Expression of major histocompatibility antigens in human chronic pancreatitis

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Abstract
T-lymphocytic infiltration of the exocrine pancreas and liver in patients with chronic pancreatitis has suggested that cell mediated immune mechanisms may play a part in the pathogenesis of this disease. As expression of major histocompatibility (MHC) antigens is a prerequisite for organ specific autoimmunity, the expression of HLA class I (β2-microglobulin) and class II (HLA-DR) determinants have been analysed, together with the presence of T-lymphocytes, in 93 patients (64 men and 29 women, mean age 40-66 years) having an operation for chronic pancreatitis. Ethanol (63 patients), recurrent acute pancreatitis (12), congenital lesions (2), and unknown (16) were suggested to be the causes of the disease. Immunohistochemical staining of formalin fixed and paraffin wax embedded tissue sections used conventional immunohistochemical techniques with specific anti-serum samples. No MHC expression was identified in 10 historically normal pancreatic control specimens or in four cases of chronic pancreatitis secondary to obstruction by neuroendocrine tumours within the head of the pancreas. β2-microglobulin expression by pancreatic exocrine epithelial cells was seen in 76 chronic pancreatitis specimens (82%) while HLA-DR was present in 61 (66%). Simultaneous expression of both class I and II determinants was seen in 53 (57%) of cases. MHC determinant expression was not found in 10 cases (11%) of chronic pancreatitis. In the positive specimens, expression was confined to ductal and ductular (interlobular and intralobular) epithelium with no staining of acinar cells. Staining was not related to the suspected cause of the disease or age. T-lymphocytes were more prominent in chronic pancreatitis mean (SEM) (131 (15) cells per high powered field) than controls (5 (1), p<0.01). Aberrant MHC expression by exocrine pancreatic epithelial cells occurring in the presence of an appreciable T-cell infiltration confirmed that the appropriate cellular conditions were present for cell mediated cytotoxicity to contribute to the pathogenesis of chronic pancreatitis.

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Chronic pancreatitis is a progressive inflammatory disease characterised by intractable pain, irreversible morphological change, and loss of pancreatic exocrine and endocrine function. The radical operations that may be required to control pain often exacerbate both exocrine and endocrine loss. While alcohol is the most frequently implicated cause, non-alcohol associated (idiopathic) cases account for between nine and 41 per cent of all chronic pancreatitis, the precise incidence depending upon the population studied. Rarer causes include hypercalcaemia, hyperlipidaemia, hereditary factors and, in tropical countries, protein and fat malnutrition. In individual patients, however, the precise cause and pathogenesis often remain unknown. In this report, we explore the possible role of autoimmune mechanisms in the cause and pathogenesis of chronic pancreatitis.

In alcoholic chronic pancreatitis, changes in pancreatic juice may lead to precipitates of protein and calcium, which obstruct the ducts, causing ulceration, fibrosis, and subsequent atrophy of exocrine pancreatic epithelial distal to the obstruction. This hypothesis has evolved after the examination of 'end stage' chronic pancreatitis in which calcium deposition is common, but does not identify the events that start the damage nor does it explain the development of the disease in non-alcoholics. Moreover, many non-alcoholics do not develop chronic pancreatitis. Chronic pancreatitis clearly comprises more than one cause, and Sarles has emphasised the morphological distinction between the chronic calcifying form and 'primary inflammatory pancreatitis.' Klöppel and Mailliet discovered protein plugs and calcification only infrequently in early chronic pancreatitis, suggesting that these are unlikely to be causal factors but rather epiphenomena occurring during progression of the disease.

Previous studies of liver biopsy specimens in patients with severe chronic pancreatitis suggest that immunological mechanisms could play a part in its pathogenesis. Even when alcohol was the suspected cause, hepatic parenchymal damage was minimal compared with the severity of pancreatic disease, showing that alcohol may not be the sole injurious factor. T-lymphocyte infiltration in the absence of hepatocyte damage would be consistent with the transit of inflammatory cells originating within the inflamed pancreas. Autoantibodies reactive with pancreatic acinar cells and ductal antigens have been described in patients with chronic pancreatitis, as have activated lymphocytes specifically sensitised to crude pancreatic antigens, supporting the participation of autoimmune mechanisms in some types of chronic pancreatitis.

During the development of organ specific autoimmunity, initial events entail aberrant expression of major histocompatibility complex (MHC) molecules by cells that do not express such determinants. Class I antigens are expressed by most nucleated somatic cells but not by exocrine pancreatic cells. Normal acinar and ductal pancreatic epithelial cells do not express MHC class II determinants either, but...
some expression has been reported in two small series of patients with chronic or 'obstructive' pancreatitis. Possible autoimmune recognition of pancreatic epithelial cells occurring early in some types of chronic pancreatitis does not exclude alcohol from being an important cause or oppose the hypothesis that proteinaceous and calcific deposits contribute to the general pathogenesis of the disease. Confirmation of autoimmune mechanisms, however, might help define a population of patients who could benefit from non-surgical management of chronic pancreatitis.

Patients and methods

PATIENTS

Pancreatic specimens were reviewed from a personal series (RCNW) of 102 patients having an operation for chronic pancreatitis from 1977 to 1990 at Bristol Royal Infirmary or the Hammersmith Hospital, London. Of these, biopsy samples from nine patients were considered either not to contain sufficient material to permit an accurate histological diagnosis or sufficient epithelial elements to assess immunostaining adequately. These sections were excluded from this study. The 93 patients now reported comprised 64 men and 29 women with a mean age of 40-6 years (range 20-72 years). The main cause of the disease was ethanol (65 patients). Causes in the other 28 patients included previous acute pancreatitis in 12, choledochal cyst in one and pancreas divisum in another. No cause was identified in the remaining 16 patients.

CHRONIC PANCREATITIS SPECIMENS

Wherever possible, pancreatic resection specimens were received fresh and dissected in a standard manner before fixation. Blocks of tissue for examination were taken transversely across the pancreas through cellular regions intermediate between the resection margins and the macroscopically most fibrous and scarred parts of the specimens. These blocks were then bisected vertically to allow examination of the parenchyma both in the regions surrounding the central ducts and at the periphery of the tissue. Blocks taken from densely fibrotic or heavily calcified regions of the specimens were avoided.

CONTROL PANCREATIC SPECIMENS

Ten morphologically normal pancreatic specimens and four specimens of chronic pancreatitis secondary to obstruction by neuroendocrine tumours within the head of the pancreas were used as control tissues. The normal tissues were obtained from patients undergoing distal pancreatic resections for small non-obstructing pancreatic neuroendocrine tumours, patients with suspected chronic pancreatitis but normal pancreatic histology, or where the distal pancreas was excised whole in a radical cancer operation. The tissue blocks were taken transversely across the pancreatic specimens through the regions immediately adjacent to, but at 0-5 cm from, resection margins.

IMMUNOHISTOCHEMISTRY

Class I MHC expression was determined by identification β2-microglobulin, the non-polymorphic light chain component of human class I histocompatibility antigens, using polyclonal rabbit antibody (Dako Ltd, High Wycombe, UK) diluted to 1:1000 in 120 mmol/l sodium phosphate buffered saline (PBS, pH 7-4). Monoclonal mouse anti-human HLA-DR (Dako Ltd), which reacts with the β-chain of all products of the HLA gene subregions DP, DQ, and DR, was diluted to 1:150 in PBS to identify class II MHC expression. Monoclonal antibody UCHL1 (anti-human T cell) was obtained as a gift from Dr Peter Beverley, University College Hospital, London and used at a 1:10 dilution. Human tonsillar tissue was used as the positive control for the three antibodies used.

Formalin fixed and paraffin wax embedded sections (2 μm thickness) were dewaxed in TCF 30 (Infrakem Ltd, Wigan, UK), brought to water through graded ethanols and digested with trypsin (0-1% wt/vol in aqueous 0-1% CaCl2 for five minutes at 37°C). Endogenous peroxidase was blocked with 0-3% hydrogen peroxide (H2O2 in distilled water for 30 minutes at room temperature). A peroxidase anti-peroxidase technique was used to detect bound β2-microglobulin. HLA-DR staining was performed by the avidin-biotin-peroxidase complex method using the Vectastain ABC kit (Vector Laboratories, Burlingame, USA). A conventional indirect immunoperoxidase technique was used for detection of bound HCHL1. In all cases, sections were incubated with primary antibodies overnight at 4°C. The resultant immune peroxidase complexes were developed in 0.5% (vol/vol) 3,3' diaminobenzidine hydrochloride (DAB, Aldrich, Gillingham, Dorset, UK) in PBS containing 0-03% (vol/vol) H2O2. Sections were counterstained with haematoxylin and
magnified in Perox (Histolab, Hemel Hempstead, UK). Expression of α2-microglobulin and HLA-DR were assessed as either being positive or negative in the epithelial elements (acinar, ductular, or ductal), mononuclear cells, and islets. Lymphocyte morphometry was performed using a 10×10 mm graticule within the microscope ocular at ×400 magnification. For each specimen, the number of T-lymphocytes per high power field (hpf) in three areas of maximal infiltration was counted and a mean value calculated.

**STATISTICS**

Statistical analyses were performed using χ2 and unpaired Student’s t tests.

**Results**

**HISTOPATHOLOGICAL ASSESSMENT**

Tissues from the 93 patients included in this study contained the characteristic features of chronic pancreatitis. In all cases, the disease was focal in distribution. Lobulo-acinar destruction was severe in some foci while many adjacent lobules were morphologically unaffected. Islets of Langerhans seemed to be spared during this process and were prominent, particularly in those regions where exocrine epithelial structures had been destroyed. Tissue sections were chosen to show the early rather than later stages of chronic pancreatitis. In these, neither protein plugging of major ducts nor calcification of minute ducts were identified in the specimens included in this study. Sections of endstage chronic pancreatitis comprising only a few residual epithelial elements together with nerves and vascular structures in dense fibrous connective tissue, together with focal calcification, were avoided in the assessment of MHC expression.

**CONTROL IMMUNOHISTOCHEMICAL STUDIES**

No class I or class II MHC expression by exocrine epithelial cells of any of the 10 normal control specimens, or by the four cases of obstructive chronic pancreatitis, was identified. Only a few scattered foci of T-lymphocytes were seen in the interlobular stroma of these specimens.

**CLASS I MHC (α2-MICROGLOBULIN) EXPRESSION**

Of the 93 cases of chronic pancreatitis studied, class I MHC expression was seen in the epithelial elements of 76 (82%) specimens. Expression was focal and varied. Positive staining was confined to intralobular and interlobular ductules (Fig 1) and interlobular ducts; there was no staining identified in acinar cells. In 27 biopsies; this expression affected both ductular and ductal epithelium, while in another 44 cases β2-microglobulin staining was limited to either ductules (28 specimens) or ducts (16 specimens). This pattern of distribution was not assessed in the remaining five biopsies because of the absence of either of these structures. We could not identify a predominance of either a ductular or ductal pattern of β2-microglobulin expression in this series (χ2=3.01, 0.10>p>0.05).

Class I MHC expression was not related to the sites of maximal lymphocytic infiltration, although concurrent β2-microglobulin expression by epithelial cells and adjacent lymphocytes was identified in 10 specimens. In 79 cases (85%), mononuclear cells expressing β2-microglobulin were present, contrasting with only two cases (20%) among controls (p<0.001). In all specimens, both of controls and chronic pancreatitis, islet cells exhibited positive staining for β2-microglobulin. This provided a convenient internal positive control within each tissue section for this antibody.

**Analysis of the data for alcohol intake showed** 50 of 63 patients (79%) with a possible alcoholic cause of the disease to express β2-microglobulin while in the non-alcohol group, 26 of 30 patients with chronic pancreatitis (87%) expressed this determinant. The differences between the groups were not significant (p>0.5).

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**TABLE I**

Comparison of major histocompatibility antigen expression and causes of chronic pancreatitis (n=93)

<table>
<thead>
<tr>
<th>Alcohol</th>
<th>Previous acute pancreatitis</th>
<th>Congenital</th>
<th>Idiopathic</th>
<th>Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>No patients</td>
<td>63 (79%)</td>
<td>10 (83%)</td>
<td>1 (50%)</td>
<td>16 (94%)</td>
</tr>
<tr>
<td>HLA-DR positive</td>
<td>29 (61%)</td>
<td>6 (50%)</td>
<td>1 (300)</td>
<td>14 (83%)</td>
</tr>
</tbody>
</table>

Statistical analysis; *p<0.5 v total, **p<0.1 v total, †no analysis, ‡p<0.05 v total.

**TABLE II**

MHC expression according to age group

<table>
<thead>
<tr>
<th>Age group (yr)</th>
<th>No</th>
<th>β2m positive*</th>
<th>HLA-DR positive</th>
</tr>
</thead>
<tbody>
<tr>
<td>&lt;40</td>
<td>39</td>
<td>31</td>
<td>25</td>
</tr>
<tr>
<td>40-55</td>
<td>44</td>
<td>37</td>
<td>29</td>
</tr>
<tr>
<td>≥55</td>
<td>31</td>
<td>8</td>
<td>7</td>
</tr>
<tr>
<td>Total</td>
<td>103</td>
<td>76</td>
<td>61</td>
</tr>
</tbody>
</table>

*p>0.5 v no in each age group of population studied.
CLASS II MHC (HLA-DR) EXPRESSION

HLA-DR expression was identified in 61 (66%) specimens. This followed a similar pattern of distribution to $\beta_2$-microglobulin. No expression was seen in any acinar cell, positive staining being confined to ductular and ductal epithelium. In 40 cases, HLA-DR was identified in both these structures while expression was restricted to ductules in 12 cases and to ducts in seven cases. Absence of either structure precluded analysis of distribution in the remaining two cases. There was also no relation between areas of HLA-DR expression and foci of lymphocyte infiltration. Concurrent staining of both epithelial elements and adjacent lymphocytes was seen in eight biopsy specimens (Fig 2). Mononuclear cells expressed HLA-DR in 88 (95%) cases compared with five (50%) in controls ($p<0.001$). In five cases, a population of islet epithelial cells and adjacent intra-islet endothelial cells were HLA-DR positive. In these tissues, however, not all the islets contained positive cells.

Analysis of the data for alcohol intake showed 29 of 63 patients (61%) with a possible alcoholic cause of the disease to express class II MHC determinants while in the non-alcohol group, 22 of 30 patients with chronic pancreatitis (73%) expressed this determinant. The differences were marginally significant (0.05 $p>0.02$).

COMBINATION OF CLASS I AND II DETERMINANTS

Simultaneous expression of both class I and II MHC determinants was seen in 53 (57%) cases. Twenty two specimens showed only $\beta_2$-microglobulin staining while eight cases were only HLA-DR positive. There was no relation between either $\beta_2$-microglobulin or HLA-DR expression and the underlying cause (Table I). MHC expression was also not age related (Table II).

CHARACTERISATION OF T-LYMPHOCYTES

T-lymphocytes were identified in all tissue sections studied. These cells were more prominent in chronic pancreatitis specimens, however, with an estimated density mean (SEM) of 131 (15) cells per hpf compared with controls 5 (1), $p<0.01$. The density of lymphocyte infiltration varied among specimens. A wide variety of morphological patterns of T-lymphocyte distribution was identified (Fig 3). The proportion of T-lymphocytes to total mononuclear cells per hpf ranged from 19% to 78%. No direct correlation was seen between T-lymphocyte densities in 38 pancreatic specimens and the periportal regions of corresponding liver biopsy specimens taken at the same operation.11

Discussion

Abnormal expression of class I or class II MHC determinants, or both by exocrine epithelial cells has been found in 89% of cases of chronic pancreatitis. In addition, all cases of chronic pancreatitis contained significantly increased numbers of T-lymphocytes compared with controls. Although no $\beta_2$-microglobulin expression was identified in control exocrine tissues despite positive staining of islet cells, 63% of chronic pancreatitis specimens showed positive interlobular and intralobular ductular staining, not previously described.15,16 Class I determinants are present on all nucleated cells,17 their expression is variable,17 but a detailed analysis of these determinants failed to detect class I HLA-ABC heavy chain antigens in the exocrine pancreas.18 Thus, expression of $\beta_2$-microglobulin in chronic pancreatitis was abnormal and sufficiently enhanced to suggest a state of hyperexpression analogous to that proposed for type I insulin dependent diabetes mellitus.18

In agreement with earlier reports,19,21 this study found that normal pancreatic epithelial cells did not express HLA-DR antigens, yet such expression was seen in two thirds of the chronic pancreatitis specimens. There was simultaneous
expression of class I and II antigens in 53 specimens (57%), while HLA-DR expression alone was rare (9%). We also found a substantial increase in MHC expression by mononuclear cells in chronic pancreatitis specimens and such expression by T-lymphocytes is necessary for the antigen recognition process. Confirmation that these modules are expressed at readily detectable values by pancreatic exocrine ductular epithelial cells, in association with a considerable T-lymphocyte infiltration, establishes the presence of known cellular components required for the start of an autoimmune reaction in a large proportion of patients with chronic pancreatitis.

Participation of immune mechanisms in the cause and pathogenesis of chronic pancreatitis has been suspected as increased serum concentrations of immunoglobulins (IgA, IgM) and changes in concentrations of circulating T-lymphocytes were reported in 1974. Later, Nerenberg et al detected raised values of a circulating antibody monospecific for a pancreatic acinar antigen. Initial events during development of organ specific autoimmunity entail aberrant expression of MHC class II antigens, thus allowing presentation of cell specific antigens to potentially autoreactive T-lymphocytes. These cells recognise protein fragments and synthetic peptides in association with class I and II molecules. Antigens synthesised endogenously by target cells preferentially yield complexes with class I molecules. By contrast, antigens acquired exogenously and taken into target cells by endocytosis preferentially yield class II complexes. While class I antigens participate in antigenic presentation to cytotoxic T-lymphocytes, class II MHC is necessary for similar presentation to helper T-cells. De novo expression of class II MHC (HLA-DR) has been shown in diseases such as primary biliary cirrhosis and type I insulin dependent diabetes mellitus, conditions now considered to be immune mediated.

Our data add to the current understanding of the cause and pathogenesis of chronic pancreatitis. Sarles et al suggested that in the earliest stages, before any cellular damage is detected, there is luminal precipitation of calcium salts and a special pancreatic stone protein. The molecular basis for this process is unknown but could include direct ethanolic toxicity, cellular adaptation to ethanol, or cytotoxic damage to the cell membrane. Each mechanism implies change at a cellular level before the lithogenic process. In a recent histopathological and immunohistochemical study, Suda et al concluded that protein plug obstruction of the ductal system is not the main cause in the genesis of ethanolic pancreatitis, and our morphological findings support this conclusion. Any discrepancies between these findings and those previously reported could reflect a spectrum of different causal factors or studies conducted at different stages in disease progression. In our series, absence of calcification from tissues containing early changes of chronic pancreatitis agrees with the recent findings of Klöppel and Mallet.

We have not yet defined the characteristics of those patients who showed abnormal pancreatic expression of MHC antigens, nor have we yet ascertained whether expression is restricted to a specific subpopulation of patients. There was no relation between abnormal MHC expression and either age or the currently accepted causes of chronic pancreatitis. Our findings do not support a clear differentiation between ethanolic and non-ethanolic pancreatitis. In fact, the classification of chronic pancreatitis based on the causes and morphological patterns may have to be reviewed in the light of the present and further immunological studies. Our findings suggest that a cell mediated immune reaction represents one stage in a 'multistep' pathogenetic process and could act as a common pathway in different types of chronic pancreatitis. In some cases of chronic pancreatitis, different causes might induce abnormal MHC expression and allow an autoimmune reaction to develop. This process seems to begin in the ductal epithelium and the interlobular and intralobular ductules. Alternatively, MHC expression is secondary to lymphocyte infiltration and activity. While α and β interferons are produced by fibroblasts, the more potent γ interferon is a T-cell derived lymphokine. All three classes of interferons can enhance both class I and II MHC expression, and in vitro studies have shown interferon induced class I expression in class I negative cell lines. What is less certain, however, is the ability of interferon to induce de novo synthesis of MHC gene products that are not spontaneously expressed in vivo. Whether aberrant MHC determination or expression represents a primary epithelial or lymphocyte dysfunction, its presence implicates a cell mediated cytotoxic reaction in the pathogenesis of chronic pancreatitis.

This study has identified the expression of cellular components that may contribute to an autoimmune mechanism in the pathogenesis of chronic pancreatitis. It is not suggested that autoimmunity is either the primary or even the main cause in this disease. Elucidation of the mechanisms by which primary injury is sustained by pancreatic exocrine epithelial cells will require a separate study specifically designed to answer that question. 'Aberrant' expression of MHC determinants by epithelial cells requires cautious interpretation, because it might just arise in the vicinity of an inflammatory focus. Similarly, spontaneous ('normal') expression of HLA-DR antigens does not point to a predisposition to autoimmune disease. Nevertheless, we have not seen calcific obstruction of pancreatic ducts early in the disease process. Expression of HLA class I and II determinants by exocrine epithelial cells occurred in a regional, sublobular manner and its distribution was incompatible with a purely obstructive cause. Furthermore, such expression was found of the epithelial cells in morphologically normal lobules without either a significant inflammatory component or evidence of epithelial destruction. Additional studies are now required to elucidate the nature and timing of this cell mediated immune reaction. If subsequently proved, it holds exciting treatment prospects for changing the clinical management of this difficult disease.
Expression of major histocompatibility antigens in chronic pancreatitis

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