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Aldehyde “Adduction” Explains Synergy of Smoking and Alcohol in Promoting Pancreatitis



See “Malondialdehyde-acetaldehyde extracellular matrix protein adducts attenuate unfolded protein response during alcohol and smoking-induced pancreatitis” by Bhatia R, Thompson CM, Clement EJ, et al, on page 1064.

Pancreatitis is the inflammatory disorder of the pancreas, which often forms a disease continuum starting with a so-called sentinel attack of acute pancreatitis (AP), followed by recurrent AP (RAP) episodes, and eventual progression to chronic pancreatitis (CP).¹ The risk of

recurrence after the first AP attack is ~10% to 30%, and the risk of progression from RAP to CP is ~25% to 35%.^{2–4} Progression after the sentinel episode of AP to RAP and ultimately to CP is often associated with environmental or genetic risk factors, or both.^{3,5,6}

The most common environmental risk factor is alcohol, which is responsible for 43% to 76% of pancreatitis etiology.^{3,7} Smoking has been identified as an independent risk factor responsible for 61% of pancreatitis cases, and it can also amplify the hazardous effects of alcohol.^{3,7–10} The mechanisms by which these risk factors synergize are currently being explored.¹¹ In a 2017 study, Lugea et al¹² demonstrated that cigarette smoke together with alcohol

contribute to CP pathogenesis through endoplasmic reticulum (ER) stress in pancreatic acinar cells. In this issue of *Gastroenterology*, findings of Bhatia et al¹³ provide an even closer mechanistic link between chronic ethanol and smoke exposure and ER stress-induced pancreatitis.

Activation of ER stress mechanisms has been recognized as a key pathologic event in both the development of AP and progression to CP.^{14,15} During ER stress, cells initiate the unfolded protein response to cope with accumulating unfolded or misfolded proteins, or both, in the ER. Main modules of the unfolded protein response include the (1) protein kinase RNA-like ER kinase (PERK), the (2) activating transcription factor 6 (ATF6), and the (3) inositol-required enzyme 1 (IRE1), which are localized to the ER membrane and become activated during ER stress. Among these 3 distinct branches, alcohol activates the IRE1 arm, leading to higher levels of the spliced form of the transcription factor X-box binding protein 1 (XBP1) in the cytoplasm. Spliced XBP1 then translocates to the nucleus and increases the expression of chaperone proteins as part of the adaptive response. Continued ER stress and a defective XBP1 response activate the PERK branch, which leads to the upregulation of the proapoptotic transcription factor cytidine-cytidine-adenosine-adenosine-thymidine (CCAAT)-enhancer-binding protein homologous protein (CHOP, also known as DNA damage-inducible transcript 3 [DDIT3]) and results in cell death. Toxic chemicals in the cigarette smoke extract were shown to promote cell death via dysregulation of the unfolded protein response.^{16,17} Lugea et al¹² discovered that cigarette smoke reduced spliced XBP1 protein expression levels, thereby suppressing the adaptive response to alcohol, which resulted in sustained PERK activation and increased CHOP expression, followed by acinar cell death and pancreatitis. Nevertheless, the exact molecular mechanism leading to the downregulation of spliced XBP1 remained unclear.

To tackle this knowledge gap, in the present study, Bhatia et al¹³ performed proteomic analysis on a mouse model of secretagogue-induced RAP/CP with chronic alcohol and cigarette smoke extract exposure. They identified stable malondialdehyde-acetaldehyde adducts on the deposited extracellular matrix proteins in the pancreas of mice with RAP/CP and in pancreatic samples from patients with CP. Curiously, they found that the stable malondialdehyde-acetaldehyde adducts generated by alcohol and smoking significantly inhibited XBP1 expression, leading to unresolved ER stress and acinar cell death. In addition, they observed that stable malondialdehyde-acetaldehyde adducts lingered after the cessation of alcohol and smoking and delayed pancreatic recovery by suppressing the expression of cyclin-dependent kinases, cell cycle-associated proteins, and regeneration markers.

Cigarette smoking and alcohol consumption are the main sources of acetaldehyde and malondialdehyde exposures. With respect to smoking, acetaldehyde is formed from the pyrolysis of tobacco, and malondialdehyde is the by-product of interaction of cellular lipids with cigarette smoke.¹⁸ Similarly, alcohol metabolism yields acetaldehyde via the

action of alcohol dehydrogenase and produces malondialdehyde through lipid peroxidation.¹⁸ High concentrations of acetaldehyde and malondialdehyde react with proteins and form hybrid adducts. Stable malondialdehyde-acetaldehyde adducts were previously shown to stimulate proinflammatory responses in liver¹⁹ and lung¹⁸ diseases; however, this is the first report on their role in the context of pancreatitis. Although not studied directly, the authors speculate that malondialdehyde-acetaldehyde adducts on extracellular matrix proteins activate scavenger receptors, thereby leading to acinar cell death and fibroinflammatory responses.

Alcohol abuse often concurs with cigarette smoking, and >80% of patients with alcohol-induced CP have been identified as smokers.²⁰ Because the organ damage caused by these substances is reversible to a certain extent, participating in cessation programs ought to be a straightforward and immediate response in disease management. On the other hand, the present study indicates that pancreatic regeneration is significantly delayed long after the cessation of exposure to cigarette smoke extract and alcohol. Thus, therapeutic interventions should aim at the prevention of pancreatic damage and specifically inhibit pathways activated by the pancreatitis-causing toxic substances. In this regard, the discovery of Bhatia et al¹³ identifies malondialdehyde-acetaldehyde adducts as novel therapeutic targets in the prevention and treatment of pancreatitis associated with alcohol abuse and smoking.

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Needle-free Nonalcoholic Fatty Liver Disease Prognostication: Moving One Step Closer



See “Liver stiffness on magnetic resonance elastography and the MEFIB index and liver-related outcomes in nonalcoholic fatty liver disease: a systematic review and meta-analysis of individual participants,” by Ajmera V, Kim BK, Yang K, et al, on page 1079.

The high prevalence, phenotypic diversity in the patient population, variability in disease progression rates, and increased risk of cardiovascular disease makes risk stratification of nonalcoholic fatty liver disease (NAFLD) challenging.¹ Liver fibrosis is the key histological feature predictive of liver-related outcomes in NAFLD.² However, the limitations of liver biopsy, the current gold standard for staging hepatic fibrosis, are well-known.³ Thus, developing effective noninvasive tests (NITs) for disease staging and determination of progression risk has long been a holy grail of the NAFLD field.

Several NITs initially developed for staging chronic hepatitis C infection, such as the Fibrosis-4 index (FIB-4) and vibration controlled transient elastography (VCTE), have been adapted, with different cutoff values, for noninvasive fibrosis assessment in NAFLD.^{4,5} Other metrics, such as the NAFLD Fibrosis Score and BARD score, have been developed specifically for NAFLD and incorporate metabolic risk factors such as diabetes and obesity.^{6,7} However, these NITs of liver fibrosis in NAFLD have several limitations of their own. First, NITs have a high negative predictive value (scores below the low cutoff value), but low positive predictive value (scores above the high cutoff value) for advanced fibrosis. Second, some patients may be classified in the intermediate risk category between the low- and high-cutoff values.⁸ This situation necessitates the use of additional NITs or liver biopsy for disease staging. Thus, there remains a strong clinical need for developing additional tools and risk stratification pathways for determining disease stage and prognosis in NAFLD.