing burden between 2007 and 2017, with an increased mortality and disability-adjusted life years (DALYs) during this decade of 24.9% and 21.6%, respectively (3). Similar patterns were observed for pancreatitis, with 17.6% increase in mortality and 13.0% increase in DALYs during the same time interval (3). Acute pancreatitis presented in

2017 an incidence and prevalence of 1300.9 and 118.6 per 1,000, respectively (4).

The authors investigated the consequences of glycosylation alterations with subsequent CA19-9 elevation in animal models (mouse) and pancreatic organoids. The expression of human FUT3 and  $\beta$ 3GALT5 genes in murine cells generated similar profiles with human CA 19-9 carriers. Induction of CA19-9 expression in mice produced histological signs of acute pancreatitis, such as interstitial edema, lymphocyte infiltrations, and collagen deposition. After 28 days of continuous CA19-9 expression, the disease progressed to chronic pancreatitis, with acinar atrophy, metaplastic ductal cells, and persistent fibro-inflammation (*Figure S1*) (1). FUT3 expression was associated with removal of the terminal galactose moieties present in rodents but not in humans, and increased levels of Lewis antigens (1).

Glycosylation is a post-translational process, through which carbohydrate molecules or ‘glycans’ are added to the cell membrane proteins and lipids, with impact in cellular adherence and mobility (5). In patients with pancreatic cancer were observed significant alterations of sialyl Lewis A (CA 19-9), but also of other important glycans, such as sialyl Lewis X antigen (sLex), truncated O-glycans (Tn

**Table 1** Search strategy used in PubMed/Medline and Web of Science

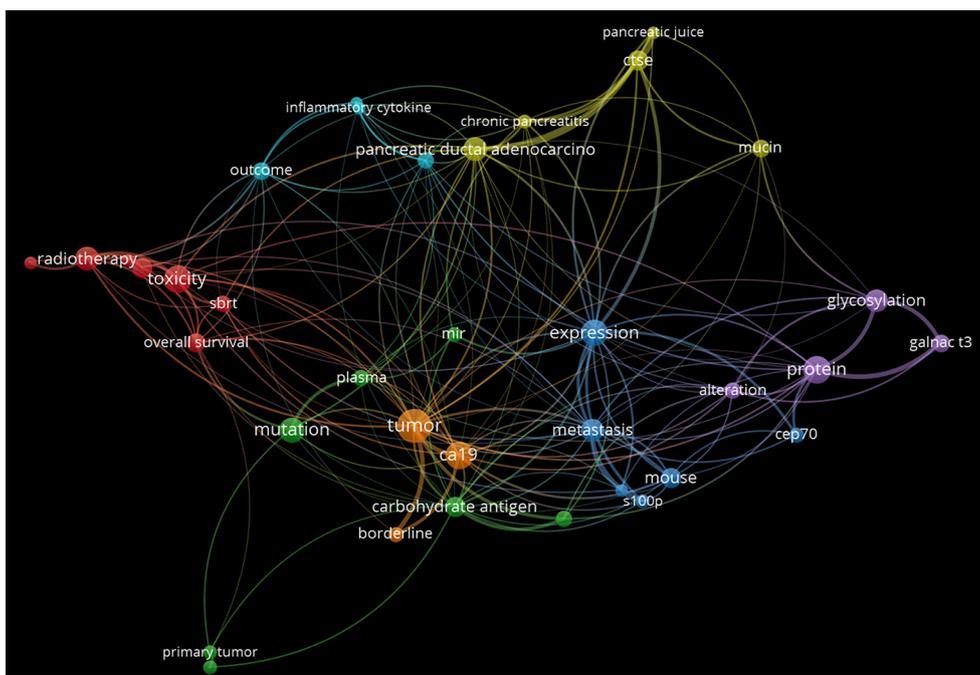
Search	Query	Items found
PubMed/Medline		
#6	Search ((((((pancreatic cancer[Title/Abstract] OR pancreatic ductal adenocarcinoma[Title/Abstract])) OR pancreatitis[Title/Abstract])) AND (((Glycosylation alterations[Title/Abstract] OR CA 19-9[Title/Abstract]) OR carbohydrate antigen 19-9[Title/Abstract] OR sialyl-Lewis[Title/Abstract])) AND ((pathogenesis[Title/Abstract] OR targeted therapy[Title/Abstract]))	8
#5	Search (pathogenesis[Title/Abstract] OR targeted therapy[Title/Abstract])	442,904
#4	Search (((Glycosylation alterations[Title/Abstract] OR CA 19-9[Title/Abstract]) OR carbohydrate antigen 19-9[Title/Abstract] OR sialyl-Lewis[Title/Abstract])	5,786
#3	Search (((pancreatic cancer[Title/Abstract] OR pancreatic ductal adenocarcinoma[Title/Abstract])) OR pancreatitis[Title/Abstract])	92,229
#2	Search pancreatitis[Title/Abstract]	57,123
#1	Search (pancreatic cancer[Title/Abstract] OR pancreatic ductal adenocarcinoma[Title/Abstract])	38,636
Web of Science		
#6	#5 AND #4 AND #3	55
#5	#2 OR #1 <i>Indexes=SCI-EXPANDED, SSCI, A&amp;HCI, CPCI-S, CPCI-SSH, BKCI-S, BKCI-SSH, ESCI, CCR-EXPANDED, IC Timespan=All years</i>	130,655
#4	TOPIC: (pathogenesis) OR TOPIC: (targeted therapy) <i>Indexes=SCI-EXPANDED, SSCI, A&amp;HCI, CPCI-S, CPCI-SSH, BKCI-S, BKCI-SSH, ESCI, CCR-EXPANDED, IC Timespan=All years</i>	671,302
#3	TOPIC: (Glycosylation alterations) OR TOPIC: (CA 19-9) OR TOPIC: (carbohydrate antigen 19-9) OR TOPIC: (sialyl-Lewis) <i>Indexes=SCI-EXPANDED, SSCI, A&amp;HCI, CPCI-S, CPCI-SSH, BKCI-S, BKCI-SSH, ESCI, CCR-EXPANDED, IC Timespan=All years</i>	9,871
#2	TOPIC: (pancreatitis) <i>Indexes=SCI-EXPANDED, SSCI, A&amp;HCI, CPCI-S, CPCI-SSH, BKCI-S, BKCI-SSH, ESCI, CCR-EXPANDED, IC Timespan=All year</i>	56,273
#1	TOPIC: (pancreatic cancer) OR TOPIC: (pancreatic ductal adenocarcinoma) <i>Indexes=SCI-EXPANDED, SSCI, A&amp;HCI, CPCI-S, CPCI-SSH, BKCI-S, BKCI-SSH, ESCI, CCR-EXPANDED, IC Timespan=All years</i>	79,292

and sTn), N-glycans, proteoglycans and galectins, or O-GlcNAcylation (6).

This changing of the tumor cells surface increases cell mobility, with extravasation of the circulating tumor cells into tissues using mechanisms similar with leucocytes populating an inflammatory situs (5). Glycans, such as sialyl Lewis X and sialyl Lewis A, significantly interact with the function of E-selectin ligand, transmembrane adhesion molecules expressed on the surface of vascular endothelial cells (5).

In acute pancreatitis (AP), the genetic background may amplify the deleterious effects of predisposing

environmental etiological factors (7). The involved genetic pathways may be or not pancreas specific, and activated in the early or late-phase of the disease. In the early stage of the disease are involved the *PRSS*—cationic trypsinogen, *SPINK1*—serine protease inhibitor Kazal type 1, *CTRC*—*chymotrypsin C* (in premature trypsinogen activation) and interleukin genes, antioxidant enzyme genes, ACE genes—angiotensin-converting enzyme, *MIF*—migratory inhibitory factor, *iNOS*—inducible nitric oxide synthase, *COX-2*—cyclooxygenases 2, *MYO9B*—myosin IXB (in NF- $\kappa$ B activation). In the late stage of the disease, genetic pathways related to severity are *TLR* genes—toll-like receptors,



**Figure 1** Text analysis of the title and abstracts of papers retrieved in the Web of Science database, using the search strategy from *Table 1*, using the VOSviewer software. Maybe observed important inter-relationships between ‘CA 19-9’ ‘expression’, primary ‘tumor’, ‘borderline’ and ‘metastatic’ tumors as words in the abstracts of the retrieved papers.

*CD14*, *MCP-1*—monocyte chemoattractant protein-1, *HBD* genes—human  $\beta$ -defensin 2, and *MBL2*—mannose-binding lectin 2. The complications seems to be related to *TNF- $\alpha$*  genes—tumor necrosis factor- $\alpha$ , *IL-10*—interleukin 10, and *TLR-4*—toll-like receptor 4 (7).

On the other hand, the epigenetic regulatory mechanisms play an important role in the control of the inflammatory process in AP, with a rapid and limited spatial increase in the H3K14ac, H3K27ac, H4K5ac and (8).

The micro-RNAs with decreased concentration in AP were miR-92b, miR-10a, and miR-7, while increased concentrations were found for miR-126-5p, miR-148-3p, miR-216a-5p, miR-551b-5p, miR-375 (in patients with severe AP), and miR-216a-5p, miR-551b-5p, and miR-375 in mild AP (9). The miR-21 is an important biomarker for patients with pancreatic cancer (10), but seems to modulate also the inflammatory response during AP, through upregulation of *Pias3* and downregulations of *Hmgb1* (11).

Engle *et al.* revealed increased CA19-9 expression in more than 93% of the resected chronic pancreatitis human specimens, especially at the ductal level and less significant in the acinar compartments (1). Loncle *et al.* revealed that transition from chronic pancreatitis to pancreatic cancer

is promoted by the *Il 17*, using as a downstream pathway *REG3 $\beta$ -JAK2-STAT3* inflammatory pathway (12).

CA19-9 expression is associated with hyperactivation of the epidermal growth factor receptor (EGFR) signaling. CA19-9 expression was associated with elevated levels of endogenous fibulin-3 (FBLN3), a glycoprotein with five EGF-like domains, which activated the EGFR pathway (1). Administration of antibodies directed against CA19-9 reduced the inflammatory process of the pancreas, the serum amylase and lipase levels, and decrease the hyperactivation of the EGFR pathway (1). Given the presence of *Kras* gene mutation in more than 90% of patients with pancreatic cancer, the role of EGFR, Ras/Raf/MEK/ERK, PI3K/PTEN/Akt/mTORC1/GSK-3, Janus kinase/Signal Transducer and Activator of Transcription pathways were extensively studied for a better understating of disease pathogenesis (13). The EGFR family of receptors are significantly involved in malignant transformation processes, such as prevention of apoptosis, drug resistance, cancer stem cells and metastasis (13,14). Increasing evidence supports that EGRF signaling is involved the metaplasia process, which converts the acinar cells in progenitor-like ductal cells. This process of acinar to ductal metaplasia may be observed in

pancreatic cancer initiation in oncogenic *Kras* and patients with chronic pancreatitis (15,16). EGFR inhibitors, such as gefitinib (Iressa) and erlotinib (Tarceva) block the epithelial-to mesenchymal transition, decreasing the metastatic potential of pancreatic cancer cells (14). Some drugs that sensitize the pancreatic cancers cells to EGRF inhibitors were described, such as rhein (17) or alantolactone (18). Inhibition of PI3K $\gamma$  significantly reduced acinar cells injury and necrosis in a murine model of AP (19). Karki *et al.* revealed that *Mist1* gene transcription and protein accumulation were significantly reduced during the process of acinar-to-ductal metaplasia found in patients with AP (20). Constitutive mice with expression of Cre-inducible *Mist1* transgene (*iMIST1*) during AP presented a dramatic increase of organ injury secondary to acinar cells death (20).

The CA19-9 expression in mice also harboring *Kras* oncogene, generates aggressive forms of pancreatic cancer, with anaplastic primary tumors and widespread metastases, associated with decreased median survival (202 versus 460 days) (1). There is a cross talk between Ras and EGFR downstream pathways, which seems to have important role in driving metastasis (13).

The present evidence revealed engineered anti-CA 19-9 antibodies, produced with the aim to specifically target pancreatic tumors and increase the accuracy of imagistic methods (21-23). In a murine study, the authors developed three specifically antibodies ( $^{89}\text{Zr}$ - $^{88}\text{DFO}$ -5B1,  $^{88}\text{FL}$ -5B1, and  $^{89}\text{Zr}$ - $^{88}\text{dual}$ -5B1), directed against CA 19-9 for Positron Emission Tomography (PET), near-infrared fluorescent optical imaging and multimodal imaging of pancreatic cancers (24). Currently, a phase 1 study evaluates  $^{89}\text{Zr}$ -DFO-HuMab-5B1 (MVT-2163) and HuMab-5B1 (MVT-5873) for tumor imaging using PET scanning in patients with pancreatic cancer and other CA 19-9 positive malignancies (25).

Targeted therapies using  $^{225}\text{Ac}$ -labeled tetrazine radioligand and a Trans-cyclooctene-bearing anti CA 19-9 antibodies (5B1) were described for  $\alpha$ -radioimmunotherapy of pancreatic ductal adenocarcinoma, reducing hematotoxicity while maintaining the therapeutic effects (26).

Human monoclonal antibodies to sialyl-Lewis<sup>a</sup> were generated and characterized from blood lymphocytes of people immunized with sLe<sup>a</sup>-KLH vaccine (27). The 5B1 and 7E3 antibodies increased the median survival of animals engrafted with Colo205 tumor cells. Treatment with 5B1 antibodies cured 40–60% of mice, while the mortality was 100% in untreated mice within 155 days. Both antibodies presented increased activity through a

complement-dependent cytotoxicity mechanism; the 5B1 antibody presented also an increased antibody-dependent cytotoxicity (27).

In conclusion, further basic and clinical research is needed in the area of pancreatology, due to the severity of pancreatic diseases and their increased morbidity and mortality. The recent findings revealed the genetic pathways and epigenetic factors involved in pathogenesis of acute pancreatitis, chronic pancreatitis and pancreatic cancer, offering the hope for more precise and effective treatments in patients with pancreatic diseases.

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## Footnote

*Conflicts of Interest:* The authors have no conflicts of interest to declare.

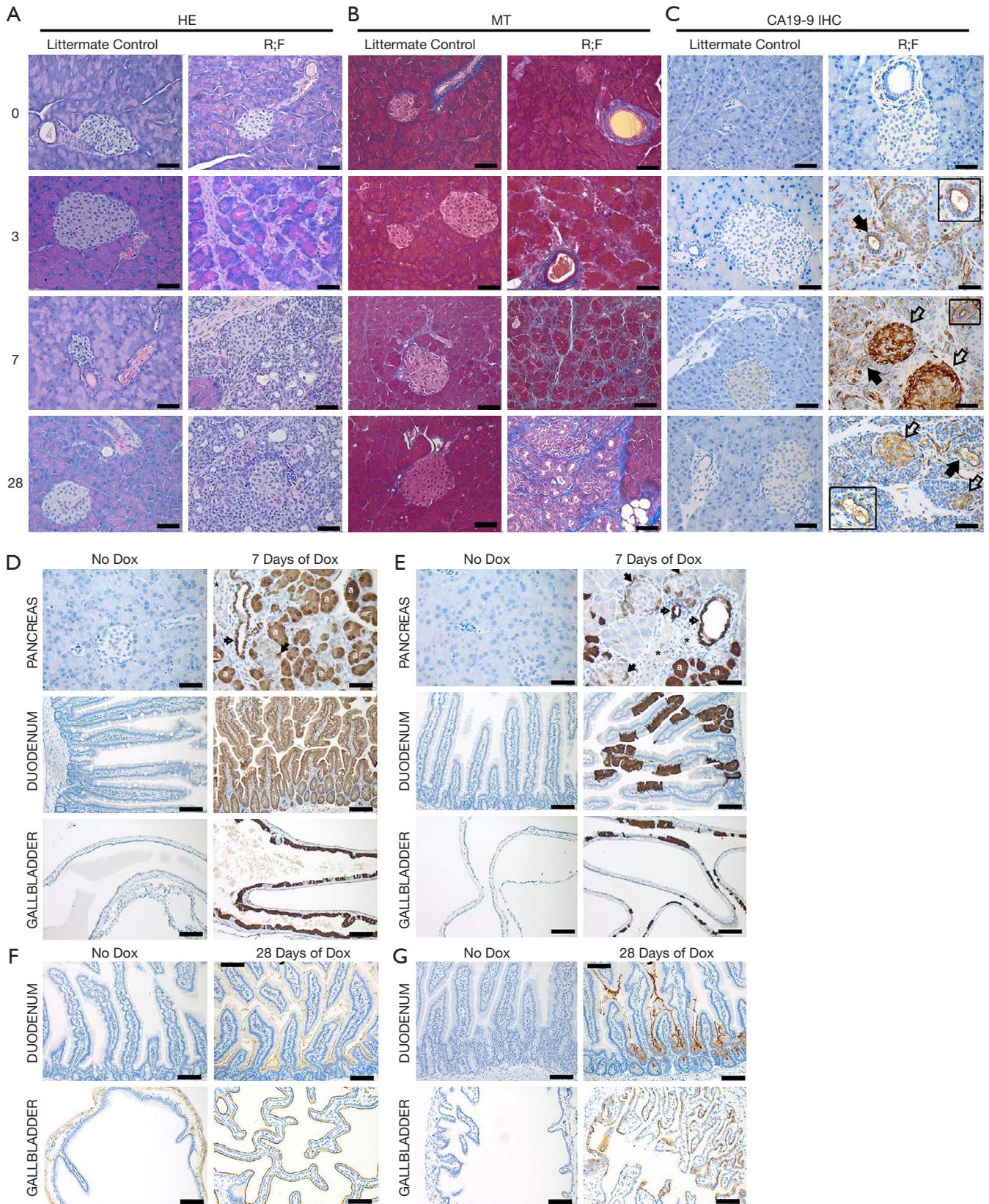
*Ethical Statement:* The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

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**Figure S1** Mouse model comparison for pancreatitis. (A) H&E, (B) Masson's Trichrome (MT), and (C) CA19-9 IHC of mouse pancreata from genetically negative control littermates and R;F, mice treated with Dox. Closed arrows indicate ducts and open arrows indicate islets. eGFP IHC on untreated and 7-day Dox-treated (D) R;F and (E) C;RLSL;F mouse pancreata, duodenum, and gall bladder. Closed arrows indicate positive islet cells, open arrows indicate positive ducts, \* indicates negative vessels and "a" indicates positive acinar cells. CA19-9 IHC on untreated and 28-day Dox treated (F) R;F and (G) C;RLSL;F mouse duodenum and gallbladder. Scale bars = 50  $\mu$ m [From ref (1)].