Autoantigens in primary biliary cirrhosis

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Abstract
The autoimmune liver disease primary biliary cirrhosis (PBC) is characterised by serum autoantibodies directed at mitochondrial and nuclear antigens (seen in most patients and a subset of patients, respectively). The antimitochondrial antibodies (AMA) characteristic of PBC are directed at members of the 2-oxoacid dehydrogenase components of multi-enzyme complexes; in particular, the E2 and E3 binding protein (E3BP) components of the pyruvate dehydrogenase complex (PDC). The presence of autoantibodies reactive with PDC-E2 and/or E3BP is strongly predictive of the presence of PBC. Therefore, the detection of these antibodies plays a very important role in the diagnosis of PBC. Originally demonstrated using immunofluorescence approaches, AMA can now be detected by the use of commercially available enzyme linked immunosorbent assays (ELISAs). Although the ELISA based approaches have advantages in terms of laboratory practicality, they are slightly less sensitive for the diagnosis of PBC than immunofluorescence (occasional patients with PBC show reactivity with PDC related antigens not present in the antigen preparations available for use with ELISA). Therefore, immunofluorescence should continue to be available as a complementary diagnostic test for use in occasional patients. In a subset of patients with PBC, autoantibodies are directed at increasingly well characterised nuclear antigens. Antinuclear antibody (ANA) positive patients are typically AMA negative. There are no significant differences in disease phenotype between AMA positive and AMA negative groups. At present, the clinical detection of ANA is mostly by Hep2 immunofluorescence, although ELISA kits for individual nuclear antigens are increasingly becoming available.

Keywords: liver cirrhosis; biliary; autoimmunity; antibody

Primary biliary cirrhosis (PBC) is a chronic cholestatic liver disease characterised by inflammatory damage to the biliary epithelial lining the small intrahepatic bile ducts. Recent epidemiological studies have suggested that PBC, which typically, although not exclusively, affects middle aged women, is a much more common clinical problem than had previously been realised. The most recent estimates suggest that it affects up to one in 700 women over the age of 40. It is increasingly evident that PBC is an important cause of morbidity and mortality in the affected population. Over 80% of patients are symptomatic (the most pervasive symptoms being fatigue and pruritus), whereas progression of the disease to cirrhosis, with associated complications including variceal haemorrhage, contributes to the greatly increased standardised mortality ratio seen in patients with PBC.4 6

Recent years have seen important advances in the treatment of PBC. For patients with end stage disease, orthotopic liver transplantation is highly effective, with patients reporting good quality of life postoperatively.7 6 Reversal of the histological features of PBC (or at least slowing down of the progression of disease), thereby reducing the symptomatic burden, and the risk to life and requirement for transplantation that are associated with advancing disease, remains the goal of medical treatment. The hydrophilic bile acid ursodeoxycholic acid (UDCA), given orally, has been shown to be effective in slowing disease progression and has entered widespread clinical use.9–12 It is likely, however, that UDCA is most effective when given early in the disease course. The importance of early treatment emphasises the necessity of early diagnosis. Fortunately, the autoimmune nature of PBC, and the specificity of its characteristic autoantibodies, are now well established. What is perhaps less well appreciated, however, is the sheer strength of the serological associations in PBC, and the resulting power of autoantibody screening in isolation as a diagnostic test for this disease. The purpose of this leader is to review the current data regarding the autoimmune response and its diagnostic detection in PBC, with emphasis on clinical applications. I will discuss each of the major PBC autoantigen families in turn outlining the nature of the antigen, the nature of the immune response mounted to it, and the clinical applications and relevance of detecting this response.

The M2 mitochondrial autoantigens

NATURE OF THE ANTIGEN(S)

It has been appreciated since the 1960s that most patients with PBC have high titres of antibodies in their serum reactive with non-organ, non-species specific antigens present within mitochondria.13–14 The detection of these antimitochondrial antibodies (AMA) using immunofluorescence remains one of the most important diagnostic tests for PBC. Subsequent experimental approaches further localised the mitochondrial antigens to the inner mitochondrial membrane (and the M2 mitochondrial subfraction)15 before, in the late 1980s, Gershwin and colleagues, using patient serum to screen an expression library, identified a cDNA clone encoding the dominant

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mitochondrial antigen. This antigen was subsequently identified as the dihydrolipoamide acyltransferase (enzyme 2 (E2)) component of the pyruvate dehydrogenase complex (PDC). Subsequently, it has been shown by enzyme linked immunosorbent assay (ELISA) and immunoblotting that antibodies reactive with PDC-E2 are present in the serum of 95% of patients with PBC, confirming that it is to this antigen that the dominant autoantibody response in PBC is mounted.

PDC is a member of the 2-oxoacid dehydrogenase family of multienzyme complexes, the other members of which are 2-oxoglutarate dehydrogenase complex (OGDC) and branch chain 2-oxoacid dehydrogenase complex (BCOADC). PDC, OGDC, and BCOADC all have fundamental roles in cellular metabolism. These complexes are highly conserved throughout evolution, and are present in all organisms from rickettsia upwards. All three complexes share a common structure, with multiple repeats of E1 (2-oxoacid dehydrogenase) and E3 (dihydrolipoamide dehydrogenase) subunits built around an E2 (and in the case of PDC only, E2 and E3 binding protein (E3BP)) core. All the component subunits are nuclear encoded, and undergo post-translational transport and active uptake into the mitochondria, where complex assembly occurs. Some component modification occurs within the mitochondria, most notably the addition of a lipoyl acid group to the E2 (and where present E3BP) components. Lipoic acid plays a crucial role in the enzymatic function of each complex.

The E2 chains of PDC, OGDC, and BCOADC share a highly segmented structure. They comprise, from the N-terminus, one to three lipoyl domains, a peripheral subunit binding domain, and a large core forming acyltransferase catalytic domain, all linked together by long (25–30 residues) segments of flexible polypeptide chain rich in alanine and proline.

The three dimensional structures of PDC-E2 of both prokaryotic and eukaryotic origin have been determined by means of nuclear resonance spectroscopy. A common structure consisting of two four stranded β-sheets forming a flattened barrel, with the lipoylation site (a lysine residue physically exposed at the tip of a tight turn in one of the β-sheets), and the N-terminal and C-terminal residues close together at the other end of the molecule, is seen. Little is known, as yet, regarding the three dimensional structures of the other PDC component subunits.

THE AUTOIMMUNE RESPONSE IN PBC

Although, as will be discussed below, immunofluorescence remains a valuable clinical tool, further characterisation of the antigen specificity of the AMA response in PBC was not possible using this technique. This characterisation was performed using immunoblotting and ELISA to detect responses to purified antigens. Different groups have adopted individual approaches to the purification of antigen for use in these experiments. These divide broadly into recombinant and biochemical purification (from PDC rich tissue) based approaches. Each approach has advantages and disadvantages, and it must be appreciated that some of the apparent differences in responses described in studies from different groups are merely a result of the different approaches taken to antigen preparation.

Immunoblotting of serum from patients with PBC against biochemically purified, tissue derived PDC showed that 95% of patients who are AMA positive by immunofluorescence have antibodies reactive with PDC-E2. Immunoblotting and ELISA of serum from patients with PBC against PDC of human origin has confirmed that these are true antibodies showing reactivity with self antigens. Antibodies reactive with E3BP (formerly known as protein X) are present in all patients showing reactivity with PDC-E2. The absence of sequence data related to PDC-E3BP precluding synthesis of recombinant antigen, and the failure of biochemical approaches to separate PDC-E3BP from PDC-E2, have, until recently, prevented the isolation of PDC-E3BP and the characterisation of specific autoreactive responses to it. Thus, almost all work on the dominant autoreactive responses in PBC has been directed at PDC-E2.

The recent cloning and expression of human E3BP has allowed this imbalance to be redressed. The anti-PDC-E2 totally removes antibodies reactive with E3BP but not vice versa. The degree of crossreactivity at the B cell level between PDC-E2 and E3BP has meant that serological approaches cannot tell us to which of these antigens the initial breakdown of tolerance occurs (the wider distribution of B cell epitopes within the PDC-E2 than E3BP is suggestive but not more) that tolerance breakdown is initially to an epitope in PDC-E2 that is cross-reactive with E3BP, with subsequent epitope spreading to non-crossreactive epitopes. A recent study, demonstrating a complete absence of T cells in the peripheral repertoires of patients with PBC who are reactive with recombinant human E3BP (in contrast to the responses seen to recombinant PDC-E2 in > 90% of the same patients) suggests that the breakdown of tolerance is principally, if not entirely, to PDC-E2.

Antibodies reactive with the E1α and E1β subunits (which are not crossreactive with PDC-E2 or PDC-E3BP) are seen at lower frequency (40% and 10% of patients with PBC, respectively). Antibodies are seen to the E2 components of OGDC and BCOADC in approximately 90% and 50% of patients with PBC, respectively. Antibodies are not seen to the E3 subunit common to PDC, OGDC, and BCOADC, and to the E1 subunits of OGDC and BCOADC. OGDC and BCOADC do not have E3BP.

Attempts to map the B cell epitopes within PDC-E2 using different experimental approaches (trypsinised whole PDC-E2, truncated recombinant polypeptide, and sequence...
specific peptides) have generated largely consistent results. The dominant B cell epitope within PDC-E2 is conformational in nature and spans the inner (of two) lipoic acid binding domains.\textsuperscript{38}–\textsuperscript{40} Significantly lower (100 times) titre responses are seen to the outer lipoil domain. Responses have not been reported to the other domains of PDC-E2. The lipoic acid attached to the inner lipoil domain of PDC-E2 itself appears to constitute part of the epitope, the AMA response to lipoylated recombinant PDC-E2 being of significantly higher titre and affinity than that seen to unlipoylated antigen.\textsuperscript{41} The role played by lipoic acid in the B cell autoepitope of PDC-E2 has been confirmed in studies of the antigen specificity of human monoclonal antibodies secreted by hybridomas derived from peripheral blood B cells from patients with PBC.\textsuperscript{42} Two (of five) hydridomas were found to be specific for lipoylated recombinant human PDC-E2 inner lipoil domain, with no binding being seen to unlipoylated antigen. The other three monoclonal antibodies showed greatly reduced binding to unlipoylated antigen. These findings suggest that the lipoic acid residue forms a part, of variable importance, of the dominant B cell epitope within PDC-E2. In a proportion of B cells, the lipoic acid constitutes an important part of the epitope that its absence abrogates antigen–antibody binding.\textsuperscript{43}

Less is known about epitope localisation within PDC-E3BP. Although a pronounced response is seen to the lipoic acid binding domain, which shows considerable homology with the inner lipoil domain of PDC-E2,\textsuperscript{44} some patients also report to the non-homologous "catalytic" domain. The dominant epitopes within OGDC-E2 and BCOADC-E2 appear, as is the case with PDC-E2, to be conformational in nature and to be contained within the lipoic acid binding domain.\textsuperscript{45} The lipoic acid group appears to constitute part of the dominant B cell epitope within PDC-E3BP,\textsuperscript{46} but not those of OGDC-E2 and BCOADC-E2.\textsuperscript{47–49}

Important questions that remain unanswered include the mechanism by which immune self tolerance to ubiquitous and highly conserved antigens such as PDC, OGDC, and BCOADC breaks down, and the reasons why tolerance breakdown to these antigens results in a disease with such a restricted tissue distribution (limited to the biliary, salivary, and lacrimal epithelium). Another important issue requiring clarification is the precise role played by different effector mechanisms in the pathogenesis of target cell damage in PBC and in particular the question of the direct role, if any, played by anti-PDC antibodies.\textsuperscript{50} The observation that placental transfer of IgG anti-PDC can occur, and is associated (in two cases at least) with liver damage (albeit not typical PBC) in the baby raises the possibility that anti-PDC antibodies play a direct role in disease pathogenesis.\textsuperscript{51}

One observation of potential relevance for our understanding of the tissue tropism of PBC is that the subcellular distribution pattern of PDC is abnormal in PBC, with aberrant cell surface expression of the complex (which is normally exclusively mitochondrial in distribution) being seen in the affected biliary and salivary epithelia.\textsuperscript{52–54} The mechanism giving rise to this aberrant cell surface expression of PDC is unclear, but the end result could be that the cells are vulnerable to antibody dependent pathways of cell cytolysis. An alternative potential mechanism whereby anti-PDC antibodies could contribute directly to target cell damage is suggested by the recent observations that IgG antibodies can penetrate living cells and interfere with cellular function.\textsuperscript{55} As yet no data are available regarding the penetration of IgG anti-PDC into epithelial cells and the resulting effects, if any, on cell metabolism function and integrity.

Although data on IgG anti-PDC penetration into epithelial cells in PBC are lacking there is strong, if indirect, evidence to suggest that dimeric IgA anti-PDC can enter and be exported across epithelial cells as part of the normal physiological process of IgA transcytosis.\textsuperscript{56–58} Moreover, in vitro systems at least, IgA anti-PDC taken up into cultured epithelial cells has been shown to colocalise with PDC-E2 within the cell.\textsuperscript{59} In viral infection model systems, it has been shown previously that IgA specific for viral antigens can, while undergoing transcytosis, bind to viral antigens within infected epithelial cells and export them from the cell.\textsuperscript{60} It has been proposed, therefore, that IgA anti-PDC could, while undergoing transcytosis, bind to nascent PDC component polypeptides and export them from the cell, thereby leading to chronic PDC depletion and metabolic damage to the cell.\textsuperscript{61} This model would, if correct, help to explain both the aberrant expression of PDC on the epithelial cell surface (it is taken there by transcytosing IgA anti-PDC) and the tissue tropism of the disease (it is restricted to those tissues that export IgA by the process of transcytosis). Although this model is intriguing there remains, at present, little hard evidence to support it.

An alternative possibility is that AMA, in fact, play little or no direct role in target cell damage in PBC and are merely an “immunological overspill” phenomenon resulting from the generation of a (increasingly well characterised) T cell response that is itself responsible for the damage.\textsuperscript{62–64} If autoreactive T cell responses do indeed play a dominant role in the pathogenesis of PBC it is perhaps surprising that the major histocompatibility complex (MHC) associations seen in the disease are relatively weak.\textsuperscript{65} One potential explanation for this finding comes, however, from the observation that the dominant T cell autoepitope in PBC contains anchor residues allowing it, theoretically at least, to bind to multiple human MHC (HLA) alleles.\textsuperscript{66–68} Data from a recently described mouse model of PBC (experimental autoimmune cholangitis (EAC)), in which sensitisation of SJL/J mice with PDC induces immunological and histological features characteristic of PBC, appear to confirm the view that T cell mediated responses are responsible for target cell damage. The generation of autoreactive anti-PDC antibodies is a rapid
and non-specific feature of the sensitisation of mice with PDC, which is non-predictive of the development of bile duct lesions. In contrast, the generation of an autoreactive anti-PDC T cell response correlates completely with the presence and kinetics of histological damage.58 67–69

CLINICAL RELEVANCE
In contrast to their uncertain pathophysiological relevance, the importance of anti-M2 AMA as diagnostic markers for PBC cannot be overstated. Anti-M2 AMA are among the most sensitive and specific of all serological disease markers. More than 95% of patients with liver biopsy and serum liver biochemical (a typically cholestatic pattern) features of PBC will be AMA positive by immunoblotting, and will have serum anti-PDC antibodies detected by immunoblotting. An ELISA developed by our group (using biochemically purified PDC-E2/E3BP as antigen) has been shown to have a sensitivity of 93% and a specificity of 96% for the diagnosis of PBC.70 The combination of AMA and cholestatic liver function tests (LFTs) is therefore highly suggestive of the diagnosis if PBC. Liver biopsy is still recommended, however, to confirm the diagnosis and, in particular, to allow histological staging. The strength of the association between AMA in PBC is such, however, that the presence of the autoantibody in isolation is strongly suggestive of the presence of histological features of PBC in the liver. In 1986, we reported the outcome of investigations performed on a series of patients who were AMA positive, but who had no symptoms suggestive of PBC, and who had normal LFTs. Twenty four of these 29 patients were found to have liver histological features diagnostic of, or compatible with, PBC.71 In 1996, we reported on the outcome of this cohort of patients after follow up of a median of 17.8 years.72 Of the original 29 patients, 22 (76%) had developed typical symptoms of PBC and 24 (83%) had developed abnormal LFTs with a typical cholestatic picture. Interestingly, although five patients had died, none had died of liver related causes, and of the 10 who had undergone a further liver biopsy for clinical reasons, none had developed cirrhosis (stage IV PBC). Therefore, the conclusions from these two studies are that the presence of AMA is strongly suggestive of the presence of histological features of PBC in the liver, even in the absence of symptoms or abnormal LFTs, and that, over time, such patients will go on to develop “classic” clinical PBC. However, patients presenting in this manner appear to have a slowly progressive form of the disease. Our current practice is to monitor annually for LFTs all patients who are AMA positive and have normal LFTs. If, during follow up, patients develop cholestatic LFTs, we would then proceed to liver biopsy to confirm and stage the disease before starting UDCA treatment. We believe that aggressive monitoring of patients with such early disease is warranted because the efficacy of UDCA treatment is highest when started early in the disease process.

It is unclear whether the titre of AMA seen in patients with PBC has any predictive value for disease severity. One early ELISA study suggested that there was a positive association between disease stage and IgG AMA titre.73 This observation appeared to support earlier ones made using indirect immunofluorescence based approaches to IgG AMA detection, which similarly showed increasing AMA titre with advancing disease.73 74 More recently, an association between serum IgA AMA titre and liver inflammatory activity in both the untreated and UDCA treated state has been reported.75 A further study has, however, failed to show any significant association between AMA titre detected by either immunofluorescence or ELISA and disease severity as assessed by the Mayo score (a well validated disease prognosis model).76 My feeling is that any correlation between AMA titre and disease activity/severity/progression is limited and of no practical relevance in the clinical assessment of patients.

One practical issue in the serological detection of AMA now faced by diagnostic laboratories and clinicians is the choice of assay. Indirect immunofluorescence remains the “gold standard” for the serodiagnosis of PBC (having close to 100% specificity77) but is labour intensive and observer dependent. Newer approaches such as ELISA and assays based on the ability of AMA to inhibit the enzymatic function of PDC, a phenomenon that can be relatively easily adapted for use in a colorimetric assay,78,80 are less labour intensive. The question is, however, are they as clinically useful? As described above, an ELISA using biochemically purified PDC-E2/E3BP has been shown to be 93% sensitive and 96% specific for the diagnosis of PBC. The use of this PDC-E2/E3BP preparation has the advantage that it contains both the major antigenic proteins and retains physiological conformation—important when the dominant epitopes are conformational in nature. The disadvantage of this ELISA is that it requires tissue derived antigen and is thus not practical for widespread laboratory use. Therefore, the use of ELISAs based on recombinant antigens is theoretically attractive. It appears, however, that although the specificity of ELISAs using recombinant antigen can be very impressive, sensitivities can be lower than those reported for immunofluorescence or ELISA using whole antigen.78 81 The reasons for this lower sensitivity are twofold. First, there is the issue of the conformational nature of the epitopes, reduced or lost in recombinant molecules. Second, there is the problem of the (small) subgroup of patients with clear cut PBC who have AMA that react exclusively with non-PDC-E2 mitochondrial antigens (PDC-E1, PDC-E3BP, OGDC-E2, and BCOADC-E2). Although “designer” recombinant molecules have been synthesised containing OGDC-E2 and BCOADC-E2, in addition to PDC-E2, these cannot be used for patients responsive to PDC-E1 only (up to 2% of the total PDC population). In the future, PDC function inhibition based assays might be useful to deal with
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200 kDa.87 88 This protein has been identified as a protein with a molecular mass of about 200 kDa.

Most of the autoantibodies from patients with PBC that label the nuclear envelope recognise the lamin B receptor.96 Autoantibodies from four of these rare patients have been shown to recognise a conformational epitope contained within amino acids 1–60 of the nucleoplasmic N-terminal domain.99 So far, autoantibodies against the lamin B receptor also appear to be PBC specific and are more often present in the sera of patients who are AMA negative.

M-ANA

PBC sera giving a multiple nuclear dot immunofluorescence pattern have been shown to bind to proteins of 95 kDa100 and 78–92 and 96–100 kDa101 in immunoblots. Two nuclear proteins have now been shown to react with

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these MND-ANA. The first to be cloned was Sp100. This cDNA encodes a protein of unknown function with sequence similarities to several transcriptional transactivating proteins, including human immunodeficiency virus 1 (HIV-1) nef proteins. Large scale screening of anti-Sp100 autoantibodies in sera from patients with various diagnoses has been performed by ELISA using recombinant proteins: anti-Sp100 antibodies were found in 27% (50 of 184) of patients with PBC, but in none of the patients with other liver diseases. Anti-Sp100 antibodies are occasionally found, however, in rheumatic disorders. 

Epitope mapping experiments have shown that one domain of Sp100 (containing the sequence showing similarity with the HIV nef proteins) is recognised by all anti-Sp100 positive sera. Two major immunodominant epitopes have now been identified at amino acids 296–311 and 332–351. A second protein that colocalises to dot like nuclear domains, and which is aberrantly expressed in promyelocytic leukaemia cells (PML), has also been shown to react with MND-ANA in PBC sera. PML, as is the case with Sp100, shows sequence similarity with transcription factors, suggesting a possible role in transcription regulation. Interferons have been shown to increase the concentrations of both Sp100 and PML mRNA and protein. Autoantibodies against PML are as highly prevalent and specific for patients with PBC as those against Sp100. Again, a single, and as yet unconfirmed, study has suggested that patients with PBC who are Sp100 and PML positive might have a worse prognosis than antibody negative patients (in this case, antibody positive patients were shown to be more likely to progress from early (stage I/II) to late disease (stage III/IV) over a follow up period of two years than anti-Sp100/anti-PML negative patients.

The role played by all the PBC specific ANA in target cell damage is, as is the case with AMA, unclear. Anti-Sp100 and anti-gp20, as well as classic AMA persist post-liver transplantation, even in the absence of disease recurrence.

Non-M2 mitochondrial antigens

It has been postulated by the Tubingen group that there are other mitochondrial antigens in addition to the M2 family that are important in PBC. The importance of the responses to these antigens and, indeed, the identities of the antigens themselves are controversial. Berg and colleagues initially described three antigens (M4, M8, and M9) associated with the outer mitochondrial membrane to which antibody responses are seen in PBC. Retrospective studies suggested that patients who were anti-M2 and anti-M9 positive at diagnosis had a relatively benign subsequent disease course, whereas patients who were anti-M4 and anti-M8 positive at presentation had a worse outcome. A prospective study carried out by the same group has seemingly confirmed this finding. Although a serological marker predictive, at presentation, of the future disease course would be of the utmost value in the clinical management of patients, doubts remain about the importance of the data relating to anti-M4, anti-M8, and anti-M9. The greatest problem with regard to these studies is that they have not, as yet, been independently replicated. This is a result, in part, of the complex nature of the experimental approaches needed to detect these antibodies. Antibodies to M4, M8, and M9 are not detectable by standard immunofluorescence. Instead, detection requires the use of complement fixation or ELISA in the case of M4 and M8, and ELISA or immunoblotting in the case of M9. In the absence of positive identification of the M4, M8, and M9 antigens (early reports identifying M9 as glycogen phosphorylase and M4 as sulphite oxidase have not been confirmed), fresh mitochondrial membrane preparations containing the relevant antigen fraction have to be used in the above assays. This drastically limits the extent to which these assays can be standardised between laboratories. The technical problems associated with the detection of anti-M4, anti-M8 and anti-M9 antibodies, together with the uncertainties regarding the biochemical identity of the antigens, have led to the suggestion that the M4/M8/M9 nomenclature should be set aside pending clearer information on the character of these antigens and their disease associations. The potential clinical value of serological marker(s) predictive of prognosis, which would allow better targeting of treatments to high risk patients, is such that this area should be regarded as a priority for future research.

Other antigens

In addition to the well described responses to mitochondrial and nuclear autoantigens other autoantibody responses have been reported in PBC. One autoreactive response of potential clinical relevance is that directed against platelet specific antigens. Indeed, the inverse correlation between the titre of antiplatelet antibodies and platelet counts suggests that autoreactive antplatelet responses contribute (together with non-autoimmune processes such as splenic pooling in patients with portal hypertension) to the thrombocytopenia seen in PBC. It has been demonstrated recently that the antigens to which autoreactive antplatelet responses are directed in PBC are the platelet glycoprotein complexes Ib–IIa and Ib–IX. Testing for antplatelet antibodies remains a research tool at present, but the possibility of an autoimmune process should always be born in mind in patients with PBC suffering from thrombocytopenia.
Conclusions

PBC is one of the autoimmune diseases with the best characterised serology. The recogni-
tion of the prevalence of autoantibody responses and subsequent identification of the
important autoantigens has greatly contributed to our understanding of this disease. We are,
however, still learning about the disease. Perhaps the most important new development
in our understanding of the immunology of PBC, as it relates to clinical practice, is our
appreciation of the sheer scale of the association between the presence of the histological
features of PBC and serum AMA and PBC specific ANA, and the concomitant value of these
serological markers for clinical diagnosis.

It is probably true to say that what in many ways represented the advent of our under-
standing of PBC, the observation that autoantibodies are present in the serum of patients
with PBC remains, several decades later, the single most important phenomenon in the
clinical management of the disease.

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