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Mechanisms of Environmental-Induced Autoimmunity

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Abstract

Although numerous environmental exposures have been suggested as triggers for pre-clinical autoimmunity, only a few have been confidently linked to autoimmune diseases. For disease associated exposures, the lung is a common site where chronic exposure results in cellular toxicity, tissue damage, inflammation, and fibrosis. These features are exacerbated by exposures to particulate material, which hampers clearance and degradation thus facilitating persistent inflammation. Coincident with exposure and resulting pathological processes is the post-translational modification of self-antigens, which, in concert with the formation of tertiary lymphoid structures containing abundant B cells, is thought to promote the generation of autoantibodies that in some instances demonstrate major histocompatibility complex restriction. Under appropriate gene-environment interactions these responses can have diagnostic specificity. Greater insight into the molecular and cellular requirements governing this process, especially those that distinguish pre-clinical autoimmunity from clinical autoimmune disease, may facilitate determination of the significance of environmental exposures in human autoimmune disease.

Keywords

autoimmunity; innate; adaptive; xenobiotic; inflammation; animal model

INTRODUCTION

The molecular and cellular events that contribute to autoimmune diseases are complex. Nevertheless, many general principles of disease pathogenesis have been elucidated (1), however, the initiating steps that drive auto-reactivity remain for the most part poorly understood. It is for this reason that autoimmune diseases are often described as idiopathic, or of apparent spontaneous origin of unknown cause or mechanism. However, a number of environmental exposures linked to either overt autoimmune disease (2), or to subclinical autoimmunity that manifests solely as cellular infiltrates or autoantibodies (3; 4), have been identified. The role of these environmental exposures as initiating events in autoimmunity has been confirmed in certain induced experimental animal models (5). Accordingly, unlike the idiopathic autoimmunity in autoimmune-prone mice (6; 7), an essential feature of induced models is that the initiating factor is known making it possible to study the temporal and mechanistic course of events that lead to disease.

Among the possible initiating events, epidemiological studies have identified numerous infectious and non-infectious exposures associated with autoimmune diseases (8–11). These can be classified into three broad classes; biological, chemical and physical (8). Biologic agents include infection, dietary constituents, and therapeutics derived from biological components. Chemical agents comprise non-biological exposures that result from occupational, environmental, therapeutic, and lifestyle activities. Physical agents encompass radiation including sunlight, electric and magnetic fields. Autoimmunity and autoimmune diseases associated with infection (12; 13) or with therapeutics, either chemical or biological (14; 15), have been discussed elsewhere and will not be covered here. This review focuses on xenobiotics (chemicals not naturally produced by or present within an organism) that are linked to human autoimmune diseases and their corresponding experimental models used to study mechanisms of disease induction and development. Additional emphasis will focus on xenobiotics that can cause subclinical inflammation and autoimmunity but progress to clinical disease only in susceptible genetic backgrounds. This second group may offer clues to idiopathic autoimmune disease triggers because of evidence that autoimmune diseases are preceded by preclinical phases characterized by inflammatory mediators occurring prior to, or concurrent with, autoantibodies (16–19). Moreover, profiles of inflammatory mediators and autoantibodies can help distinguish between healthy individuals, autoantibody positive healthy individuals, and those with autoimmune disease (17; 18). Thus, a detailed dissection of the innate and adaptive immune sequelae that arise following autoimmunity-promoting xenobiotic exposures will likely contribute significantly to our understanding of the etiopathogenesis of human autoimmunity and autoimmune diseases.

XENOBIOTIC EXPOSURES ASSOCIATED WITH HUMAN AUTOIMMUNITY AND AUTOIMMUNE DISEASES

Although numerous xenobiotics have been suggested as triggers of autoimmunity and autoimmune diseases, definitive association has proven difficult to establish (8) often because of insufficient evidence of exposure to demonstrate causality. Consequently, only a small number of xenobiotics have been strongly implicated in autoimmune diseases. This is exemplified in the findings of a National Institute of Environmental Health Sciences (NIEHS) Expert Panel convened to determine environmental influences on the development of autoimmune diseases (8). Using defined guidelines, associations between individual exposures and autoimmune diseases were classified as “confident” or “likely” based on published evidence. Also identified were a number of exposures where the evidence for causation of autoimmune disease was considered “insufficient” but where exposure leads to features of autoimmunity. Since that review, little has changed to alter the exposure associations. Table 1 lists examples of “confident”, “likely”, or “insufficient” associations that will be discussed.

Silica

Silica, as SiO₂, is most commonly found in nature as quartz. It becomes an occupational hazard when inhaled in a particulate crystalline form in workplaces involved in dusty trades such as mining, tunneling, and sand blasting (20–22). Pulmonary deposition of silica particles (<10 μm in diameter) leads to inflammation and fibrotic nodule formation culminating in the respiratory disease silicosis (22). The link between occupational exposure to silica and autoimmune disease, specifically rheumatoid arthritis (RA), was made by Caplan and colleagues (23; 24). Following those initial observations, numerous reports linked silica dust exposure to systemic lupus erythematosus (SLE), RA, systemic sclerosis (SSc), and anti-neutrophil cytoplasmic antibody (ANCA)-related vasculitis (8; 25). Evidence of a breakdown in immune tolerance, including autoantibodies, can occur with silica-exposure in the absence of silicosis (26), and exposure intensity rather than cumulative dose may be important in development of autoimmune disease (25). Recently, autoimmune diseases associated with silicosis have been reported following exposure to dust from the fabrication of artificial stone (>90% silica) (20; 27; 28). Such exposure is associated with acute and accelerated silicosis (20) and incidence of a heterogeneous pattern of autoimmune features and diseases (27; 28). This is similar to autoimmune disease in miners following exposure to very high amounts of silica-containing dust (25; 29; 30). Gene-environment interactions have not been examined for silica-induced autoimmunity, but susceptibility to silicosis is associated with polymorphisms in family with sequence similarity 13 member A (FAM13A) (31) and tumour necrosis factor alpha (TNF-α) (32).

Smoking

Smoking is a well-known risk factor for RA (8; 33), particularly for seropositive RA, which includes either rheumatoid factor (RF) or anti-citrullinated protein antibody (ACPA). Both smoking status and intensity increases risk (19). Another well-known risk factor for RA are certain human leukocyte antigen (HLA) class II alleles which share a conserved amino acid

sequence in their antigen binding site called the HLA shared epitope (SE) (34). Meta-analysis has identified significant association between the SE, smoking exposure, and ACPA, suggesting a gene-environment interaction between smoking and HLA linked autoantibody responses in RA (35). Weaker and less consistent associations have been documented between smoking and SLE, multiple sclerosis (MS), and Crohn's disease (8). Single studies in SLE have also reported gene-environment interactions between smoking and interleukin (IL)-33 (36), integrin subunit alpha M (*ITGAM* or CD11b) (37), and N-acetyltransferase 2 (*NAT2*) (38).

Solvents

Several independent meta-analyses have supported an increased risk of SSc with solvent exposure (39; 40). A large number of different solvents have been implicated (39; 41; 42) along with a higher prevalence of disease in males compared to unexposed cohorts (43; 44). Disease severity was also greater in those with exposure (44). Exposure is primarily occupational (39) involving inhalation or dermal adsorption, with greatest risk from aromatic solvents, trichloroethylene (TCE), halogenated solvents, and ketones (40). A less established association has been described for solvent exposure and MS (8; 45) and the gene environment interaction of solvent exposure, smoking and MS-linked HLA genes (46) was found to dramatically increase the risk of MS (47).

Mercury

The association between mercury exposure, autoimmunity and autoimmune diseases has been recently reviewed (3; 4). Mercury exists in elemental, organic and inorganic forms and exposure can occur via occupation, dietary contamination, therapeutic or cosmetic agents, and fossil fuel emissions (3). Although current evidence for mercury induction of autoimmune disease is insufficient (4; 8), mercury exposure is linked to markers of inflammation and autoimmunity (3; 4; 48). Studies of artisanal gold miners and fish consumption in South America identified a significant association between anti-nuclear autoantibodies (ANA) and mercury levels in blood or hair (3; 4), as well as increased levels of inflammatory cytokines with mercury exposure (49; 50). Several other cohorts, including the National Health and Nutrition Examination Survey (NHANES), and the Long Island Study of Seafood Consumption, have produced conflicting data on the relationship between autoantibodies, cytokines and mercury levels (3; 51), but this may reflect the low level of exposure in those groups. Several reports have identified nephrotic syndrome as an outcome of mercury exposure most often from skin whitening/lightening cream use leading to membranous glomerulonephritis and sometimes autoantibodies (3).

Asbestos

Asbestos refers to a group of naturally occurring silicate minerals broadly classified into the serpentine (chrysotile) and amphibole (crocidolite, amosite, tremolite, anthophyllite, actinolite) groups (52). Amphibole fibers are needle-like and brittle, while serpentine fibers are curved and softer. Exposure may occur via different routes, but inhalation is the most common and can lead to asbestosis as well as malignancy (53). Amphibole asbestos exposure appears to be more closely associated with autoimmunity than the serpentine form (54). Recent analysis of the association between asbestos exposure and autoimmunity (8; 55;

56) concluded that while epidemiological evidence for a causal role is insufficient, it does support a higher than expected risk of systemic autoimmune disease. Additionally, the presentation of rheumatological symptoms can be atypical, making diagnosis of specific autoimmune diseases difficult (57). A caveat is that, particularly with mining, asbestos exposure may also include concomitant exposure to crystalline silica (8). Autoantibodies, especially ANA, have been reported in cohorts exposed to asbestiform amphibole-contaminated vermiculite (Libby, Montana) (58; 59), mined crocidolite (blue asbestos) (Wittenoom, Western Australia), and fluoro-edenite in quarried rock (Biancavilla, Sicily) (60). Studies of the Libby community also found an increased risk of developing connective tissue diseases (61) which is supported by a recent study showing elevated mortality due to systemic autoimmune diseases such as RA, SSc, and SLE (62).

MECHANISMS LEADING TO XENOBIOTIC INDUCED AUTOIMMUNITY AND AUTOIMMUNE DISEASES

While the ability of certain environmental agents to induce autoimmunity is well established (63–66), to what extent idiopathic and induced disease are mediated by common mechanisms is unclear (67; 68). For example, the same agent can induce different autoimmune disorders (i.e., silica-associated RA, SSc and SLE) (8), while multiple agents can produce a similar clinical picture (i.e., different medications leading to similar lupus-like syndromes) (63). Moreover, it is unclear if a xenobiotic-induced response is an exacerbation of underlying disease, a xenobiotic specific response, or a combination of both. This is reflected in the diagnostic criteria for idiopathic autoimmune diseases which are not designed to delineate similarities and/or differences in biological markers that may be useful as diagnostic criteria for xenobiotic-induced autoimmune disease (69).

Pollard et al (70) have argued that xenobiotic exposure leads to tissue damage and the release of damage associated molecular patterns (DAMPs), including nucleic acids, as well as self- and modified self-antigens. This results in the engagement and activation of Toll-like receptors (TLRs) and other innate sensors, which stimulate the production of inflammatory mediators and the induction of inflammation. This innate response promotes presentation of self- and modified self-antigens to non-tolerant lymphocytes, followed by expansion of autoreactive B and T effector cell populations, and the production of autoantibodies. These events often occur at the site of exposure where they can be linked to the development of ectopic or tertiary lymphoid structures (TLS) and/or hypertrophy of secondary lymphoid organs. The expansion of autoreactive lymphocytes and their subsequent migration to target tissues such as the kidney in SLE, or the joints in RA, results in disease. The occurrence of these events following specific xenobiotic exposures are shown in Figure 1 and Table 2 and discussed below.

Location, location, location.

Although the proposed general mechanism above may not appear dissimilar to that proposed for idiopathic disease (1), significant differences exist in the identity of a triggering exposure and its location. It is immediately noticeable from Table 1 that the lungs are an important site of xenobiotic exposure. Recent reviews have argued that initiation of autoimmunity may

occur at mucosal surfaces such as the lung for RA, as a result of smoking (71; 72), or epithelial surfaces such as the skin in SLE, following ultraviolet (UV) light (72). While it is clear that smoking and UV exposures are not the only risk factors for these diseases, they do highlight the growing appreciation of xenobiotic exposure at these body surfaces as a key step in the initiation of autoimmune reactions.

Tissue damage, inflammatory mediators, and inflammation

Inhalation of xenobiotics results in responses by a number of cell types in the lung that can result in cell death, cell activation, and release of reactive oxygen species, peptides, proteases, chemokines and cytokines (73). Inhalation of particulate matter such as crystalline silica, asbestos fibers, and the particulate phase of smoke results in their ingestion by alveolar macrophages and cell activation. Moreover, attempts to clear these degradation resistant particles can lead to frustrated phagocytosis which results in lysosomal damage and release of proteases and inflammatory mediators (73; 74). The cellular stress elicited by these events together with cell death leads to release of DAMPs including heat shock proteins, nucleic acids, and other cellular constituents recognized by pattern recognition receptors (PRR) such as TLRs (74–77). The subsequent innate inflammatory response leads initially to neutrophil recruitment and influx of blood monocytes and lymphocytes which is reflected in increased cellularity of bronchoalveolar lavage fluid (BALF). The inability to clear and/or degrade particulate material deposited in the lung, particularly in the alveoli, leads to persistent tissue damage, chronic inflammation and fibrosis (78). Notably, significant gene-environment interactions are at play in the inflammatory responses to smoking (75), silica (21), and mercury (3), as the levels of responses vary depending on the mouse strain.

IL-1 is a major contributor to the initial pulmonary inflammation induced by exposure to smoke, silica, and asbestos as blockade of IL-1 or deletion of IL-1 receptor significantly diminishes the inflammatory response as well as the production of other proinflammatory cytokines such as IL-6 and TNF- α (74; 79). The generation of IL-1 is determined, in part, by activation of the NACHT, LRR and PYD domains-containing protein 3 (NALP3) inflammasome leading to cleavage of pro-IL-1 β and also pro-IL-18 which allows release of mature IL-1 β and IL-18 and the amplification of inflammation in the lung (76; 77). In the case of silica-induced lung inflammation, IL-1 α (a DAMP, constitutively present in epithelial and mesenchymal cells) is released from alveolar macrophages following cell damage and functions as an ‘alarmin’ preceding the production of inflammatory mediators, including IL-1 β , and the infiltration of neutrophils (80; 81).

Lysosomal damage, following subcutaneous injection of inorganic mercury (HgCl₂), may be important for early events in mercury-induced inflammation because inhibition of cathepsin B by CA-074 reduces mercury-induced mRNA expression of NLRP3, IL-1 β , IL-6 and TNF- α (82). Although It is not known if pulmonary inflammation due to solvents such as TCE requires NLRP3 and IL-1 β , dichloroacetyl chloride (DCAC), a reactive metabolite of TCE, has been shown to increase expression of NLRP3 and caspase-1 in the liver (83).

With the exception of mercury-exposure (3), there is little information on the role of innate immunity in the development of xenobiotic-induced autoimmunity. However, there is

information about the inflammatory events following smoking (73; 84) and silica exposure (21) that may be relevant. Silica-induced inflammation is dependent on interferon (IFN)- γ , but not IL-12, IL-4, or IL-13. The acute inflammatory response requires IL-17, while chronic inflammation requires type I (IFN and interferon regulatory factor 7 (IRF7)). Silica-induced inflammation and fibrosis can be uncoupled because the innate immune responses of inflammation, neutrophil accumulation, IL-1 β release, and granuloma formation require myeloid differentiation primary response 88 (MyD88), while development of lung fibrosis (collagen deposition) does not. Smoking induced inflammation is also dependent on MyD88 signaling, and increased proinflammatory cytokine production (IL-1, IL-6, IL-8, TNF- α) leading to accumulation of T helper type 1 (Th1) and Th17 CD4+ T cells (75; 84). In contrast, mercury-induced autoimmunity is dependent on endosomal TLR trafficking and signaling, and IFN- γ (3) while type I IFN, IRF7, and NLRP3 are not required.

An additional step which likely plays a role in development of autoimmunity following xenobiotic exposure is the formation of neutrophil extracellular traps (NETs) during NETosis. NETosis is a form of neutrophil cell death that contributes to pathogenesis in autoimmune diseases such as SLE, RA, and ANCA-associated vasculitis (85). Neutrophils undergo NETosis in response to both non-sterile and sterile stimuli (85) including mercury (3), smoking (84), and silica (86). Many of the cellular constituents externalized during NETosis (e.g. DNA, chromatin) are recognized as autoantigens. Additionally, protein arginine deiminase (PAD) 4, which is involved in the release of decondensed chromatin of neutrophils via citrullination of histones (87), contributes to the production of citrullinated autoantigens and the induction of ACPA in RA. NETosis is therefore an important mechanism for self-protein modification, one of several by which xenobiotic exposure can modify self-proteins.

Self-protein modification

Xenobiotics associated with autoimmunity and autoimmune diseases are typically highly reactive and elicit adverse biological responses leading to oxidative stress, cell death, and inflammation. A common consequence of these reactions is post-translational modification of self-proteins via direct interactions or indirect effects mediated via enzymatic, including proteolytic, reactions as a result of an induced response such as oxidative stress. Smoking results in several different forms of post-translational modifications including citrullination, carbonylation, carbamylation (homocitrullination), and lipid peroxidation (19; 88–90). Silica also induces citrullination via a PAD-dependent mechanism (91), and the generation of reactive oxygen species (ROS) (92) leading to lipid peroxidation (93) and carbonylation (94). Oxidative stress following TCE exposure results in malondialdehyde (MDA)- and 4-hydroxynonenal (HNE)-protein adducts, carbonylation and nitration (95). In addition to its high affinity for protein sulfhydryl groups, mercury exposure also leads to ROS production and protein carbonylation (96). Asbestos exposure is known to lead to protein citrullination (97) and carbonylation (98). An important aspect of these post-translational modifications is that they can alter the immunogenicity of self-proteins, and as discussed below, can lead to autoantibody responses in diseases such as RA (99) and SLE (100).

Tertiary (or ectopic) lymphoid structures

Chronic inflammation is a significant component of many diseases including autoimmune diseases (101), and can be associated with development of tertiary lymphoid structures (TLS) (102). TLS arise at sites of non-resolving inflammation (103; 104) and have been found in target organs of autoimmune diseases (105–107) where they are believed to contribute to the persistence of autoimmunity by providing a microenvironment for survival and maturation of autoreactive cells, particularly B cells (103; 108). TLS contain segregated areas of T and B cells, high endothelial venules, and a CD21⁺ follicular dendritic cell (FDC) network important for B cell responses (109). Recent evidence supporting the oral mucosal origins of autoimmune diseases (71; 72) provide a basis for the hypothesis that inhalation of autoimmune disease associated xenobiotics leads to lung damage, inflammation, and subsequent development of TLS, providing a niche for self-reactive T and B cells.

Smoke exposure induces a chronic inflammatory response in the lungs, leading to chronic obstructive pulmonary disease (COPD) (88). Smoking is responsible for over 80% of COPD in Western countries. There is extensive evidence for lung TLS in COPD (88). The progression from smoke exposure to the presence of TLS (88) begins with tissue damage and inflammatory mediator recruitment of neutrophils and macrophages that animal experimentation has shown is essential for the inflammatory process. The subsequent appearance of TLS is characterized by aggregates of mainly IgM⁺IgD⁻ B cells, surrounded by smaller numbers of primarily CD4⁺ T cells. These follicle-like structures also contain CD21⁺CD35⁺ FDCs and nearby CD138⁺ plasma cells. The severity of COPD, particularly emphysema, is positively associated with B cell follicles, and similar TLS are found in mice following chronic exposure to smoking. Several key mediators, including C-X-C motif chemokine ligand 13 (CXCL13), IL-17A, and B-cell activating factor (BAFF), are required for TLS development in cigarette smoke exposed mice (88). Importantly, smoking, lung inflammation, and the development of TLS are argued to play important roles in the preclinical phases of RA, and establish the lung as an important site for the initiation of seropositive RA (19).

Chronic silicosis is characterized by persistent inflammation and fibrosis leading to interstitial lung disease (21; 22; 110). Similar to the early inflammatory response following smoke exposure, silica-induced inflammation is solely dependent on innate immunity as it can occur independently of T, B, natural killer (NK) T, and NK cells (21). Although lymphocytes accumulate in the lung and draining lymph nodes, TLS have not, to our knowledge, been formally described in human silicosis. However, accumulations of lymphocytes in the lung and tracheobronchial lymph node have been described in experimental studies of crystalline silica exposed healthy C57BL/6 and lupus-prone mice (111–113). The typical appearance is of peribronchiolar or perivascular aggregates of B cells (Figure 2) surrounded by (111), or interspersed with (113), T cells that increase in size with exposure duration (111). Presumptive FDCs (CD21/35⁺ cells) can also be found. These lymphoid accumulations stain with anti-IgG suggesting the presence of plasma cells (113), which is supported by immunoglobulins including autoantibodies in the BALF (111; 114). This association is further supported by the observation that docosahexaenoic acid (a ω -3

polyunsaturated fatty acid with anti-inflammatory properties) prevents development of silica-induced lung TLS and autoantibodies in BALF (114).

Although mercury vapor is a major occupational hazard during artisanal small-scale gold mining (48), development of TLS in the lung has not been described in humans. However, dental amalgam-associated oral lichenoid lesions of the oral mucosa were found to show circumscribed lymphoid follicle-like structures containing B cells within a CD3⁺ T cell infiltrate (115). Dental amalgam constituents were suggested to be causally involved, although the mechanism is unknown. Subcutaneous injection of HgCl₂ in mice results in skin induration with marked expansion of the dermis and subcutaneous tissue by neutrophils, macrophages, proliferating fibroblasts and lymphocytes (116) but the presence of TLS has yet to be observed. Transoral mercury exposure leads to multiple accumulations of lymphoid-like cells in the lungs, especially adjacent to bronchioles (70), but whether these are TLS or simple inflammatory lymphoid cell aggregates has not been established.

Exposure to asbestos can lead to chronic inflammation and asbestosis, another form of lung disease characterized by interstitial fibrosis (53). Lung biopsy has shown alveolitis consisting of accumulations of macrophages and lymphocytes (117) but unequivocal evidence for TLS following asbestos exposure has not been described in human or experimental animal studies. This likely reflects the greater focus on outcomes such as fibrosis and malignancy, diagnoses made years after initial exposure. A similar focus on silicotic nodules as a diagnostic marker, years after initial exposure, may also explain the lack of information on TLS in human silicosis. The presence of lung TLS following silica exposure in experimental animals, however, has been documented (111; 114) and are present after shorter exposure times which do not allow extensive development of fibrosis or fibrotic nodules (118). Additionally, the apparent need for lymphocytes in the development of fibrosis (119) is consistent with the formation of TLS early in the disease process. The close association of lung TLS and autoimmunity in other forms of pulmonary fibrosis (120) is further evidence of a linkage between these features.

The above observations clearly support the hypothesis that xenobiotic exposures can lead to the presence of lymphoid accumulations and TLS in the lung, however the role of pulmonary TLS in the development of autoimmunity and autoimmune disease remains unclear. Accordingly, it is undecided if lung TLS are pathogenic for autoimmunity or an unrelated response to infection or fibrosis (88; 120). One feature that points to a pathological component is the presence of plasma cells within TLS and autoantibodies in BALF that characterize a particular exposure, such as ACPA production in RA subsets associated with smoking or silica exposure.

Autoantibodies

Autoantibodies are a hallmark of idiopathic autoimmune diseases with ANA being a characteristic of systemic autoimmune diseases such as SLE, SSc and RA. As xenobiotic exposure can typically exacerbate both ANA and autoimmune disease in a wide range of autoimmune-prone mice, it can be argued that genetically susceptible backgrounds are generally sensitive to the immunostimulation provided by exposure (70). In addition, xenobiotic exposure can elicit other self-antigen specificities, particular to the type of

exposure (Table 3). The similarities and differences in these responses argues that highly specific responses, can be of diagnostic utility and could yield new mechanistic insights. However there are significant gaps in our current understanding of the spectrum of autoantibody specificities that can be ascribed to xenobiotics, particularly for human exposure.

COPD, even in the absence of overt autoimmune disease, is associated with autoantibodies against many cell and tissue components including ANA. Furthermore, oxidative stress-induced protein modifications including carbonylation or oxidation of protein side chains (88), leads to anti-carbonylated protein antibodies whose titers correlate with the severity of COPD (121). A more significant protein modification for autoimmunity is the citrullination of β - and γ -actins, enolase, fibrinogen, filaggrin and vimentin, most likely mediated by smoking-induced increases in PAD-2 and 4 in the lungs. Consequently, conversion of protein arginine residues to citrulline (89) provides the autoantigens for the induction of ACPAs. Sparks and Karlson (22) have reviewed the association of smoking with preclinical stages of RA and the evidence that the lungs are critical for the development of RA, particularly seropositive RA which includes the ACPA⁺ subset. Although COPD patients without RA can be ACPA positive, the diagnostic significance of ACPAs was considerably more important in the context of RA. Furthermore, a study of 12,950 twins in Sweden found that while non-genetic factors (i.e. environment, lifestyle) are important in ACPA development, the presence of the SE had a significant impact on determining which ACPA-positive individuals developed RA (122). Another study of newly diagnosed, untreated RA patients showed that BALF had higher levels of ACPA compared to serum, and that adaptive immune cells, including B cells, were more likely to be present in bronchial biopsies of early RA patients with anti-cyclic citrullinated peptide (anti-CCP) antibodies (19). This suggests, indirectly, that the lungs are a site of ACPA production.

Consistent with the appearance of autoantibodies in preclinical phases of autoimmune diseases, silicosis can be associated with autoantibodies, including ANA, in the absence of connective tissue disease (21). Autoimmune disease-related specificities such as anti-DNA, anti-Sjögrens syndrome-A/Ro (anti-SS-A/Ro), anti-SS-B/La, anti-centromere, and anti-DNA topoisomerase 1 have been documented in silica-exposed individuals with and without a diagnosable disease such as SLE or SSc. Patients with silica-associated SSc have a greater prevalence of anti-DNA topoisomerase 1 autoantibodies, but fewer patients with silica-associated SSc and SLE had high titer ANA (>1:1,280) than in idiopathic disease. In comparison, autoimmune- and nonautoimmune-prone mice exposed to silica exhibited a broader range of ANA specificities that often included anti-ribonucleoprotein (RNP) and anti-Smith (Sm) (111–113). Silica exposure is also a risk factor for ACPA-positive RA, and this risk increases when coupled with smoking (123; 124). The mechanism of silica-mediated citrullination appears similar to smoking as *in vitro* exposure of cells to SiO₂ nanoparticles induces protein citrullination together with an increase in PAD enzyme activity (91). The finding that silica-exposed Diversity Outbred (DO) mice develop anti-CCP3 antibodies (Mayeux and Pollard, unpublished) also supports this possibility.

Exposure to TCE is also associated with ANA positivity (125). In SSc, solvent exposure including TCE is more likely to be associated with anti-Scleroderma (Scl)-70 autoantibodies

(126). Animal experimentation supports the potential of solvents such as TCE to exacerbate autoimmunity in autoimmune prone mice (95; 125), however evidence of scleroderma-like disease is lacking. TCE exposure of autoimmune prone female MRL^{+/+} mice is associated with anti-MDA- and anti-HNE-protein antibodies together with ANA, anti-ssDNA, and anti-dsDNA-antibodies possibly related to TCE-mediated oxidative stress (95).

Mercury also exacerbates autoantibody responses in lupus prone mice, as well as inducing autoantibodies in non-autoimmune prone strains. In the majority of strains tested, the specificities are primarily ANAs that target anti-chromatin and other anti-nuclear specificities (3). Mercury exposure, similar to many of the aforementioned xenobiotics, can lead to oxidative stress and carbonylation of proteins, including nuclear proteins (96), however antibodies against such modified proteins have not been demonstrated. Instead, it is the affinity of mercuric ions for sulfhydryl groups that has led to our understanding of its most characteristic response, the development of major histocompatibility complex (MHC) restricted antibodies to the nucleolar protein fibrillarin (3). Although mercury binds fibrillarin, human and mouse anti-fibrillarin autoantibodies do not bind a fibrillarin-mercury conjugate. Rather, it is mercury-induced cell death and proteolysis that leads to generation of a fibrillarin fragment that is immunogenic (3). Thus, phagocytosis and proteolysis of cellular material following mercury-induced cell death may be another primary source of neoantigenic determinants for self-reactive T lymphocytes following xenobiotic exposure.

For asbestosis, most of our understanding of autoantibodies comes from exposure to asbestiform amphibole-contaminated vermiculite in the community of Libby, Montana. The profile of ANA specificities in individuals from this area includes anti-dsDNA,-RNP,-SS-A/Ro52, and -Scl-70 (127). Autoantibodies against mesothelial cells have also been observed but anti-CCP and RF are uncommon (57). In C57BL/6 mice asbestos induces ANA including anti-dsDNA and anti-SS-A/Ro52 as well as mesothelial cell autoantibodies (127). ANA has also been induced in rats, but asbestos did not exacerbate disease in two rat models of arthritis (128). Oxidative stress is also a prominent feature in the initiation of asbestos-induced inflammation (129) and it is likely that there are protein modifications or autoimmune responses against modified self-antigens similar to silica.

As the discussion above reveals, autoantibody induction, especially ANA, by xenobiotics is a common observation, but examination of the antigenic specificities often reveals subtle, and sometimes marked, differences between types of exposures (e.g. ACPA in silica but not asbestos exposure) (Table 3). Differences can also be observed between species (e.g. humans versus experimental animals). This is not unexpected given the differences in physical and chemical properties among xenobiotics as well as their biological reactivity and the influence of genetic background. Apart from the linkage of smoking and/or silica exposure to protein citrullination and the development of ACPA, little has been done to examine the role of xenobiotic-induced self-protein modification in xenobiotic-induced autoantibody responses and autoimmune disease. This remains a missed opportunity because such modifications have been shown to generate immunogenic moieties whose cognate antibodies can facilitate epitope spreading and accelerate disease (19; 130).

The MHC link

Associations with the MHC locus, or HLA, constitute the strongest risk factors for autoimmune diseases (131). MHC associations also contribute to xenobiotic-induced autoimmunity particularly the regulation of CD4+ T cells and autoantibody responses. The most convincing association is between smoking, the HLA SE, and ACPA. The HLA SE consists of a common amino acid sequence at positions 70–74 of the HLA-DRB1 molecule with smoking increasing RA susceptibility in those carrying the HLA-DRB1 SE (35). Linkage between HLA-DRB1*15, and smoking increases the risk of MS, and this is enhanced when combined with solvent exposure (47). However, any associated antibody response remains to be identified. The anti-fibrillarin response in SSc is linked to HLA-DRB1 alleles (132), and the possibility of increased exposure to mercury (133). This is an interesting probability because the anti-fibrillarin response in mercury exposed mice is MHC linked (3), unlike responses to other nuclear antigens such as chromatin which are not restricted to mercury-induced autoimmunity. In silica-exposed miners, anti-Scl-70 positivity is more strongly associated with DRB1*0300 and DQB1*0201 alleles than in idiopathic SSc (134). In Japanese silicosis patients HLA-DQB1-0402 is more common in anti-DNA topoisomerase I positive patients than anti-DNA topoisomerase I negative patients or healthy controls (135).

These observations hint at the possibility that post-translational modification of self-antigens and MHC restriction are important factors in xenobiotic-induced autoantibody responses. This is best exemplified by the relationship between the SE, smoking and ACPA+ RA. In a similar vein, xenobiotic-induced cell death and proteolysis appears to play an important role in the mercury-induced MHC-restricted anti-fibrillarin response. A more concerted effort is needed to identify relationships between MHC and autoantibody responses following xenobiotic-exposure, and to determine the nature of the MHC antigens, and the contribution of autoantibodies to disease pathogenesis.

Disease complexity

As noted earlier, apart from epidemiological studies of exposure and disease association, diagnosis of autoimmune diseases does not discriminate between idiopathic and xenobiotic induced disease (69). Additional issues stem from poorly performed cohort studies, as well as the timeframe, often in years, between exposure and diagnosis of disease. Comorbidities, such as COPD, silicosis or asbestosis, although important clues in identifying a possible trigger for autoimmunity, also add to the complexity of understanding disease mechanisms.

The complexity of disease features that may be present in an exposed population is typified by screening of more than 6,000 community members in Libby, Montana (57). This community has a history of mining and use of asbestiform amphibole contaminated vermiculite, and almost 14% have been diagnosed with an autoimmune disease. The evolution of autoimmune diseases in these patients is atypical, with mixed diagnostic criteria suggesting that there is not one underlying autoimmune disease associated with the exposure. Moreover, the 1:1 male-to-female ratio is highly unusual given the prominent female predisposition of systemic autoimmune diseases such as SLE, SSc, and Sjögren's Syndrome (SS) (66). Other studies also reveal differences in exposure related clinical

features. Male miners, exposed to silica dust, with SLE had considerably less arthritis and photosensitivity compared with idiopathic SLE, and reduced prevalence of discoid lesions. However, the control group was not matched by age or sex, thus it is unclear if the differences reflect silica exposure or sex and/or age differences (21). In a more carefully controlled study, silica-exposed SLE patients were found to have reduced prevalence of anemia and leukopenia compared to idiopathic disease (21).

An important pathogenesis issue requiring resolution is the mechanism by which inflammation and autoreactive TLS development in an exposure site such as the lung leads to disease pathology in distant organs such as the joint in RA or kidney in SLE (72; 136; 137). Several hypotheses have been proposed including the existence of shared antigenic targets between the lungs and target tissues, epitope spreading to target specific antigen(s), and immune complex deposition in target tissues (137). Citrullinated proteins are found in the RA joint, suggesting that antigenic targets exist but it still remains unclear how events in the lung lead to pathogenesis in a target tissue such as the joint (136). This is one of the many mechanistic issues that remain to be clarified concerning xenobiotic-induced autoimmune disease.

CONCLUSION

An important commonality of xenobiotic exposures associated with human autoimmunity and autoimmune diseases is the lung as the major site of exposure. Another important feature is that the exposures are often protracted leading to persistent tissue damage. The most disease relevant exposures are also particulate material, which hampers clearance and degradation thereby facilitating a state of persistent inflammation. The failure to resolve this chronic inflammatory response allows the appearance of TLS characterized by aggregates of B and T cells which may become sites of autoreactivity. Coincident with these steps are stress response related events that lead to post-translational modification of self-proteins and the production of autoantibodies against these neoantigens. Under appropriate gene-environment interactions these responses can have diagnostic specificity such as the ACPA response in seropositive RA mediated by the HLA SE. Numerous questions still remain concerning the disease relevance not only of the xenobiotic-induced chronic inflammatory response but also the pathogenic role of pulmonary TLS particularly as to how they relate to the development of autoreactivity against modified self-proteins. Greater insight into the molecular and cellular requirements that govern this process, especially those that might distinguish pre-clinical autoimmunity from clinical autoimmune disease, may allow determination of the significance of xenobiotic exposures in human autoimmune disease.

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SUMMARY POINTS

1. Several environmental exposures, including smoking, silica dust, and solvents, have been confidently linked to autoimmune diseases. Other exposures, such as mercury and asbestos, result in autoimmunity including autoantibodies but with insufficient evidence of clinical disease. The latter may be important for identifying factors required for progression from pre-clinical to clinically significant autoimmune diseases.
2. The lungs are an important site of exposure to environmental agents linked to autoimmune diseases. The association of smoking and silica dust exposure to pre-clinical features of rheumatoid arthritis supports the growing appreciation that mucosal and epithelial cell surfaces are key sites in the initiation of autoimmune reactions.
3. IL-1 and the inflammasome are major contributors to early pulmonary inflammation mediated by the innate immune system following exposure to smoke, silica, and asbestos. Toll-like receptor signaling via MyD88, activation of Th1 CD4+ T cells and interferons also appear important for the subsequent adaptive immune response.
4. Post-translational modification of self-proteins is a shared feature of xenobiotic exposure. Several different mechanisms are operative but the most clinically important is citrullination.
5. The chronic inflammation of the lung following environmental exposure can result in formation of tertiary (or ectopic) lymphoid structures. These are comprised of focal areas of T and B cells and follicular dendritic cells that could provide a microenvironment for survival and maturation of autoreactive cells including autoantibody producing B cells.
6. Antibody responses in environmental-induced autoimmunity and autoimmune diseases include autoantibody responses against modified self-proteins. Of these, the anti-citrullinated protein antibodies found in seropositive rheumatoid arthritis following smoking or silica dust exposure have clinical and diagnostic significance.
7. The major histocompatibility complex (MHC) locus, or human leukocyte antigen (HLA), is linked to autoantibody responses accompanying environmental exposures. This is best exemplified by the association between the HLA-shared epitope, smoking, and anti-citrullinated protein antibodies. This association also exemplifies the importance of gene-environment interactions in environmental-induced autoimmunity.

FUTURE ISSUES

1. Are tertiary lymphoid structures (TLS) a common feature of all chronic environmental exposures in the lung, or are they restricted to exposures associated with autoimmune diseases?
2. Do environmental exposure induced TLS only provide a B cell survival niche, or are they important in differentiation and expansion of autoreactive T and B cells?
3. What post-translational protein modifications occur following environmental exposures, do they elicit pathogenic autoantibody responses, and do they have diagnostic potential?
4. Are major histocompatibility (MHC) restrictions common to environmental-induced autoantibody responses against post-translationally modified antigens?
5. Can greater understanding of the gene-environment interactions of environmental-induced autoimmunity help discriminate pre-clinical autoimmunity from clinical autoimmune disease?
6. Can studies of the exposome, microbiome and epigenetics help us understand the mechanisms responsible for environmental-induced autoimmunity?

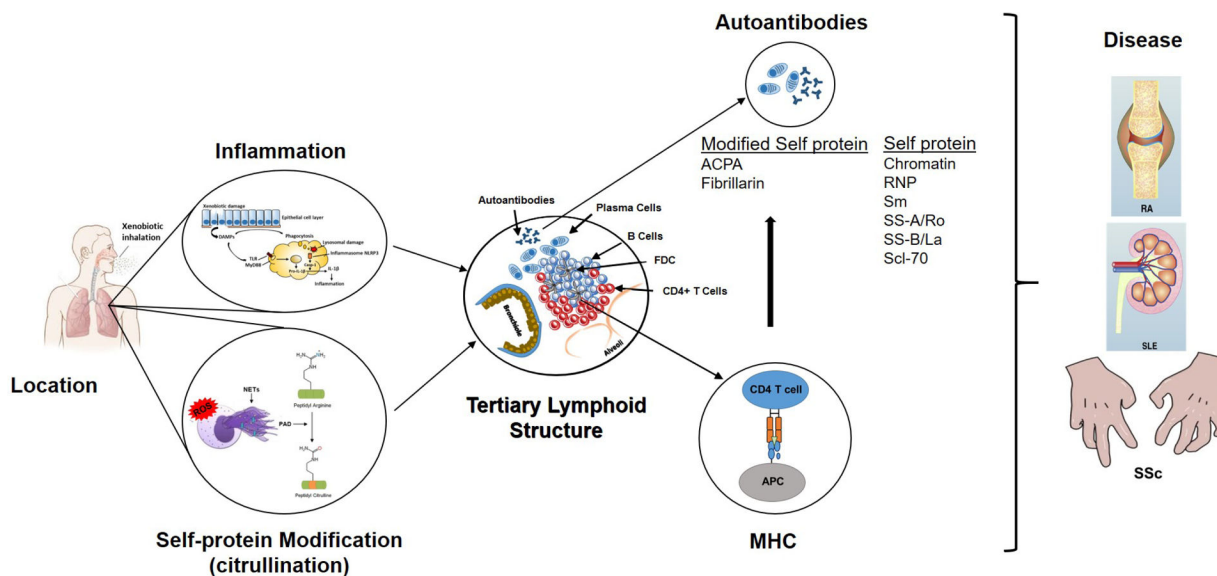


Figure 1.

Steps in the development of xenobiotic-induced autoimmunity and autoimmune disease. Information in the figure summarizes the major points discussed in the text as they relate to the seven hypothesized steps leading to autoimmunity or autoimmune disease following xenobiotic exposure. The lungs serve as a common site of xenobiotic exposure. Chronic exposure results in an inflammatory response beginning with cellular and tissue damage, DAMP activation of PRR including TLRs and expression of inflammatory cytokines. Phagocytosis of DAMPS, including particulate xenobiotics, leads to lysosomal damage, inflammasome activation and processing of pro-IL-1 β , further enhancing inflammation. Coincident with the inflammatory response are stress response related events, including NETosis and release of PAD enzymes, that leads to post-translational modification of self-proteins, particularly citrullination, and production of neoantigens. The ensuing chronic inflammatory response results in development of TLS comprised of accumulations of B cells within surrounding CD4⁺ T cells, FDCs and plasma cells. Processing and presentation of self and modified-self proteins, particularly in the context of MHC restriction, leads to autoantibodies. Under appropriate gene-environment interactions these autoantibodies can have diagnostic specificity such as the ACPA response in RA. The culmination of these steps is the development of disease pathology in distant tissues such as the joint in RA, the kidney in SLE and sclerodactyly, a skin manifestation of SSc.

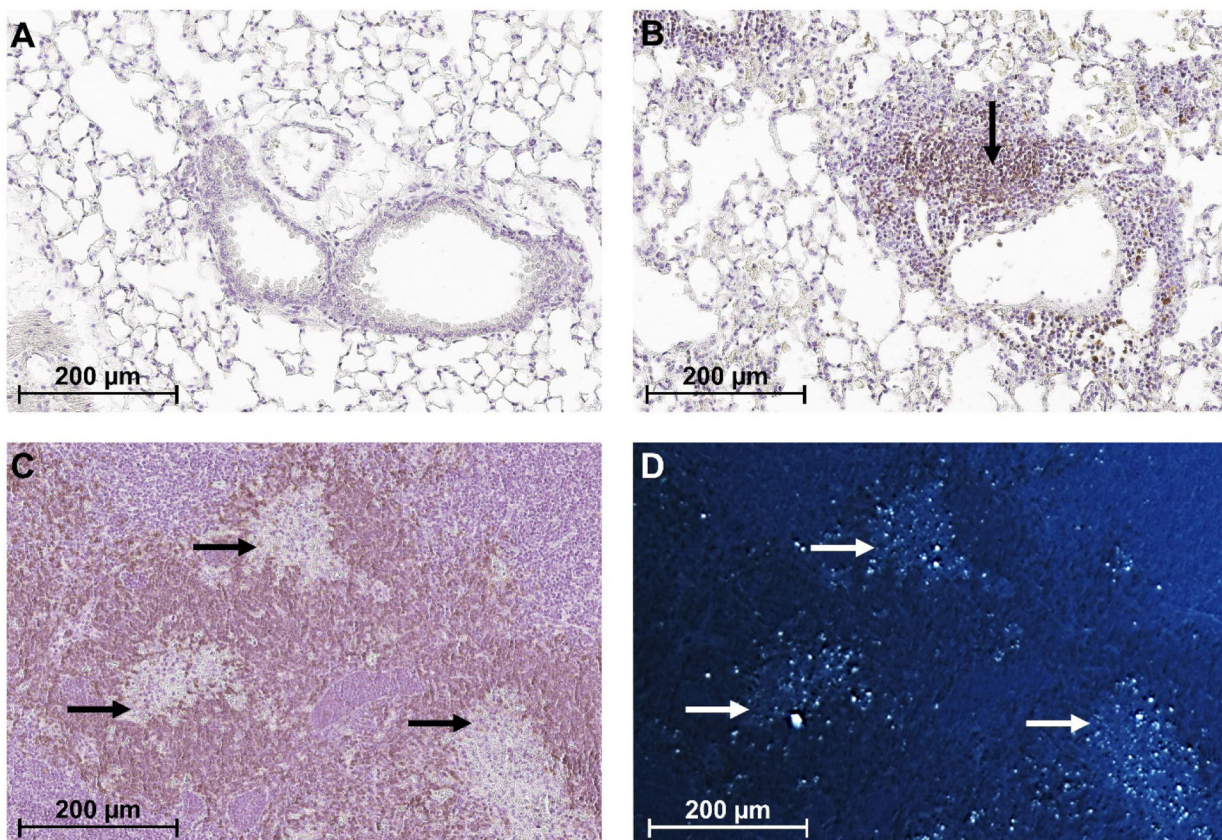


Figure 2.

B cell aggregates in the lung and lymph node of silica exposed mice. Representative images of lung sections from Diversity Outbred (DO) mice 12 weeks following one 50 μ l transoral instillation of (A) PBS or (B) 5 mg of crystalline silica (Min-U-Sil 5). IHC staining indicates B220+ B cells (black arrow in B) surrounding blood vessel in silica, but not vehicle, exposed mice. Tracheobronchial lymph node of silica exposed DO mouse imaged using (C) bright-field and (D) polarized microscopy shows areas of cell death and/or granulomas (black arrows in C) containing silica (white arrows in D) surrounded by B cells. B220+ cells are stained with DAB (brown) and hematoxylin (purple) was used as the counterstain.

Table 1.

Xenobiotic Exposures Linked to Human Autoimmunity and Autoimmune Disease.

Xenobiotic	Exposure	Exposure Site	Autoimmune Diseases or Autoimmunity ^a	Selected References
<i>Confident^b</i>				
Silica	Particulate	Lungs	SLE, SSc, RA, ANCA-related vasculitis	(8, 20, 25)
Smoking	Chemical/Particulate	Lungs	Seropositive RA	(8, 19)
Solvents	Liquid/Vapor	Skin/Lungs	SSc	(8, 39)
<i>Likely^b</i>				
Smoking	Particulate	Lungs	SLE, MS	(8)
Solvents	Liquid/Vapor	Skin/Lungs	MS	(8, 45)
<i>Insufficient evidence^b</i>				
Mercury	Vapor (gold amalgamation), Lotion (skin cream), Dietary (sea food), Particulate (dental amalgam)	Lungs, Skin, GI tract	autoantibodies, cytokines, nephrotic syndrome	(3, 4, 8)
Asbestos	Particulate	Lungs	autoantibodies, atypical rheumatological symptoms	(8, 54, 57)

^aExposures leading to autoimmunity without clinical disease are mercury and asbestos.

^bCategories of *Confident*, *Likely*, and *Insufficient Evidence* were described by Miller et al (8) essentially as follows.

Confident: Includes exposure disease associations from multiple studies from different populations using different designs; robust evidence of an overall association as identified by high-magnitude risks or the use of high-quality or established exposure assessment methods; evidence of an exposure-response gradient; and/or evidence of effect modification by disease subtype or genetics that supports biologic plausibility.

Likely: collections of research studies missing important elements, such as clarification of the temporal association between exposure and onset of an autoimmune disease, or less consistent results or were based on fewer studies.

Insufficient evidence: reported studies were too limited in design or power to allow conclusions to be drawn.

Table 2.

Postulated mechanisms for xenobiotic-induced autoimmunity and autoimmune disease.

Type of Exposure	Silica	Smoking	Solvents	Mercury ^a	Asbestos
Location	Lung	Lung	Lung/Skin	Lung/Skin/GI Tract	Lung
Inflammatory responses	IL-1 α , IL-1 β Inflammasome MyD88, IFN- γ , IL-17, type I IFN, IRF7 NETosis	IL-1 β Inflammasome IL-6, IL-8, TNF- α MyD88, NETosis	ND	Endosomal TLRs IL-6, TNF- α , IFN- γ NETosis	IL-1 β Inflammasome
Self-Protein Modifications	Citrullination Carbonylation	Citrullination Carbonylation	Carbonylation Nitration	Proteolysis Carbonylation	Carbonylation Citrullination
Tertiary Lymphoid Structure	Yes	Yes	ND	Lymphoid Accumulations	Alveolitis
Autoantibodies	ACPA	ACPA	Scl-70	Fibrillarin	Mesothelial cell
MHC association	ND	HLA-SE	HLA-DRB1	H-2 ^s	ND
Diseases	SLE, SSc, RA, ANCA-related vasculitis	Seropositive RA SLE, MS	SSc	Nephrotic Syndrome	Rheumatological Symptoms

A comparison of the hypothesized steps leading to autoimmunity or autoimmune disease following exposure to different xenobiotics. Silica, smoking, and solvents exposures are confidently linked to various autoimmune diseases while mercury and asbestos exposures although linked to features of autoimmunity, evidence for causation of autoimmune disease is insufficient.

^aIncludes information from animal studies.

ND, not determined.

Table 3.

Autoantibodies in Xenobiotic-Induced Autoimmunity and Autoimmune Disease.

Xenobiotic	Autoimmune Diseases or Autoimmunity ^a	Idiopathic ^b	Xenobiotic Human ^c	Xenobiotic Experimental ^c	Selected References
<i>Confident</i>					
Silica	SLE, SSc, RA	ANA DNA, Sm, RNP, SS-A/Ro, SS-B/La Centromere, DNA topoisomerase I RF <i>ACPA</i> ^d	ANA DNA SS-A/Ro SS-B/La Centromere, DNA topoisomerase I <i>ACPA</i>	ANA Sm RNP dsDNA RF	(21, 25, 30, 111, 112, 113) (25, 29) (123)
Smoking	Seropositive RA	RF <i>ACPA</i>	<i>ACPA</i>		(19)
Solvents	SSc	Scl-70	Scl-70	<i>MDA</i> <i>HNE</i> ssDNA dsDNA	(126) (95)
<i>Insufficient evidence</i>					
Mercury	autoantibodies, cytokines, nephrotic syndrome	NA	ANA	ANA Fibrillar Chromatin	(3)
Asbestos	autoantibodies, atypical rheumatological symptoms	NA	ANA dsDNA RNP SS-A/Ro52 Scl-70 Mesothelial cell	ANA dsDNA SS/A-Ro52 Mesothelial cell	(127)

Note: This table is not a comprehensive listing of all autoantibodies found in a particular idiopathic autoimmune disease. Rather it is a comparison of autoantibody responses found with a specific xenobiotic exposure and whether the same response has been identified in idiopathic disease.

^aExposures leading to autoimmunity without clinical disease are mercury and asbestos.

^bIdiopathic disease means no known association with an environmental exposure.

^cAntibodies in humans or experimental animal models following exposure.

^dIn italics, antibodies against post-translational protein modification.

^eAbbreviations: NA, not applicable.