New Insights into the Pathogenesis of Pancreatitis

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Abstract

**Purpose of review**—In this article, we review important advances in our understanding of the mechanisms of pancreatitis.

**Recent Findings**—The relative contribution of intra-pancreatic trypsinogen activation and NFκB activation, the two major early independent cellular events in the etiology of pancreatitis, have been investigated using novel genetic models. Trypsinogen activation has traditionally held the spotlight for many decades as it is believed to be the central pathogenic event of pancreatitis. However, recent experimental evidence points to the role of trypsin activation in early acinar cell damage but not in the inflammatory response of acute pancreatitis through NFκB activation. Further, chronic pancreatitis in the caerulein model develops independently of trypsinogen activation. Sustained activation of the NFκB pathway, but not persistent intra-acinar expression of active trypsin, was shown to result in chronic pancreatitis. Calcineurin-NFAT signaling was shown to mediate downstream effects of pathologic rise in intracellular calcium. IL-6 was identified as a key cytokine mediating pancreatitis-associated lung injury.

**Summary**—Recent advances challenge the long-believed trypsin-centered understanding of pancreatitis. It is becoming increasingly clear that activation of intense inflammatory signaling mechanisms in acinar cells is crucial to the pathogenesis of pancreatitis, which may explain the strong systemic inflammatory response in pancreatitis.

**Keywords**
Trypsin; NFκB; acute pancreatitis; chronic pancreatitis

Introduction

Pancreatitis, whether acute or chronic, is a cause of significant morbidity and mortality(1). There is at this time no targeted treatment for pancreatitis owing to the lack of understanding of its pathogenesis. The notion of pathologic autodigestion by the digestive enzymes synthesized and secreted by the pancreas as a result of premature intra-pancreatic activation, seems a natural possibility and forms the basis of the century-old autodigestion-centered theory of pancreatitis (2, 3) (Figure 1). Concerted research efforts in the 1980s and 1990s focused upon the mechanisms of premature digestive enzyme activation, and these led to significant advances in our knowledge of intracellular events in the initiation and progression of pancreatitis (4–7). Pathologic calcium signaling, subcellular redistribution leading to colocalization of lysosomal and zymogen components, early activation of NFκB pathway, identification of the processes of ER stress and autophagy in acinar cells, effects of redox signaling and oxidative stress, pathologic role of cellular and extracellular pH changes and the role of bide duct epithelial cells during pancreatitis have all been reviewed in our
previous articles (2, 3). In this article, we review important recent advances in the field since
the last review period. These articles are intended to collectively present a comprehensive
discussion of the pathogenesis of pancreatitis.

We present first the developments relating to the existing trypsin-centered theory of
pancreatitis and examine them in detail with a focus on their implications and then discuss
alternative hypotheses.

Following this, we discuss growing evidence supporting a novel paradigm of pancreatitis in
which inflammatory signals from acinar cells trigger intense local and systemic
inflammatory responses.

The trypsin-centered theory

Activation of trypsinogen to trypsin, the first step in the physiologic digestive enzyme
cascade activation that occurs in the duodenum, has been confirmed to occur pathologically
within the pancreas in most experimental models of pancreatitis (6–10). A plausible
scenario, which is implicitly assumed by the trypsin-centered theory of pancreatitis (Figure
1), is that premature trypsinogen activation is directly pathogenic in pancreatitis (2, 8).

However, several other possibilities exist (8): the detectable premature trypsinogen
activation – 1) may not be pathologically significant quantitatively, or 2) may be adequately
cleared by protective responses within the acinar cell, or 3) may trigger pathologic acinar
cell inflammatory signaling without direct cell damage, or 4) may cause some direct cell
damage, trigger protective and pathologic cellular pathways such as autophagy and ER
stress, and activate pathologic inflammatory signaling, or as a few experts have suggested
(5, 11) may even be a protective response. Alternatively, other potential important trypsin-
independent pathologic events may be present in sufficient degree to cause pancreatitis
independently or in conjunction of trypsin-related injury (12). Thus, the causality of
premature trypsinogen activation and pancreatitis cannot be an obvious assumption without
rigorous proof. Experiments using protease inhibitors, with their questionable target
specificity in suppressing trypsin activity, failed to provide any resolution (13, 14).
Fortunately, genetic models specifically lacking pathologic trypsinogen activation (such as
mice lacking trypsinogen-7 (T−/−) (12) or cathepsin B (CB−/−) (15) have now emerged,
permitting the design of exciting experiments that permit unambiguous reexamination of the
trypsin-centered theory of pancreatitis.

Premature trypsinogen activation and early acinar injury in acute pancreatitis

The extent of acinar necrosis was reduced by half in T−/− mice during caerulein-induced
acute pancreatitis (12). Similar results had previously been seen in CB−/− mice (15).

However, local and systemic inflammation comparable to that seen in wild type mice (who
do have pathologic trypsinogen activation (12, 15)) developed in the absence of significant
pathologic trypsinogen activation in both T−/− and CB−/− mice. Thus, premature
trypsinogen activation seems to contribute to a component of local injury but inflammation,
both local and systemic, which is the hallmark of acute pancreatitis responsible for high
morbidity and mortality, progresses independently of trypsinogen activation during acute
pancreatitis (3, 12).

In vitro studies using T−/− acini showed no induction of cell death with supramaximal
caeerulein stimulation, which induces necrosis in acini from wild type mice (12). However,
apoptosis occurred with transfection-mediated intra-acinar expression of active trypsin (in
mouse acini) (16) or of mutated trypsinogens (in rat AR42J cells) (17). Thus both necrosis
and apoptosis are induced by active trypsin, but the mechanism is not known although
involvement of lysosomal membrane permeability has been postulated (Saluja A,
unpublished observations). Both caspase-dependent and -independent mechanisms have been demonstrated (16).

However, trypsin-induced acinar cell death in experimental conditions does not by itself validate the belief that premature trypsinogen activation is sufficient to induce the systemic complications of acute pancreatitis. In an interesting mouse study using inducible expression of active trypsin specific to pancreatic acini, only rapid inductions with maximal expression (in homozygotes) resulted in acute pancreatitis, while gradual repetitive inductions or milder expression (in heterozygotes) were not sufficient (18). Thus, it is apparent that not all detectable trypsin activity necessarily implies pathogenicity. However, most experimental pancreatitis studies continue to mechanistically link the pathologic events being investigated to detectable trypsinogen activation (19–21).

The response of pancreatic acini to trypsinogen activation seems to be rapid induction of cell death, and this may be a primarily protective response (18). Nonetheless, when overwhelming, as in caerulein-induced pancreatitis, cell death resulting from premature trypsinogen activation accounts for the early pancreatic injury during acute pancreatitis that is notably lacking in T−/− and CB−/− mice (12, 15) (Figure 2).

**Trypsin and pancreatitis susceptibility genes**

The biggest support for the trypsin-centered theory comes from identification of mutations in the cationic trypsinogen gene *PRSS1* found in hereditary pancreatitis (22), an uncommon form of pancreatitis with autosomal dominant inheritance. Over the last decade, several additional mutations have been associated with familial as well as alcoholic and non-familial forms of chronic pancreatitis (23–26). Five pancreatitis susceptibility genes have now been well established: 1) cationic trypsinogen (*PRSS1*), 2) cystic fibrosis transmembrane conductance regulator (*CFTR*), 3) pancreatic secretory trypsin inhibitor or serine protease inhibitor kazal type 1 (*SPINK1*), 4) chymotrypsinogen C (*CTRC*), and 5) calcium-sensing receptor (*CASR*) (23, 26).

Increased trypsinogen activation resulting from the recognized mutations in these susceptibility genes has been postulated to be the mechanism of pancreatic injury (23–26). This hypothesis is the most plausible mechanistic explanation, but it lacks any direct evidence, as is well illustrated by investigations on *PRSS1*, the best-studied among these genes. Increased trypsinogen activation has been demonstrated in the test tube for only a few of the *PRSS1* mutations (27, 28), but their expression in human embryonic kidney (HEK) cell lines suggests that intracellular trypsinogen activation may not occur (29). No evidence of increased trypsinogen activation was found in mice expressing the most common p.R122H *PRSS1* mutant (30). It has also been recognized that, in a few cases, these mutations may be incidental (31, 32). Further the trypsinogen activation hypothesis of hereditary pancreatitis does not explain incomplete penetrance, intermittent nature of the disease, and lack of progression to CP in some individuals despite recurrent episodes (33–35). However, emerging epidemiologic and genetic data continue to be linked to trypsinogen activation (23, 25, 36–39). On the contrary, it is becoming increasingly clear that with the exception of hereditary pancreatitis and cystic fibrosis, a direct simplistic genetic mechanism may not exist but that instead, a complex interplay between genetic, environmental and developmental factors governs the susceptibility, progression and severity of pancreatitis (23, 40).

**Premature trypsinogen activation and chronic pancreatitis**

Traditionally, the pathogenesis of chronic pancreatitis has been mostly deduced as a natural extension of acute pancreatitis. Despite the tendency to clump these entities together,
assumptions from acute pancreatitis are speculative at best, since the relationship between acute and chronic pancreatitis is not clear (33, 41–44). Nevertheless, as for acute pancreatitis, the popular belief persists that intra-pancreatic trypsinogen activation is responsible for the pathogenesis of chronic pancreatitis. This theory is primarily supported by the observation that PRSS1 mutations of hereditary pancreatitis, and as described earlier, bear a complementary relationship with the proposed trypsinogen activation theory of other pancreatitis susceptibility genes (24, 26, 45).

Interestingly, prolonged expression of active trypsin in mouse pancreatic acinar cells in an in vivo genetic model resulted in acinar cell loss but was not sufficient to cause chronic pancreatitis (18). Conversely, T−/− and CB−/− mice lacking pathologic intra-acinar trypsinogen activation developed chronic pancreatitis indistinguishable from that in wild type mice, suggesting that the pathogenesis of chronic pancreatitis is independent of trypsinogen activation (46). These data, although limited by lack of validation in multiple experimental models, challenge the existing trypsin-centered theory of chronic pancreatitis. Further confirmation of the results just described awaits the development of alternative experimental mouse models of chronic pancreatitis where existing genetic manipulations such as knockout of trypsinogen-7 can be applied.

Inflammatory signaling in acinar cells

Pancreatic acinar cells are unique in the human body - they have the highest rate of protein synthesis and produce large quantities of digestive enzymes (47). Conceivably, these cells have unique properties to maintain their functionality, stability and self-protection. One of these seems to be their inflammatory properties (3). Emerging evidence from acinar cells points to early and sustained activation of inflammatory signaling that is responsible for the intense local and systemic inflammatory response seen in acute pancreatitis (Figure 2) and the development of the chronic inflammation and fibrosis of chronic pancreatitis (Figure 3). The NFκB pathway is the best described inflammatory signaling pathway in the pancreatic acinar cell (48).

Activation of NFκB in acinar cells

Though the activation of NFκB within the acinar cell during pancreatitis had been previously recognized (49–51), it is only recently that its significance is becoming apparent. The matter has been debated as a potential consequence of premature trypsinogen activation for over a decade (2, 52), but recent data from T−/− and CB−/− mice (12) and in vitro expression of active trypsin within acini using transfection (16) confirm that intra-acinar NFκB activation occurs very early in pancreatitis independent of trypsinogen activation. Further, activation of NFκB in acinar cells has been shown to result in severe acute pancreatitis with local injury and systemic inflammatory response in several genetic models including adenovirus-mediated transfection of activated NFκB in the acinar cells by intraductal injection (53) and in two separate models of inducible expression of active IKK2 targeted to pancreatic acinar cells (54, 55). In addition, inhibition of NFκB in genetic models, e.g. mice with constitutional and global deletion of NFκB p50 subunit (56) and pancreas-specific expression of inducible inactive IKK2 (54), as well as pharmacologic studies (48, 57, 58), have been shown to result in attenuated pancreatitis response with severity inversely proportional to the degree of NFκB inhibition. In another recent genetic model, pancreas-specific inducible expression of NFκB p65/RelA subunit led to compensatory expression of IkBα leading to unclear phenotype at baseline but increased severity of acute pancreatitis with even mild caerulein stimulation (55). When this model was combined with inducible pancreas-specific expression of active IKK2 that resulted in inducible p65 and active IKK2 co-expression in the pancreas (55), spontaneous severe acute
pancreatitis was the result. This highlights the role of induction of the NFκB pathway in causing acute pancreatitis.

**Sustained intra-acinar NFκB activation in chronic pancreatitis**

Sustained activation of NFκB in the acinar cells was seen in the caerulein model of chronic pancreatitis in WT as well as T−/− and CB−/− mice lacking significant intra-acinar trypsinogen activation (46). Chronic intra-acinar NFκB activation was further confirmed in human chronic pancreatitis (46). Based on these data, it was suggested that sustained activation of inflammatory pathways in pancreatic acinar cells results in chronic pancreatitis (46) (Figure 3).

Recent experimental data support this theory of sustained intra-acinar NFκB activation. Inducible expression of NFκB p65/RelA in pancreatic acini led to induction of severe chronic pancreatitis within 6 weeks of weekly episodes of caerulein-induced mild acute pancreatitis (55). These repeated episodes of mild acute pancreatitis failed to result in chronic pancreatitis in the wild type mice (55). Further, the prolonged expression of inducible active IKK2 in pancreatic acini resulted in spontaneous severe chronic pancreatitis (55).

Persistence of the stimulus, which may be any of the recognized etiologies (such as alcohol, bile exposure, increased ductal pressure, intracellular accumulation of mutated proteins, autoimmune sensitization, etc), may drive the sustained intra-acinar activation of the inflammatory pathways in chronic pancreatitis (46). In line with this theory, it has been demonstrated in alcohol-related pancreatic injury that withdrawal of the stimulus may lead to disease reversal at least in the early stages (59). A complex interplay of genetics, environmental and structural factors may influence the effects of the pathogenic stimuli (46). This novel paradigm of chronic pancreatitis, contrary to the prior theories, does not require trypsinogen activation, necrosis-fibrosis sequence, acute-to-chronic pancreatitis progression or a sentinel episode of acute pancreatitis (33, 41, 42, 45).

**Constitutive vs inducible NFκB model paradox**

In contrast to observations in the inducible models of NFκB described above (with the exception of one constitutive mouse model with global deletion of p50), Algul et al (21, 60, 61) report paradoxical results in models with constitutive NFκB genetic modifications.

Pancreas-specific truncation of p65/RelA led to increased severity of acute pancreatitis (60) as well as chronic pancreatitis (61) induced with caerulein. Mice with deletion of IκBα in the pancreas demonstrated constitutive NFκB activation, which attenuated the pancreatitis response, and this effect could be reversed by combined deletion of p65/RelA and IκBα (21). This group has proposed a protective role for NFκB activation in the acinar cell due to induction of pancreatitis associated protein (PAP) and serine protease inhibitor 2A (spi2A, which has been linked to diminished trypsinogen activation) (21, 60).

The authors argue that NFκB activation in the myeloid cells of the inflammatory infiltrate is responsible for inflammation in acute pancreatitis (21, 60) and in fibrosis and stellate cell activation in chronic pancreatitis (61). Interestingly, deletion of p65/RelA in the myeloid cells alone did not lead to any difference in pancreatitis response compared with that in wild type mice (61). While the paradox remains to be definitively resolved, one possible explanation may be disruption of the broad spectrum of NFκB -dependent processes (62, 63) in the pancreas in the constitutive models, which probably remain unaffected or minimally affected in the inducible/conditional models. Such an example was recently demonstrated in pancreatic acinar cells in a constitutive model of IKK1 deletion, which led to reduced autophagic degradation independently of NFκB activity and increased spontaneous
pancreatitis by induction of ER stress and oxidative stress (64). Undoubtedly, the diversity of NFκB-regulated processes and lack of a clear understanding of the fine tuning between physiological and pathological responses (63) lend great complexity to these experimental models.

**Cytokines**

Zhang et al (65) produced a model of severe acute pancreatitis with lung injury by using IL6-deficient mice and a modified protocol of caerulein pancreatitis where mild acute pancreatitis was repeated consecutively for five days. They showed here that IL6 mediates pancreatitis-induced lung injury through complexation with its soluble receptor (sIL6R), a process known as trans-signaling (65). The IL6-sIL6R complex led to STAT3 activation in the pancreatic acini while myeloid NFκB activation induced IL6 production (65). IL22 was found to be protective in acute pancreatitis and identified as a potential therapeutic option (66). Aryl hydrocarbon receptors (AhR) on CD4 cells were required for IL22 production and AhR-deficient mice developed severe pancreatitis which could be rescued by exogenous administration of IL22 (66). Overexpression of IL22 receptors on acinar cells occurred in acute pancreatitis (66).

**Other pathologic mechanisms**

Pathologic calcium signal is generated in pancreatic acinar cells as an immediate response to pathologic stimuli (discussed in detail in our previous reviews (2, 3)). This signal has been shown to be transmitted through calcineurin activation (20). When inhibited pharmacologically and via genetic deletion, calcineurin was shown to reduce bile-induced pancreatitis in mice (20). Further downstream target of calcineurin activation was identified as the family of 4 transcription factors: nuclear factor of activated T-cells (NFAT c1–4). Inhibition of NFATc3 was shown to attenuate pancreatitis (19).

ER stress and activation of unfolded protein response, recently recognized in acute pancreatitis (2, 3, 67), were shown to be sustained in chronic pancreatitis (unpublished observations, Sah and Saluja). Pancreatic acinar cells, owing to their dependence on high ER volume and functionality, are especially susceptible to perturbations in ER homeostasis.

Autophagy and lysosomal dysfunction are being investigated intensively as crucial cellular events in pancreatitis (68). SPINK3/PST1 (mouse correlate of SPINK1 in humans) has been shown to affect autophagy as well as several other processes including inflammation and proliferation (38, 39, 69), which may explain the attenuated pancreatitis observed in mice overexpressing SPINK3/PST1 (70, 71). Retarded progression of autophagy has been linked to pathogenesis of pancreatitis (68, 72), an effect demonstrated recently in numerous models such as pancreas-specific IKK1-deleted mice (64), interferon regulatory factor 2 (IRF 2)-deleted mice (73) and mice overexpressing IL22 (74).

The multiple pathologic cellular events identified in acute pancreatitis, i.e. autophagy, ER stress, oxidative stress, lysosomal and mitochondrial dysfunction may be interdependent and may eventually contribute to sustained acinar cell inflammatory signaling that leads to acute pancreatitis, as described in our prior article (Figure 4 in ref. (3)) (Figure 2).

Pancreatic acinar cells, consequent to the activation of inflammatory signaling, lead to generation of NGF, H⁺ and other neurotrophic factors that activate receptors in the pancreatic afferents such as TRPV1, TRPA1 and PAR2, and this leads to generation of neurokines and bradykinins, which induce neuro-inflammation (75). The role of neuro-inflammation in pancreatitis is not fully characterized apart from generation of pain, although it may lead to augmentation of the pancreatic inflammatory response. Recently,
TRPV1 and TRPA1 channels have been shown to be important in chronic inflammation (76).

**Conclusion**

Genetic models of trypsin expression and models lacking trypsinogen activation have led to exciting results that challenge the century-old trypsin-centered theory of pancreatitis. It is becoming increasingly clear that activation of intense inflammatory signaling mechanisms in acinar cells is crucial to the pathogenesis of pancreatitis, which may explain the strong systemic inflammatory response in pancreatitis.

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**References**


Key points

- Premature trypsinogen activation leads to acinar cell death, which, in turn, contributes to a component of acinar injury in acute pancreatitis.

- Pathogenesis of local and systemic inflammation occurs in acute pancreatitis independently of premature trypsinogen activation, challenging the century-old trypsin-centered theory of pancreatitis.

- Early, sustained intra-acinar NFκB activation is responsible for the intense inflammatory response of acute pancreatitis.

- Chronic pancreatitis in the caerulein model develops independently of intra-acinar trypsinogen activation. Intra-acinar trypsin activation alone is not sufficient to result in chronic pancreatitis.

- Sustained activation of inflammatory pathways by persistent pathogenic stimuli such as NFκB in the acinar cell is responsible for chronic pancreatitis.
Figure 1. The trypsin-centered theory of pancreatitis
Intra-acinar trypsinogen activation is the central event in this theory and is responsible for local injury and systemic inflammation. Other pathologic events like ER stress, autophagy, lysosomal dysfunction, pH alterations, oxidative stress, bile duct dysfunction, etc (see text) are linked to the central event (trypsinogen activation) in this theory.
Figure 2. A schematic of the pathogenesis of acute pancreatitis
Activation of inflammatory pathways such as NFkB in the acinar cell leads to intense inflammatory reaction responsible for local injury and the systemic inflammatory response in acute pancreatitis. Trypsinogen activation seems to cause early acinar damage during acute pancreatitis. Other pathologic events like ER stress, autophagy, lysosomal dysfunction, pH alterations, oxidative stress, bile duct dysfunction, etc (see text) potentially augment and sustain the acinar cell inflammatory signalling leading to the intense inflammatory response.
Figure 3. Pathogenesis of chronic pancreatitis
Sustained activation of inflammatory pathways such as NFκB leads to the development of chronic pancreatitis. The recognized etiological associations of chronic pancreatitis are shown in the box, which may be the pathologic stimulus for sustained activation of acinar cell inflammatory pathways. The effect of these etiological associations seems to be modulated by genetic and environmental factors. See text for details.