

Autoreactive B cells in rheumatoid arthritis Kristyanto, H.

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General introduction

ACPA-positive rheumatoid arthritis: the role of autoreactive B cells and targeting strategies

1 Rheumatoid arthritis and its burden

Rheumatoid arthritis (RA) is a chronic autoimmune disorder which is characterized by persistent inflammation of the joints leading to progressive joint destruction, deformity, and immobility. The American College of Rheumatology and the European League Against Rheumatism joint working group has set out criteria for the classification of patients with at least 1 joint with clinical synovitis which cannot be better explained by another disease. These patients are classified as having 'definite RA' when fulfilling at least 6 criteria out of 10 which include the number and kind of joints involved (maximal score of 5); the presence and levels of RA-related autoantibodies, i.e. Rheumatoid Factor (RF) and Anti-Citrullinated Proteins Antibodies (ACPA) (maximal score of 2); the evidence of systemic inflammation as shown by increased acute-phase reactants (score 1); and persistence of symptoms (for at least 6 weeks, score 1).¹ The signs and symptoms of RA include painful, warm and/or swollen joints, joint stiffness particularly after inactivity, fatigue, and occasionally fever and loss of appetite. Moreover, RA is associated with increased risk of lung involvement and cardiovascular morbidity as well as premature death.^{2,3,4,5}

There are six histologic characteristics of rheumatic joints: (1) increased vascularization due to vasodilation and neo-angiogenesis; (2) aggregation of fibrin in the joint cavity, forming rice bodies; (3) accumulation of neutrophils in the synovial fluid; (4) infiltration of synovial stroma by dense perivascular inflammatory cells, including B cells, T cells, plasma cells and macrophages; (5) osteoclast activity in underlying bone; and (6) formation of pannus tissue, a mass of synovium and synovial stroma which grows over the articular cartilage. Osteoclast activity and pannus formation lead to bone and cartilage erosion in severe RA, respectively.⁶

RA affects between 0.3% and 1% of the population world-wide, predominantly women, and is more frequent in developed countries.⁷ As RA tends to develop during productive years of adulthood, the societal impact and economic burden are particularly high. It is estimated that RA costs the National Health Service of the UK approximately £560 million a year in health care costs with additional £1.8 billion a year for work-related absence and disability.⁸

2 RA subsets: ACPA positive and negative

RA originates from a heterogenous immune-pathogenesis that initiates, promotes and maintains the disease. One of the serological markers of a distinct subset of RA

is ACPA. ACPA discriminate two subsets of RA which differ in genetic background and clinical course.^{9-11,12,13} ACPA are present in 46% to 80% of RA patients and serve as highly specific biomarkers for RA.^{14,15} These autoantibodies are detectable years before the onset of arthritis, are associated with early, rapidly progressive disease, and prognosticate erosive joint destruction in established RA.^{16,17} In patients who achieve drug-free remission or treatment reduction, relapses occur more frequently in ACPA-positive individuals, suggesting persistent immunological dysregulation in spite of clinical improvement.¹⁸ Due to rapid disease progression and the lower chance of reaching sustained drug-free remission, ACPA-positive RA patients have higher annual medical expenditures compared to their ACPAnegative counterparts.¹⁹ The focus of this thesis will be on ACPA-positive RA.

Next to clinical, epidemiological and genetic differences, there are also distinctive features in the synovium and circulating chemokines that differentiate these two subsets of RA. The synovium from ACPA-positive RA shows higher numbers of B and T cell infiltrates compared to ACPA negative RA, with the formation of germinal centre-like follicular structures in some but not all patients with ACPA-positive disease. Moreover, ACPA positive RA patients have higher serum levels of CXCL13, a B cell chemotactic factor.²⁰ ACPA-positive RA patients also respond better to B cell depletion therapy and T cell co-stimulation inhibitor compared to ACPA-negative patients.^{21,22} These differences suggest a pivotal role for B and T cells in the pathogenesis of ACPA-positive RA.

3 Antibodies

Antibodies or immunoglobulins (Ig) are Y-shaped proteins that collectively form the humoral part of the adaptive immune response. In physiological terms, antibody production aims to neutralize pathogens. Antibodies are composed of a pair of heavy and light chains, which form two fragment antigen-binding (Fab) variable regions and one fragment crystallizable (Fc) region. While the Fab region is remarkably diverse to allow for specific binding towards divergent pathogens, the Fc region has a constant structure with limited isotypes and subclasses. Antibody isotypes include early produced IgM, scarcely circulating IgD, most prevalent IgG, mucosaassociated IgA, and parasite-fighting IgE.

The Fc region determines the effector functions of an antibody by its distinct capacity to activate complement and by binding specific Fc receptors (FcR) expressed by effector cells such as macrophages, natural killer cells, neutrophils and mast cells. Antibody binding towards different FcR is regulated by the presence of

different glycan groups in the Fc regions.²³ Upon binding to their cognate antigens, antibodies form immune complexes (ICs) that enhance complement and FcR activation. Consequently, pro-inflammatory cytokines and other mediators that contribute to inflammation are produced. In addition to this Fc tail-dependent effector cell activation, antibodies can also function directly by neutralizing the infective capacity of pathogens, by agonistically activating receptors or by disturbing intermolecular interaction.

IgM and IgA antibodies have a half-life of five to six days while IgG has a 21-day half-life due to recycling by the neonatal FcR (FcRn). Antibody effector functions are also regulated by the presence of soluble FcR which binds to the antibody without subsequent FcR signaling and by binding to the inhibitory FcR, i.e., Fc-gamma-receptor 2b or CD32b.²⁴⁻²⁶

Antibodies are produced by two distinct late-phase differentiated types of B cells, namely plasmablasts and plasma cells which have different characteristics. Plasmablasts are rapidly produced, short-lived and proliferating cells while plasma cells can either be short-lived or long-lived and can reside as quiescent producers of antibodies in tissues and bone marrow.²⁷ Both cell types are the product of B cell differentiation in the germinal centre at secondary lymphoid organs where B cells undergo substantial changes in B cell receptor (BCR) variable domain genes due to somatic hypermutation. In addition, an alternative pathway of B cell extrafollicular development supports rapid production of antibodies at the site of inflammation. During the course of adaptive humoral immune responses, the antibody repertoire evolves to increase its binding capacity towards the cognate antigens, to broaden the binding targets for efficient clearance of pathogens, and to diversify the isotype usage in order to expand the effector functions of the antibodies. These processes, which are respectively termed as avidity maturation, epitope spreading and class switch recombination, occur primarily in the germinal center.

4 ACPA

ACPA are a group of antibodies that recognize a variety of self-proteins including fibrinogen, filaggrin, vimentin, myelin basic protein, histone and alpha-enolase in their citrullinated forms. The commonality of the majority of these autoantibodies is their binding to a single synthetic peptide, i.e. CCP2. When tested for their fine specificity, most ACPA bind with varying affinity to various citrullinated proteins, indicating that they cross-react. In addition, ACPA can also recognize and thus frequently cross-react with proteins or peptides that undergo other post-translational



modifications.28,29

Citrullination or deimination is a post-translational modification of a positively charged arginine residue to the neutrally charged citrulline residue in a peptide or protein backbone, catalysed by a family of enzymes called peptidylarginine deiminases (PADs). The alteration of amino acid charge leads to changes in protein folding and molecular interactions. PADs are expressed in five isoforms where each one has tissue-specific distribution and distinct cellular substrates.^{30,31} PAD function is tightly regulated by calcium homeostasis. Under physiological conditions, PADs play roles in the regulation of gene transcription, skin keratinization, brain plasticity, the formation of neutrophil extracellular traps and the regulation of chemokines function such as CXCL8 and CXCL12.^{32,33} Aberrant expression and activation of PADs are observed in many pathologic conditions such as cancer; neurodegenerative diseases including Alzheimer's and prion diseases; and autoimmune diseases such as multiple sclerosis, psoriasis and RA.³⁴

Despite ubiquitous citrullination in many physiologic and pathologic conditions, the presence of ACPA is strongly associated with RA. Moreover, there is so far only a limited number of citrullinated proteins that have been identified as binding targets of ACPA. It is still unclear whether ACPA targets *in vivo* are the physiologic substrates of PADs.³⁵ It has been hypothesized that aberrant citrullination by PADs may lead to new citrullinated epitopes that are not tolerized by the immune system. Subsequently these new epitopes could evoke an autoimmune response in susceptible individuals. As rheumatic joints contain numerous citrullinated proteins, it is likely that ACPA can bind to antigens in the joint *in vivo* and evoke inflammatory immune responses.^{36,37}

5 Effector functions of ACPA

Autoantibodies can exert their activity by directly binding to organ-specific targets which induces agonistic or inhibitory effects on a particular receptor, thereby activating alternative signalling pathways or disturbing essential molecular interactions.³⁸ These organ-specific autoantibodies play central roles in autoimmune diseases such as neuromyelitis optica, Grave's disease, bullous pemphigoid, myasthenia gravis, alopecia areata, and a subset of Sjögren's syndrome.³⁹⁻⁴⁴ Other autoantibodies do not have organ-specific targets but bind to ubiquitous autoantigens which lead to systemic disease manifestations. An example of these autoantibodies are anti-double stranded DNA antibodies which contribute to systemic lupus erythematous through IC deposition in the kidney, complement

activation and subsequent tissue injury. Unlike these autoantibodies, however, the disease-causing roles of ACPA are still under debate.

The study of ACPA pathogenicity in human RA is challenging, partly due to the strict definition of ACPA which are defined as antibodies that bind to citrullinated antigens but not to the unmodified "parent" peptide or protein. This means that the isolation of ACPA for functional studies requires strict controls that ensure binding of purified ACPA to citrullinated antigens but not arginine control antigens. Moreover, ACPA isolation could generate antibody aggregation, immune complex formation, co-purification of rheumatoid factor and introduction of pyrogens which can lead to various false positive immunological and non-immunological read-outs.⁴⁵ Consequently, there has been considerable debate as to the reproducibility of data with regard to ACPA effector functions derived from isolated polyclonal ACPA preparations.

To circumvent some of the issues regarding the isolation of patient-derived polyclonal ACPA, monoclonal ACPA have been produced and tested for their pathogenicity. Infusion of monoclonal ACPA which bind citrullinated fibrinogen into an RA mouse model induced worsening of arthritis and increased interleukin-6 levels in the joints and serum.⁴⁶ In addition, Fab fragment of two monoclonal ACPA, namely D10 and B02, have been shown to induce osteoclast activation and bone resorption through binding with a PAD-mediated citrullinated target on the osteoclast precursors which leads to autocrine IL-8 production.⁴⁷ This study suggests a direct activating role of ACPA by binding to a citrullinated target on osteoclast precursors. Moreover IL-8 produced by ACPA monoclonal-activated osteoclast induces pain behaviour in mice through activation of sensory neurons.⁴⁸ One fundamental caveat of these studies is the uncertainty of whether these D10 and B02 monoclonals are true ACPA since previous work describing the generation of these monoclonals have been retracted due to the inability of these monoclonals to bind citrullinated peptides.⁴⁹ Furthermore, two cohorts of arthralgia and early RA did not show independent association between ACPA and bone erosion or pain.^{17,50} In the arthralgia cohort, it is suggested that ACPA indirectly increases bone erosion through the induction of local inflammation in RF-dependent manner.⁵⁰ These findings indicate the putative role of ACPA as a part of inflammatory network in the development of RA.

The evidence is clear that ACPA positivity is associated with rapidly progressing, severely destructive RA. Moreover, in individuals without RA, ACPA positivity is already associated with joint complaints.⁵¹ ACPA effector functions have been

researched extensively *in vitro*. It has been reported that ACPA IC bind to Fcgamma-receptor IIa on macrophages and induce tumor necrosis factor alpha (TNF α) production.⁵² As the ACPA response matures prior to disease onset, ACPA may bind to a growing number of diverse citrullinated proteins. These characteristics enable ACPA to form large and numerous ICs which could potently stimulate TNF α production by macrophages.⁵³

Moreover, ACPA often co-exist with another autoantibody group, namely the anti-IgG antibodies rheumatoid factor (RF). RA patients who have both autoantibodies exhibit higher disease activity and systemic inflammation compared to patients with either autoantibodies.⁵⁴ *In vitro* studies show that IgM RF is able to augment the pro-inflammatory cytokine response of ACPA-IC-activated macrophages.^{54,55} Besides binding to and activating macrophages, ACPA IC can also bind to activated neutrophils which are abundant in RA synovial fluid.⁵⁶ This binding can intensify joint inflammation. Moreover, IC in the joint can activate FcR on the neurons which induces pain sensation.⁵⁷ Apart from activating effector cells, ACPA are also able to activate complement systems *in vitro*.⁵⁸ Accordingly, the number of individual synovial fluid ACPA reactivities in the IC, but not monomeric ACPA in the free form, associate with joint inflammation and destruction.⁵⁹ These findings support inflammation inducing and/or enhancing roles of ACPA primarily via the formation of IC, in particular in the synovial fluid.

Despite these findings, the clinical outcomes of RA are not mirrored by the levels of these autoantibodies. For instance, clinical benefits of medication are generally not coupled with a decrease in ACPA levels.⁶⁰ Moreover, plasmapheresis was shown not to benefit chronic RA.⁶¹ In addition, the presence of ACPA alone is not associated with bone erosions in the patients with clinically suspect arthralgia.⁵⁰ Therefore, it is possible that ACPA might propagate joint inflammation once it is started and that the underlying immune dysregulations play a more dominant role in the establishment of synovitis, which is the main clinical correlate of RA.

6 The evolution of the ACPA response during RA development

ACPA can be detected years before the onset of RA. During this period, the ACPA response is not stable and can also revert to a negative state.⁶² In some individuals, however, it undergoes substantial changes. Most of the knowledge about the evolution of the ACPA response comes from the study of unaffected first degree relatives (FDR) of RA patients and from blood donors who later developed RA. FDR have a higher prevalence of ACPA positivity than the general population and share

genetic as well as environmental risk factors for RA, including a high prevalence of HLA-SE alleles and, in some populations, smoking.⁶³ FDRs are more likely to have joint symptoms than healthy controls, without fulfilling the criteria of definite RA.⁶⁴ In this "pre-RA" stage, ACPA titres are generally low with more limited epitope recognition and isotype usage compared to established RA.⁶⁴⁻⁶⁶ Notably, the most prominent ACPA class at this stage is IgA, indicating that a mucosal trigger could breach the tolerance towards citrullinated antigens. ACPA-positive FDR gradually expand their ACPA binding targets until the onset of disease, indicating a maturation of the ACPA response through epitope spreading.⁶⁷ Accordingly, unaffected FDRs who harbour ACPA with a broad epitope recognition profile are more likely to develop RA in the future than those with limited recognition profile.^{63,66}

ACPA avidity also increases over time until disease onset. Thereafter, no further avidity changes take place.⁶⁸ Compared to antibodies against vaccination antigens such as tetanus toxoid, however, ACPA from patients with established RA have substantially lower avidity despite undergoing extensive somatic hypermutation.^{69,70} Another major difference between ACPA and protective antibodies is that the majority of ACPA IgG contain glycans in their variable domain (VD).⁷¹ This unique feature is thought to result from a selective advantage inferred by the introduction of N-glycosylation sites during the development of ACPA-expressing B cells.^{70,72} This modification is an acquired feature of the ACPA response which occurs before the onset of disease. Generally, unaffected FDR exhibit low ACPA IgG VD glycosylation. However, those with extensive VD glycosylation are six times more likely to develop RA in the future.⁷³ ACPA IgG also undergo changes in the glycosylation pattern at the Fc tails at the onset of RA. Based on murine studies, it is assumed that the newly acquired Fc tail glycosylation pattern renders these autoantibodies more inflammatory.⁷⁴

The evolving nature of the ACPA response before the onset of disease can be explained by a two-hit theory (Figure 1a and 1b). In this model, the first hit occurs when tolerance towards citrullinated antigens is breached, resulting in the appearance of a limited ACPA response without clinical manifestations. This event is likely to occur in the mucosal organs. The following, second hit triggers maturation of the ACPA response, resulting in increased titers, avidity maturation, epitope spreading, isotypes diversification and the acquisition of VD glycosylation prior to disease onset. These events require close interaction between ACPA-expressing B cells and T cells in at least two discrete episodes that are reminiscent of a prime-boost strategy during vaccination. Eventually, this conceptual model might be too



Figure 1. The development and roles of ACPA-expressing B cells in the aetiopathogenesis of RA (A). ACPA-expressing B cells develop due to the breach of tolerance towards citrullinated selfantigen, resulting in the production of low affinity ACPA. (B) ACPA-expressing B cells are reactivated due to several environmental and genetic factors with the help of T cell, resulting in activated memory phenotype and maturation of pro-inflammatory ACPA. (C) Mature ACPA, together with activated macrophages and synovial fibroblasts, elicit the development of joint inflammation resulting in undifferentiated arthritis. (D) ACPA-expressing activated B cells migrate to the synovium, propagate and sustain inflammation, resulting in RA.

simplistic, however, as the dynamics of changes in ACPA characteristics suggest that multiple events/hits are required that induce a step-wise maturation of the ACPA-response ('multiple hit model'). Nonetheless, upon formation of a "mature" ACPA response, inflammation and other clinical manifestations may follow.

7 The Aetiology of ACPA-positive RA

Numerous pre-clinical, epidemiological, genetic, epigenetic and therapeutic studies have provided evidence and clues on the distinct causes of the development of ACPA and ACPA-positive RA. Intricate interactions between environmental and genetic factors are presumably responsible for the breach of tolerance towards citrullinated autoantigens, which gives rise to ACPA positivity and the development of RA which is preceded by an enhanced ACPA response (Figure 1b).

7.1 The breach of tolerance towards citrullinated autoantigens

In a large population-based cohort study from the Netherlands, ACPA were detectable in around 1% of the general, mostly Caucasian population. ACPA positivity is more prevalent in women, the elderly, smokers, people with joint complaints, RA patients and first-degree relatives of patients with rheumatic diseases. When RA patients are excluded, ACPA positivity still associates with older age, smoking and joint complaints.⁵¹ In a Swiss cohort, the presence of ACPA peaks in women around perimenopausal age which indicates that pronounced hormonal decline may support the breach of tolerance towards self-antigens, including citrullinated proteins.⁷⁵ Accordingly, hormone replacement therapy associates with a reduced risk of developing ACPA-positive RA but not with ACPA-negative RA in post-menopausal women.⁷⁶

In a Swedish twin study, heavy cigarette smoking is significantly associated with ACPA positivity without RA.⁷⁷ Similarly, exposure to air pollutants such as sulphur dioxide is weakly associated with the presence of ACPA in a large Canadian cohort.⁷⁸ Indeed, smoking has been shown to increase protein citrullination in the bronchial mucosal and alveolar compartments as well as to alter cytokine balance.^{79,80} Based on these epidemiological studies, ageing, the decline of ovarian function and mucosal exposure to air pollutants are linked to the breach of tolerance towards citrullinated self-antigens. These associations could be explained by the accumulation of citrullinated self-proteins related to ageing and smoking as well as by the dysregulation of immunological tolerance related to endocrine changes and immunosenescence.⁸¹⁻⁸⁵



The association of autoimmunity with smoking, ageing, female gender as well as endocrinological transition is, however, not exclusive for the development of ACPA. Data from multinational cohorts showed that smoking is associated with concurrent multiple RA-associated autoantibodies and not with a specific autoantibody.⁸⁶ Smoking is also associated with the presence of other disease-specific autoantibodies including anti-myeloperoxidase and anti-topoisomerase I antibodies.^{87,88} Moreover, many autoimmune diseases are more prevalent in women while the course of the disease is tightly linked to major female endocrinological changes, namely puberty, pregnancy and menopause. These life events are known to have considerable influence on both the innate and adaptive immune systems.⁸⁴ Ageing, on the other hand, is associated with the involution of the thymus, the major T cell central tolerance organ; a decrease in B cell lymphopoiesis and quality control; and increased levels of autoantibodies.⁸⁹⁻⁹³

As the B cell receptor (BCR) repertoire originates from random recombination of gene segments in the bone marrow, mature naïve B cells that recognize citrullinated antigens may arise stochastically. Moreover, new evidence shows that some germline encoded B cell receptors already have propensity to weakly bind citrullinated and other post-translationally modified proteins.^{46,94} These ACPAexpressing B cells may subsequently get selected to become plasmablasts that eventually produce low levels of ACPA as a result of increased citrullination dysregulation of immunological tolerance. Autoimmunity and against citrullinated antigens is, then, born. At this stage, however, the ACPA response is still 'immature'. The emergence of ACPA is necessary but not sufficient for the development of ACPA-positive RA. Accordingly, some ACPA-positive individuals may never develop RA in their lifetime, and without additional triggers and/or genetic predisposition the ACPA response is likely to disappear again at this stage.⁶² The development of autoimmune disease from autoimmunity can take several years and requires interaction between additional genetic and environmental factors and, in particular, the presence of specific helper T cells, as indicated by HLA association which is restricted to ACPA-positive RA but not ACPA only.

7.2 The development of ACPA-positive RA 7.2.1 Environmental factors

Bacterial and viral dysbiosis has been hypothesized to induce ACPA-positive RA in genetically susceptible individuals.⁹⁵⁻⁹⁷ Some studies link ACPA-positive RA with severe chronic periodontitis, an inflammation of the gums and supporting structures

of the teeth caused by bacterial dysbiosis. Both RA and periodontitis share similar risk factors, namely the HLA-DRB1 shared epitope, smoking, and ageing.⁹⁸

Two periodontal pathogens have been suggested to increase citrullination of selfas well as bacterial proteins which, in theory, can subsequently activate ACPAexpressing B cell by means of both the citrullinated antigens and pathogen-associated molecular patterns (PAMPs). These bacteria are *Porphyromonas gingivalis* and *Aggregatibacter actinomycetemcomitans*. *P. gingivalis* contains a bacterial citrullinating enzyme (termed *Prophyromonas* PAD or PPAD) and bacterial enolase. Anti-*P.gingivalis* antibodies are increased in RA patients and are detectable years before the onset of RA.⁹⁵ Moreover, a positive correlation between the serum levels of anti-PPAD IgG and ACPA IgG has been reported in Japanese subjects.⁹⁹ This correlation is, however, could not be shown for subjects in the US.¹⁰⁰

It has been hypothesized that humans develop antibodies against PPAD-mediated citrullinated bacterial proteins which eventually cross-react with human citrullinated self-proteins due to molecular mimicry.^{101,102} To test this hypothesis, several groups have assessed the cross-reactivity of ACPA from RA patients with citrullinated *P. gingivalis*-derived proteins and peptides. Using patient-derived anti-citrullinated alpha-enolase peptide antibodies, Lundberg *et al* showed that these autoantibodies bind a citrullinated bacterial enolase peptide which shares sequence similarity with human enolase.¹⁰³ At the protein level, however, Muñoz-Atienza *et al*, could not identify ACPA binding specificity towards auto-citrullinated *P. gingivalis* proteins.¹⁰⁴ In addition, PPAD has been shown to citrullinate human fibrinogen and enolase which are two binding targets of ACPA, in a manner different from human PAD.^{105,106} These findings suggest possible roles of *P.gingivalis* in the development of RA, particularly via PPAD-mediated bacterial and human protein citrullination, although the mechanistic basis for this hypothesis is still under debate.

Aggregatibacter actinomycetemcomitans (Aa), on the other hand, does not contain bacterial PAD. Instead, it produces leukotoxin A (LtxA), a pore-forming toxin, which increases calcium influx in neutrophils, triggers hypercitrullination of intracellular neutrophils proteins, and releases the hypercitrullinated cargo in a similar fashion as neutrophil extracellular trap (NET) formation.⁹⁶ These hypercitrullinated neutrophil proteins are thought to trigger immune responses towards citrullinated proteins in susceptible individuals. In a study from the US, exposure to the bacterial LtxA was found, indeed, associated with ACPA, suggesting a putative role of Aaas a trigger for the breach of tolerance towards citrullinated proteins in RA.⁹⁶ This association, however, was absent in a large cohort of Dutch RA patients.¹⁰⁷



To further complexify the possible mechanisms triggering ACPA immune responses, it was found that ACPA can also cross-react with antigens bearing other post-translational modifications (PTM) such as acetyl-lysine and homocitrulline.⁹⁴ This cross-reactivity is likely to be more relevant as it can directly induce a breach of tolerance. Apart from ACPA, other anti-modified proteins antibodies (AMPA) such as anti-carbamylated proteins antibodies (anti-CarP) and anti-acetylated proteins antibodies (AAPA) are also prevalent in RA.^{29,108} Due to this cross-reactivity, AMPA are thought to originate from a common B-cell response which differentiates into a cross-reactive AMPA immune response with varying avidity for different PTM antigens.²⁹ Importantly, it has been shown in a murine model that such diversification can occur *in-vivo*. Immunization with either acetylated or carbamylated murine proteins induced both AAPA and anti-CarP antibodies.²⁹ These findings have opened many possible originating antigens which can commence and activate the ACPA immune response, particularly because acetylation in bacterial proteins is commonplace.¹⁰⁹

The mucosal bacterial origin of ACPA-positive RA theory is further supported by the fact that some antibiotic therapies are also effective in treating ACPA-positive RA. Double-blind trials have shown that minocycline, a tetracycline antibiotic, is associated with improved clinical outcome in both early and established seropositive RA.¹¹⁰⁻¹¹³ Furthermore, the patients who benefit most from minocycline are those possessing HLA-SE which is a major susceptibility gene in ACPA-positive RA.¹¹⁴ Other anti-bacterial therapies such as levofloxacin and clarithromycin have also been associated with clinical improvements in RA.¹¹⁵ These drugs, however, have also anti-inflammatory properties which are independent of their antibacterial effects.¹¹⁵⁻¹¹⁸ It is therefore not clear whether the clinical response of these antibiotics occurs primarily due to the anti-bacterial or anti-inflammatory properties of these drugs. Nonetheless, these findings indicate putative roles of mucosal bacteria in the development of ACPA-positive RA which may be different in each population or individuals.

7.2.2 Genetic and epigenetic factors

Apart from environmental factors, genetic elements also play roles in the origin of ACPA-positive RA. It is estimated that the heritability of ACPA-positive RA is approximately 68%.¹¹⁹ Although the variation in the susceptibility genes may vary in different populations and occur in only a fraction of RA patients, they can provide insights in the development of ACPA-positive RA. Single nucleotide polymorphisms (SNPs) that are known to render people susceptible for developing

ACPA-positive RA include those at *HLA*, *PTPN22*, *PAD4*, *SLC8A3*, *CD28*, *CTLA-*4, *CD40*, *FCGR2*, *FCRL3*, *IL6R*, *TRAF1/C5* and *CCR6* genes.¹²⁰ Using this information, a chain of events can be made to decipher the mechanism by which ACPA-positive RA develops (Figure 1b and 1c.

Excessive and/or ectopic protein citrullination may induce the development and activation of ACPA-expressing B cells. SNPs in the genes involved in protein citrullination, including the citrullinating enzyme peptidylarginine deiminase 4 (*PADI4* and sodium/calcium exchanger membrane protein *SLC8A3*, associate with ACPA-positive RA.^{121,122} A SNP at *PADI4* also associates with progressive joint destruction in ACPA-positive RA in Japanese patients.¹²³ The excessive citrullinated peptides can then be presented by antigen presenting cells (APC, which include ACPA-expressing B cells and dendritic cells, to CD4⁺ T cells with the help of human leukocyte antigen (HLA class II.¹²⁴

HLA polymorphisms have been estimated to contribute 11-37% to RA heritability which can either increase or decrease the susceptibility of getting ACPA-positive RA.^{125,126} A group of HLA-DRB1 alleles that bears similar sequence motifs comprising of five amino acids in residues 70-74 of the DR chain, termed the shared epitope (SE, has a major influence on ACPA-positive RA susceptibility. ACPA-positive individuals with HLA-SE are 4-6 times more likely to develop RA than those without these alleles.¹³ Moreover, HLA-SE alleles are linked to the levels of ACPA, indicating a direct role of these genes in the production of ACPA.¹²⁷⁻¹²⁹ HLA-SE alleles also associate with ACPA-IgG VD glycosylation, a hallmark of mature ACPA response in RA.¹³⁰ Remarkably, there is no significant positive association between HLA-SE with ACPA positivity in individuals without RA.⁷⁷ These findings suggest that HLA-SE alleles have central roles in the activation and maturation of the ACPA positivity to ACPA-positive disease.

Activation of T cells via antigen presentation requires not only peptide-HLA molecule complexes but also co-stimulatory molecules including CD80 and CD86. These co-stimulatory signals are then received by two competing receptors, the stimulatory CD28 and the inhibitory CTLA-4 on T cells. Gain-of-function SNPs of *CD28* and loss-of-function SNPs of *CTLA-4* are known to render people susceptible to develop RA.¹³¹⁻¹³³

Once activated, CD4⁺ T cells produce CD40 ligand which activates ACPA-expressing B cells and induces BCR somatic hypermutation. Subsequently, epitope

spreading, affinity maturation and VD glycosylation of ACPA may occur. These events mark the maturation of the ACPA immune response. Accordingly, SNPs at CD40 genes as well as the genes related to CD40 signalling such as TRAF1and TNFAIP3, have been associated with the development of joint destruction in ACPA-positive RA.¹³⁴⁻¹³⁶ The activation of both T and B cells requires intracellular signalling regulations which calibrate the threshold of activation. RA susceptibility genes which are involved in the modulation of lymphocyte activation are PTPN22 and FCRL3.¹³⁷ Individuals who have a combination of a PTPN22 variant and ACPA positivity confer a much higher relative risk to develop RA than those who have either risk factor, suggesting pathogenic roles of hyperactive lymphocytes, especially ACPA-expressing B cells, in the development of ACPA-positive RA.¹³⁸ A genetic variance on PRDM1, a transcription factor essential for B cell differentiation into plasmablasts/cells (PB/C), also associates with seropositive RA, indicating putative pathological roles of autoreactive PB/C in the development of RA.¹³¹ Activated B cells are indeed found in the RA synovial tissue. One of the chemokine receptors which regulates migration towards inflamed organs is CCR6. SNP of this receptor as well as chemokine CCL21 are associated with ACPA-positive RA.^{136,139-141}

Once mature ACPA are produced at a certain level and bind to citrullinated antigens in the joints, they form immune complexes that activate effector cells through FcR binding as well as complement activation, resulting in joint inflammation. Accordingly, polymorphisms in the FcR and complement factor-expressing genes, *FCGR2A*, *FCGR3A* and *TRAF1-C5*, have been shown to associate with ACPApositive RA in Caucasian subjects.^{131,142-144}

Other than polymorphisms at the susceptibility genes, epigenetic factors, such as DNA methylation at *PCDHB14* is associated with ACPA positivity, while DNA methylations at *PCDHB5* and *EXOSC1* are associated with ACPA-positive RA.^{145,146} This supports the distinct mechanisms of the breach of tolerance towards citrullinated antigens and development of ACPA-positive RA also at epigenetic levels.

There are some overlaps between the genetic basis of ACPA-positive and -negative RA.^{13,120,147} This indicates that B cell-independent factors also contribute to the pathogenesis of RA while the ACPA response enhances the progression and the chronicity of the disease (Figure 1c). These overlaps include a genetic variant at *C5orf30* which encodes a 206-aa protein that regulates tissue damage response and activation of synovial fibroblasts and macrophages.^{13,148,149}

A *C5orf30* polymorphism increases the invasiveness and the production of gamma interferon-inducible protein-10 (IP-10) by synovial fibroblasts.¹⁵⁰ Synovial fibroblasts physiologically constitute the inner layer of synovium which is essential for the production of the components of the synovial fluid. These cells also produce optineurin to inhibit osteoclast differentiation in inflammatory conditions.¹⁵¹ In RA, however, these cells display an altered phenotype, grow uncontrollably and produce IL-6, RANKL as well as matrix metalloproteases which mediate joint inflammation and bone erosion.^{152,153} The altered phenotype may stem from the upregulation of endothelium-derived Notch3 signalling.¹⁵⁴ Moreover, synovial fibroblast subsets may have specific pathological functions in RA. Adoptive transfer of different subtypes of synovial fibroblasts in mouse model demonstrate specialized function of fibroblast subtypes in mediating inflammation or joint damage.¹⁵⁵ Targeting synovial fibroblasts is possible using anti-cadherin-11 mAb which leads to reduction of joint inflammation in mouse models of arthritis.¹⁵²

Furthermore, polymorphisms of genes encoding cytokine receptor and signal transducer, TRAF1/C5 and STAT4, are also associated with RA irrespective of the ACPA status.^{13,120} The roles of pro-inflammatory cytokines in the development of RA are further evidenced by the association of *IL6ST* and *TRAF1* gene polymorphisms with RA as well as the success of IL-6 and TNF-alpha inhibitors for the treatment of RA regardless of the ACPA status.^{120,139,156} These data suggest that dysregulation of synovial fibroblasts, macrophages and the cytokine network in the joints may be the determinants of arthritis development which are further aggravated and sustained by the ACPA immune response.

Altogether, these susceptibility genes highlight the plethora of immune dysregulations that play roles in the development of ACPA-positive RA. The mere presence of any of these susceptibility genes, however, does not cause RA as shown in large twin studies which prove low concordance rates for ACPA-positive RA in monozygous twin. ^{157,158} This observation suggests a complex interplay between susceptibility genes and environmental factors in the development of ACPA-positive RA.

7.2.3 The perpetrators in established ACPA-positive RA: lessons from clinical trials

The importance of specific immune cells and cytokines in the pathogenesis of ACPA-positive RA has been substantiated or disapproved by RA clinical trials testing targeted therapies in the form of biological and targeted synthetic disease-modifying anti-rheumatic drugs (bDMARDs and tsDMARDSs). Inhibitors of the

IL1 receptor (anakinra), TNF-alpha (infliximab), IL-6 (tocilizumab, sirukumab). the IL-6 receptor (sarilumab), GM-CSF (otilimab) and the GM-CSF receptor (mavrilimumab) are efficacious in treating RA regardless of ACPA status.^{156,161-164} Moreover, inhibitors of IAK-STAT signalling (tofacitinib, baricitinib, filgotinib). which relay cytokine signals. also show good clinical response in both types of RA.¹⁶⁵⁻¹⁶⁹ Therefore, these clinical benefits highlight the central roles of proinflammatory cytokines in the aetiopathogenesis of RA regardless of ACPA status.

Under physiological condition, these mediators of inflammation are produced locally at the site of tissue injury which activate both local and systemic inflammatory responses. Such responses aim to enhance the clearance of pathogens and injured tissue as well as to stimulate the healing process. To this end, extra resources from the body are required and therefore compensatory mechanisms take place; for instance, the suppression of red blood cell formation and demineralization of the bone.^{170,171} To regulate this process, pro-inflammatory cytokines are normally modulated by negative feedback signalling and the activation of antiinflammatory cytokines as well as other pro-resolving mediators. In RA, however, the disbalance of the cytokine network occurs chronically due to persistent tissue injury, hyperactive immune cells and dysfunctional resolution of inflammation.172

Pro-inflammatory cytokines such as IL-1, IL-6, TNF alpha, and GM-CSF have pleiotropic effects on immune cells, synovial fibroblasts, neuron excitation, osteoclast maturation, and activation of endothelial cells all of which play roles in RA pathogenesis. These cytokines are produced mainly by activated synovial fibroblasts, monocytes and innate lymphoid cells which in turn enhance the activation, differentiation and survival of the immune cells as well as osteoclasts that play direct roles in joint inflammation and damage.¹⁷³⁻¹⁷⁶ Indeed, the levels of these cytokines are elevated in the blood and/or synovial tissue of RA patients.¹⁷⁷⁻¹⁷⁹ As a result, they promote the up-regulation of body temperature, fatigue and joint pain which characterize the symptoms of RA.¹⁸⁰⁻¹⁸³

The association of genetic variants of HLA-DR, CD28, CTLA4, PTPN22 and CD40 signalling with ACPA-positive RA as well as the abundance of T cells in RA synovium strongly suggest the involvement of T cell activation in the aetiopathogenesis of the disease.^{184,185} CD4⁺ helper T cells develop from common lymphoid progenitors in the bone marrow and migrate to the cortex of the thymus to undergo positive and negative selection in which T cells expressing T cell receptors (TCR) that bind self-proteins are largely eliminated. This central T cell tolerance is essential in ensuring that competent 26 T cells are not autoreactive.

Accordingly, mutations of *AIRE*, a gene central in T cell tolerance, cause multiorgan autoimmunity.¹⁸⁶ Once emigrating from the thymus and getting activated by antigen-presenting cells, naïve T helper cells further develop into intracellular pathogens-counteracting TH1, extracellular parasites-counteracting TH2, B cell-stimulating follicular helper T cells (TFH) and peripheral helper T cells (TPH), IL-17-producing TH17, and immune regulatory T cells (Treg).¹⁸⁷

Clinical trials targeting T cells, their subsets and cytokines to treat RA, however, have shown less promising results. The phase 1 clinical trials of a pan-T cell depletion therapy using anti-CD3 monoclonal antibody (mAb) otelixizumab to treat RA have been terminated due to a "changed risk-benefit ratio for the patients". Moreover, trials of anti-CD4 mAbs for RA have shown mixed results, with the most positive ones only resulting in modest improvement in clinical parameters.¹⁸⁸⁻¹⁹⁴ By depleting CD4⁺ T cells, it is hypothesized that not only pro-inflammatory TH17 cells but also anti-inflammatory Treg cells are affected. Therefore, a few trials have attempted to specifically target TH17-associated cytokines and to enhance Treg cell functions. Monoclonal antibodies against TH17-associated cytokines, such as anti-IL-12/23 (ustekinumab), anti-IL-23 (guselkumab) and anti-IL17A (secukinumab) failed to induce clinical improvement in RA.^{195,196} Efforts to activate Treg by targeting unique CD4 epitopes using tregalizumab have also failed to improve clinical outcome of RA patients.^{197,198}

The failure of T-cell targeting trials may stem from the fact that T cells in established RA are hypo-responsive to stimulation.¹⁹⁹ This anergic phenotype may be linked to T cell exhaustion due to persistent antigen and inflammatory signals.²⁰⁰⁻²⁰³ T cells seem to have passive roles in perpetuating joint inflammation in established RA and therefore targeting T cells at this stage of RA does not render significant clinical response. T cells are more likely to have major roles in the maturation of the ACPA response, the second hit of RA development. This hypothesis is supported by the fact that TH17-produced IL-21 and IL-22 can change the Fc-glycosylation profile of IgG.²⁰⁴ Glycosylation changes hallmark the maturation of ACPA and are strongly associated with the progression toward RA. These findings indicate that T cells may play more pivotal roles during the maturation of the ACPA response before the onset of disease rather than in the established and/or chronic phase of RA.

In contrast to T cell depletion, B cell depletion therapy using anti-CD20 mAb rituximab (RTX) is significantly more efficacious than methotrexate in reducing RA symptoms and progression.^{21,205,206} Besides for RA, RTX is authorised in Europe for

the treatment of B cell malignancies such as follicular lymphoma, diffuse large B cell non-Hodgkin's lymphoma and chronic lymphocytic leukaemia; and B cell-related autoimmune diseases such as granulomatosis with polyangiitis, microscopic polyangiitis and pemphigus vulgaris. The depletion of B cell subpopulations in peripheral blood using RTX is significant. In solid tissues, however, the depletion is incomplete.²⁰⁷

B cell development starts in the bone marrow where common lymphoid progenitors sequentially differentiate into pro-B cells, pre-B cells and immature B cells. These B cell precursors are selected based on productive rearrangement of immunoglobulin heavy (IgH) and light (IgL) chains as well as the expression of CD79a/b which together form the functional BCR complex. The expression of the transcription factor PAX5, which associates with the expression of CD19, is essential for B cell identity. Furthermore, the expression of CD20 starts from the pre-B cell stage.²⁰⁷ Transitional B cells egress from the bone marrow into the peripheral blood and mature further to become antigen-inexperienced IgD⁺ IgM⁺ naïve B cells.²⁰⁸ Upon binding to antigen, naïve B cells migrate into secondary lymphoid organs where B cells undergo class-switching recombination and somatic hypermutation (SHM) in their BCR genes with the help of TFH. B cells with high affinity to the antigen receive T cell stimulation and undergo clonal expansion as well as further development into memory B cells and antibody-secreting cells (ASCs). ASCs cease to express CD20 as well as PAX5 and some migrate back to the bone marrow or inflamed tissues as long-lived plasma cells. Therefore, RTX only selectively depletes pre-B cells up until the memory B cell subsets, but not ASC.

There are several biomarkers which can predict good and poor response to RTX. The serological biomarkers that predict good clinical response are ACPA, RF, IL-33 and increased erythrocyte sedimentation rate (ESR).²⁰⁹⁻²¹¹ A good response to RTX in RA is not only predicted by ACPA positivity but also by high levels of ACPA.^{210,212,213} Another good predictive biomarker is IL-33 which is a cytokine released from necrotic cells to alert the immune system for tissue damage or stress. In RA, IL-33 circulates at higher levels compared to healthy individuals and is decreased after anti-TNF treatment.^{214,215} The fact that IL-33 synergistically predicts positive RTX responses with the autoantibodies suggests that tissue damage may support the activation and survival of ACPA-expressing B cells.²¹¹

Poor predictors of RTX responses in RA include elevated CD20-negative ASC markers and an increased type I interferon (IFN) signature.²¹⁶⁻²¹⁹ In active RA, general type I IFN signature does not correlate with the presence and titres of ACPA

and RF.²²⁰ Moreover, people with a high type I IFN signature are more likely to develop RA, independent of the autoantibody status.²²¹ On the other hand, specific type I IFN signature genes which regulate helper T cells and B cell proliferation correlate significantly with the levels of ACPA and anti-CarP antibodies.²²² These findings suggest that type I IFN activation plays diverse roles in the development of RA, one of which is through the activation of autoreactive B cells.

All in all, clinical trials provide clear evidence for the involvement of specific cytokines and immune cells in established RA. Pro-inflammatory cytokines are central mediators of RA symptoms regardless of serological status. Moreover, CD20-positive B cells, particularly ACPA-expressing B cells, are essential for the progression of ACPA-positive RA. T cells, on the other hand, seem to be dispensable in sustaining and progressing joint inflammation in established RA.

8 ACPA-expressing B cells

8.1 The origin of ACPA-expressing B cells

Serological, genetic and clinical studies discussed above demonstrate the putative central roles of ACPA-expressing B cells in the aetiopathogenesis of ACPA-positive RA. The two hit theory of the ACPA response in RA development suggests that prior to disease onset ACPA-expressing B cells develop and mature at different time points which depend on distinctive factors (Figure 1a and 1b). In the bone marrow, B cells undergo random V(D)J recombination of BCR genes to enable the binding to infinitely diverse pathogenic antigens. The by-product of this process is the occurrence of autoreactive B cells. In fact, it has been suggested that 75% of pre-B cell clones are autoreactive, as defined by the ability to bind HEp-2 cell extract.²²³ The percentage of autoreactive BCR is decreasing during B cell development from an estimated 43% in the immature B cell stage, 40% in the transitional B cell stage and around 20% in the mature naïve B cell stage.²²³ Self-reactive B cells are, therefore, largely removed in the bone marrow at the immature B cell stage and in the blood during the transition to become mature naïve B cells.

Immunological tolerance is essential to prevent the immune system from attacking its own body. Besides autoantigen-specific B cell removal, immunological tolerance is safeguarded by receptor editing, autoantigen-specific anergy and ignorance induction.²²⁴ Switching the BCR light chain from Ig-kappa to Ig-lambda, for example, is typically effective in modifying BCR reactivity away from autoreactivity.²²⁵ Under physiologic conditions, surviving autoreactive naïve B



cells that are not deleted during B cell development adopt an anergic phenotype. such as lower IgM expression, and are prevented to become ASC by tolerance checkpoints.²²⁶ These cells are preserved in the periphery as they increase the diversity of the BCR pool which may benefit the protective immune response. infection. these autoreactive B cells mav undergo somatic Upon hypermutation (SHM) and selection processes in the germinal centre which results in decreased affinity towards self-antigens and increased binding to infectious antigens.^{226,227} In fact, broadly neutralizing antibodies against HIV-1 retain some degree of poly- and autoreactivity.²²⁸ It is hypothesized that these viral responses may arise from poly/autoreactive B cell clones which progressively lose their autoreactivity and become protective.^{229,230}

TFH cell-dependent BCR selection procedures and multiple rounds of BCR gene editing are necessary for dampening autoimmunity. BCR with high affinity to an antigen are able to bind and capture the antigen and present the processed antigen to antigen-specific T helper cells. In turn, TFH cells provide survival and activation signals to B cells. Antigen-stimulated germinal centre B cells which do not receive stimulation from TFH will undergo apoptosis.²³¹⁻²³³ On the other hand, strong T cell stimulation with weak BCR signalling may also induce FAS-dependent B cell death.²³⁴ Therefore, the requirement of balanced BCR and T cell-derived signals in the survival and activation of germinal centre B cells provides immunological checkpoints to prevent the activation of autoreactive B cells in the periphery.

This multi-step process, however, is not error-free and can be hijacked. For instance, up-regulation of anti-apoptotic signals can inhibit the elimination of autoreactive B cell clones in the germinal centre.²³² The random nature of BCR SHM, just like V(D)J recombination, can give rise to autoreactive BCR clones. In fact, autoreactive IgG⁺ MBC can be detected in the blood of healthy donors. These cells express BCR which become autoreactive only after undergoing SHM in the germinal centre.²³⁵

The germinal centre reaction normally occurs in secondary lymphoid organs (SLO) such as lymph nodes and spleen. In inflamed tissues, germinal centre reactions may also develop ectopically in inducible tertiary lymphoid organs (TLO).²³⁶ Physiologically, TLOs are able to develop local and quick humoral immune responses to foreign antigens. The immunological tolerance check-points at TLOs, however, may be less strict compared to SLOs which favours the development of autoreactive B cells.^{237,238} Persistent TLOs can be found in mice exposed to chronic cigarette smoke exposure, a risk factor of ACPA development.²³⁹ Therefore, dysregulation of immunological tolerance, which associates with cigarette smoking,

may support the development of low-affine ACPA-expressing B cells. Such cells mark the first hit in the development of the ACPA response.

8.2 The activation of ACPA-expressing B cells

The intricate genetic and environmental factors discussed previously may lead to the activation of ACPA-expressing B cells. The hallmark of this process is an enhanced T cell-dependent germinal centre activity including SHM and classswitch recombination. The BCR of ACPA-expressing MBC in RA patients is mutated extensively, contains predominantly immunoglobulin lambda light chains, and is mostly class-switched. These characteristics indicate several rounds of IgH and *IeL* gene editing in the germinal centre which lead to the expansion of ACPA repertoire.^{70,240} Moreover. approximately 90% of ACPA-IgG expressing MBCs in RA have BCR variable domain (VD) glycosylation that is introduced during SHM.⁷⁰ In mouse models of autoimmunity, VD glycans on the BCR occupy the majority of antigen binding sites and decrease the avidity of residual unglycosylated sites.²²⁷ On ACPA, however, VD glycosylation does not generally decrease the avidity towards citrullinated antigens but may affect the accessibility of antigen binding sites through steric hindrance. The high proportion of ACPA-expressing B cells with VD glycosylation indicates that this modification may provide survival advantage during maturation of ACPA response.

Remarkably, the VD glycans on ACPA contain sialic acids which can have immunomodulatory properties through binding to siglecs.^{241,242} Theoretically, the highly sialylated BCR of ACPA-expressing B cells may dampen negative selection mechanisms allowing for their survival in the periphery. Moreover, the bulky VD glycan of ACPA may interact with the BCR and other proteins which increase tonic BCR signalling.²⁴³ It has been shown that BCR signalling dysregulation can disturb B cell tolerance mechanisms. For instance, genetic variance of *PTPN22* and *PIK3CD* disturb BCR signalling and, consequently, reduce the removal of self-reactive B cells.^{244,245} Therefore it is possible that highly sialylated VD glycans provide survival advantages to ACPA-expressing B cells through dysregulation of BCR signalling and/or inhibition of immunological tolerance check-points.

The activation of MBC is also regulated by negative feedback from antibodies. Binding of antigen-antibody immune complex to both BCR and inhibitory Fc gamma receptor 2b (FcGR2b or CD32b induces apoptosis of antigen-specific MBC which limits antigen-specific humoral immune response and leaves room for other plasma cells in bone marrow.^{246,247} Down-regulation of CD32b

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expression and function on MBC impairs this negative feedback mechanism and associates with increased autoreactive IgG⁺ MBC.²⁴⁸ Therefore it is possible that ACPA-expressing MBCs modulate their CD32b signalling to remain active despite high levels of circulating ACPA and the presence of circulating antigens.

8.3 The migration and pathogenic roles of ACPA-expressing B cells

Activated B cells have effector functions locally by migrating towards inflamed or injured tissue, attracting and activating other immune cells *in situ* as well as systemically by production of antibody and cytokines. Peripheral MBCs express chemokine receptors which allow them to migrate to inflammatory and secondary lymphoid tissues following a gradient of chemokines.^{249,250} The serum levels of the chemokine CCL19 are increased in ACPA-positive RA patients, positively correlate with B cell activation markers and predict RTX response at 24 week.²⁵¹ CCL19 is primarily produced by stimulated RA macrophages and synovial fibroblasts and attracts CCR7-expressing cells to the inflamed synovial tissue.²⁵² Through the putative expression of chemokine receptors, activated ACPA-expressing MBC may migrate to inflamed and/or injured synovial tissues.

In the inflamed synovial tissue, ACPA-expressing MBC may get activated by citrullinated proteins and the inflammatory milieu to become ACPA-producing plasmablasts/cells (PB/C).^{253,254} The source of citrullinated proteins includes neutrophil extracellular traps which are present in RA synovium.²⁵⁵ ACPA-expressing MBC may also interact with specialized PD-1 high, CXCR5 negative peripheral helper T cells in RA synovium to differentiate into ACPA-producing PB/C in IL-21-dependent manner.¹⁸⁷ ACPA-producing PB/Cs are indeed present and active in RA synovial fluid.^{253,256,257} Furthermore, their survival is supported by the presence of other mononuclear cells in RA synovial fluid.²⁵⁶

In the joint, activated B cells can propagate local inflammation and induce bone erosion through the expression of co-stimulatory molecules and the production of cytokines and antibodies. The importance of the co-stimulatory molecules CD80/86 in ACPA-positive RA pathogenesis is demonstrated by the fact that ACPA positivity correlates with better treatment response to the CD80/86 blocking agent abatacept.²² CD80/86 is expressed by various antigen presenting cells and co-stimulates immune responses through an interaction with CD28 which is mainly expressed by T cells. Given the lack of clinical evidence for a role of T cell role in sustaining arthritis, the expression of CD80/86 by ACPA-expressing B cells can, theoretically, stimulate innate immune responses. Clinical study of the superagonist

anti-CD28 mAb TGN1412 led to an abrupt fatal cytokine storm which may not stem from T cell activation.²⁵⁸ Human immune cells that express functional CD28 and secrete pro-inflammatory cytokines immediately include neutrophils and eosinophils. In fact, neutrophils are the most abundant immune cells in the RA synovial fluid.²⁵⁹ Crosslinking of CD28 on these cells results in IFN gamma, IL-13 and IL-2 production.^{260,261} Moreover, Immune complex-activated neutrophils produce osteoclast-activating S100A8 and MMP-9 which play roles in bone erosion.^{262,263} These cells also release their nuclear components to form neutrophils extracellular traps (NETs) which contain citrullinated histones.²⁵⁵ Hypothetically, ACPA-expressing MBC can be activated by these citrullinated histones and differentiate into ACPA-producing PB/Cs which in turn activate neutrophils in CD86, IL-8 and ACPA-IC-dependent manners. This hypothesis argues for the bidirectional interactions between ACPA-expressing B cells and neutrophils that propagate and perpetuate inflammation as well as bone erosion.

Furthermore, BCR-stimulated B cells can produce pro-inflammatory cytokines and chemokines which are involved in bone erosion.²⁶⁴⁻²⁶⁷ Synovial B cells from RA patients spontaneously produce CCL3 and TNF-alpha which suppress osteoblast differentiation.²⁶⁵ These cells also produce RANKL which induces boneresorbing osteoclast development.^{266,267} Taken together, ACPA-expressing B cells may migrate to the inflamed synovium, get activated and remain in the synovial tissue to contribute to the progression of RA by not only producing ACPA but also co-stimulatory molecules, cytokines and chemokines which perpetuate joint inflammation and enhance bone erosion (Figure 1d).

8.4 The nature of ACPA-expressing B cells: lessons from rituximab and abatacept

The nature of ACPA-expressing B cells in RA can be deduced by studying the kinetics of clinical responses, ACPA levels and fine specificity, as well as B cell counts after treatment using RTX and abatacept. ACPA are strong positive predictors of the response to both medicines.²⁶⁸ Two infusions of 1000 mg rituximab on days 1 and 15 deplete circulating B cells at week four post first treatment. At this timepoint, synovial B cells are significantly decreased though not completely absent. Moreover, there is also a significant decrease of both synovial T cells and macrophages. The 28-joint Disease Activity Score (DAS28) as well as ACPA levels are not yet significantly reduced at this timepoint, while at weeks 16 and 24 post first treatment, the DAS28 is significantly reduced. The clinical improvement can be predicted by the reduction of synovial PB/C. At around 24 weeks, peripheral B

cells start to return. ACPA levels, however, only start to decrease significantly at 36 weeks.²⁶⁹ These observations provide evidence for the role of B cells in triggering local immune infiltration and the incapacity of ACPA in inducing arthritis. This notion is also supported by the fact that ACPA levels are frequently unaltered during disease remission.²⁷⁰

Moreover, the delay of the clinical response after CD20 depletion suggests that CD20⁺ B cells may mediate joints inflammation indirectly through activation of other immune cells and differentiation into plasma cells.²⁷¹ In fact, unaltered plasma cell counts in the joints associates with poorer response to RTX.²⁷¹

Relapse occurs after discontinuation of RTX treatment and is preceded by the reoccurrence of IgM⁺ naïve B cells from the bone marrow. Remarkably, the circulating ACPA at relapse consist of new and old antigen binding specificities. These findings suggest that the mechanism underlying ACPA responses is self-sustaining in RA and that interactions between new and residual ACPA-expressing B cells may be essential in resurgence of symptoms.^{272,273}

In contrast to RTX therapy, a decline in ACPA levels at 3 months predicts sustained improvement of RA symptoms at 12 months after treatment with abatacept, a chimeric CTLA4-Fc construct.²⁷⁴ Given that ACPA-producing plasma cells can live at least for several months, the early reduction of ACPA indicates that abatacept may not only inhibit the development of ACPA-expressing MBC into plasma cells but also directly inhibit ACPA-producing PB/C. CD80/86 proteins contain binding domains for signalling proteins in their cytoplasmic tails, suggesting the ability to transduce signals upon activation. In dendritic cells, cross-linking of CD80/86 by CD28 results in IL-6 production.²⁷⁵ Thus, the engagement of abatacept with CD80/86 on ACPA-producing PB/C may reduce their pro-inflammatory capacity which lead to a decrease in ACPA levels and, possibly, clinical improvement.

ACPA B cell responses seem to differ from responses against vaccine antigens or virus infection. Years after sustained MBC depletion, rhesus macaques still produce anti-virus antibodies which are attributed to long-lived plasma cells.²⁷⁶ MBCs, on the other hand, have the capacity to mutate their BCR to neutralize virus mutants which escape the binding to pre-existing antibodies.²⁷⁷ This division of tasks safeguards the body not only from the same pathogen but also from their evolving mutants. Accordingly, a two-year period of CD20⁺ B cell depletion in RA patients does not change the levels of protective antibodies against measles, mumps and

rubella.²⁷⁸ The levels of ACPA, however, decrease significantly in this period; even though in general no seroconversion was observed. Moreover, the positive correlation between circulating IgG levels and antigen-specific MBC is found in ACPA but not in anti-tetanus responses.²⁷⁹ These findings strongly indicate that ACPA-producing PB/C may frequently be regenerated by circulating ACPA-expressing MBC.

9 Strategies to target ACPA-expressing B cells for RA therapy

The pathogenic potential of ACPA-expressing MBC and ACPA-producing PB/C make them promising targets to treat ACPA-positive RA in a targeted manner. Depletion of CD20⁺ B cells increases the probability for developing *de novo* infections due to the reduction of protective MBCs. Specific targeting of ACPAexpressing B cells and other autoreactive B cells, on the other hand, provide a venue of next generation curative as well as preventative medicines for B cell-mediated autoimmune diseases in the era of targeted therapy. The aim of such therapies is the restoration of immunological tolerance by either inducing autoantigen-specific cellular anergy or by depletion of autoantigen-specific B cells. To this end, several strategies can be outlined: 1) blocking prominent signalling pathways on autoreactive B cells, 2) inducing specific tolerance by autoantigen-based immunotherapy, 3) targeting autoreactive B cells with autoantigen-drug conjugates, 4) employing a sequential pro-drug strategy to avoid circulating autoantibodies, 5) inhibiting autoreactive response by anti-idiotype immunotherapy, and 6) using cell-based therapies. The holy grail of immune tolerizing therapies is to reset tolerance towards self-antigens and to induce life-long protection against B cell-mediated autoimmune diseases. Passive tolerizing therapies such as the depletion of autoreactive B cells, however, may not result in sustained tolerance. Active tolerizing therapies, on the other hand, may provide sustained tolerance as a preventive and curative measure in autoimmune disease.

9.1 Signaling inhibitors

Autoreactive B cells breach immunological tolerance through dysregulation of signalling pathways which promote their survival and activation. This is well demonstrated by the association of a genetic variant of the Protein Tyrosine Phosphatase Non-Receptor Type 22 (*PTPN22*) gene with many autoimmune diseases, including RA, SLE, Grave's disease, anti-neutrophil cytoplasmic antibody-associated vasculitis, generalised vitiligo, myasthenia gravis and type 1 diabetes.²⁸⁰ The *PTPN22* R620W variant increases its degradation and enhances



BCR, BAFF receptor and CD40 signalling which leads to increased positive selection of autoreactive B cells.²⁸¹⁻²⁸⁴ Therefore, autoreactive B cells may be more sensitive to the blockage of these hyperactive signalling pathways than protective B cells. B cell signalling targets which may have clinical relevance in RA include Bruton's tyrosine kinase (BTK) and phosphatidyl-inositol-3 kinase (PI3K) (Figure 2a).

A. Autoantigen-non-specific B cell inhibition



B. Autoantigen-non-specific tolerizing therapy



Figure 2. Autoantigen-non-specific B cell inhibition and tolerizing therapy. (A) Inhibitors of signalling pathways which play roles in the activation and survival of activated autoreactive B cells such as Bruton's Tyrosine Kinase (BTK) and Phosphatidylinositol 3-Kinase (PI3K) are promising therapy for autoimmune diseases. (B) Adoptive transfer of activated regulatory t cells (Treg) or mesenchymal stem cells can induce tolerance in autoantigen-non-specific manner.

BTK plays a role in conveying BCR and Fc-gamma-R III signalling.²⁸⁵ Loss-of-function mutations at the kinase domain of BTK result in X-linked agammaglobulinemia, which is characterized by defective B cell development.²⁸⁶ In ACPA-positive RA patients, B cells upregulate the expression and function of BTK.^{287,288} In B cells, BTK inhibition suppresses not only BCR signalling but also signals from CD40 ligand (CD154, IL-21 and BAFF.²⁸⁸ Inhibiting BTK in mouse models of arthritis reduces B cell proliferation and autoantibody levels. As activated autoreactive B cells are proliferative, BTK inhibitors may have specific targeting effects on these cells. Moreover, BTK inhibitors also reduce Fc-gamma-R III-induced production of pro-inflammatory cytokines by macrophages.²⁸⁵ Therefore, in RA, BTK inhibition may selectively inhibit hyperactive autoreactive B cells while

reducing the pro-inflammatory capacity of ACPA-IC-activated macrophages.

B cell survival, differentiation and activation require metabolic reprogramming to meet the changing energy and building block requirements. The PI3K signalling pathway is crucial for this function.²⁸⁹ PI3K signalling is also essential for the expression of PAX5 and CD19 which control B cell commitment during the development from common lymphoid progenitors.²⁹⁰ Moreover, PI3K signalling has an indispensable role in BCR signalling.²⁹¹ Constitutively active PI3K-P110alpha expressed by autoreactive B cells abrogates negative selection in the bone marrow.²⁹² Furthermore, active PI3K-p110delta induces the production of autoantibodies through the suppression of pre-germinal centre peripheral tolerance.²⁴⁵ Specific inhibition of PI3k-p110gamma suppresses joint inflammation and damage in RA mouse models, reduces glomerulonephritis in SLE mouse models, and reverses autoimmune diabetes in mouse models of type 1 diabetes.²⁹³⁻²⁹⁵ Moreover, broad PI3K signalling inhibitors such as artesunate and metformin are able to suppress proliferation of chondrocytes and RA synovial fibroblasts in RA models.^{296,297} Therefore, targeting PI3K may benefit patients with ACPA-positive RA by enhancing autoreactive B cell negative selection and inhibiting synovial fibroblasts. Future studies that delineate which signalling pathways support the survival and activation of ACPA-expressing B cells will open more options on how to target these cells specifically using respective pathway inhibitors.

9.2 Autoantigen-based immunotherapy

Antigen immunotherapy has been used for more than a hundred years to induce allergen-specific desensitization in patients with allergy.²⁹⁸ By frequent administration of the allergen in increasing amounts for a long period of time, allergen-specific immune responses can be reduced. Despite differences in the immunopathological basis, autoimmune diseases may also benefit from autoantigen-based therapy.

A four double-stranded oligo-DNA abetimus is an immunomodulating agent which aims to induce tolerance for double stranded DNA and to reduce nephritic flares in SLE.²⁹⁹ Weekly infusion of abetimus was safe and reduced the levels of circulating anti-dsDNA-antibodies in a dose-dependent manner. In phase 3 clinical trials, patients with high-affinity anti-dsDNA antibodies experienced less renal flares; furthermore, abetimus reduced the need to use high-dose corticosteroids and cyclophosphamide.^{300,301} These findings show that antigen-based immunotherapy can serve as a decoy for autoantibody binding leading to reduced tissue-specific

effector functions (Figure 3a). For this purpose, autoantibody affinity to the therapeutic antigen may be essential for improving clinical parameters.



🔘 = autoantigen-specific T cell receptor gene-containing plasmid 🛛 🔘 = FOXP3-containing plasmid

Figure 3. Autoantigen-specific tolerizing therapy. (A) autoantigen-based immunotherapy can be used as decoy to prevent autoantibodies from binding to their targets and to induce antigen-specific immunological tolerance through binding with tolerogenic dendritic cells (DC) and activation of antigen-specific Treg. (B) Synthetic tolerogenic DCs can be made by conjugating autoantigen-MHC complex on nanoparticle. (C) Adoptive transfer of modified DC treated with autoantigen can induce autoantigen-specific immunological tolerance. (D) CD4+CD25bright T cells can be modified to express autoantigen-specific Tregs.

Another autoantigen-based immunotherapy, ATX-MS-1467, is a peptide mixture of four soluble epitopes from myelin basic protein for the treatment of multiple sclerosis (MS). In a mouse model of MS, ATX-MS-1467 induced the activation of antigen-specific Treg. In phase 2a trials, ATX-MS-1467 was safe and able to reduce MS-related brain lesions.³⁰² Similarly, a phase one trial showed that a peptide mixture of two soluble epitopes from thyroid stimulating hormone receptor (TSHR), ATX-GD-59, was able to suppress T cell and antibody response towards TSHR in patients with Grave's disease.³⁰³ These autoantigen-derived peptides

bind to the HLA on tolerogenic DC, supress the pro-inflammatory capacity of antigen-specific T cells, and induce the development of Treg (Figure 3a).³⁰⁴

The route of delivery of autoantigen-based immunotherapy may be crucial for the development of tolerance. Notably, oral or intra nasal immunization can induce mucosal tolerance which mimics tolerization against non-pathogenic foreign antigens such as food and airborne proteins such as pollens.^{305,306} Moreover, a combination of autoantigens with immunomodulating drugs can enhance antigen-specific tolerance. Intranasal immunization with proinsulin peptides, in combination with an anti-CD3 epsilon-specific antibody, can induce insulin-specific Tregs and reverse recent-onset type 1 diabetes in mouse models.³⁰⁷ This combinatorial therapy was more efficacious than monotherapy using the proinsulin peptide or anti-CD3 alone. Anti-CD3 treatment has been shown to expand T cells with regulatory capacities. Besides anti-CD3 antibodies, histone deacetylases inhibitors such as trichostatin A. microbial metabolites such as butyrate and proprionate, all trans-retinoic acid, low dose of IL-2 and rapamycin are known to induce Treg differentiation and enhance their immune regulatory functions.³⁰⁸⁻³¹⁶ Therefore, a mucosal route of autoantigen delivery as well as combinatorial therapy with Treg-enhancing drugs may increase the success of autoantigen-based immunotherapy.

Hitherto, robust data suggest that ACPA pathogenicity may stem from their ability to form immune complexes (IC). Due to the pro-inflammatory glycosylation profile of the Fc tail of "mature" ACPA, it is conceivable that ACPA-IC can induce FcR-dependent inflammation.⁷⁴ Therefore, administration of citrullinated antigens as a putative immunotherapy for RA must be done cautiously. Co-administration of FcR inhibitors may help preventing ACPA-IC-induced inflammation.

Another approach of antigen-based tolerizing immunotherapy mimics tolerogenic dendritic cells. Nanoparticles (NP) coated with autoantigen-derived peptides which are linked to major histocompatibility complexes (MHC or human leukocyte antigen, HLA, in human), resulting in pMHC-NP, induce the expansion of antigen-specific Tregs, suppress natural autoantigen presentation and prevent type 1 diabetes in mouse model (Figure 3b).³¹⁷ In humanized mouse models, systemic delivery of pMHC-NP induces autoantigen-specific regulatory CD4⁺ T cell type 1 (TR1)-like cells and drives the differentiation of regulatory B cells without affecting protective immunity.³¹⁸ This approach prevents the formation of IC that may worsen the disease and directly induces antigen-specific tolerance. Its efficacy in humans remains to be tested.

Chapter I

HLA molecules have complex associations with the development of ACPA-positive RA. While HLA shared epitope alleles increase the risk for RA development. other HLA-DR variants containing the amino acid sequence of "DERAA" are associated with protection from RA.³¹⁹ HLA-DERAA reduces the risk for RA development possibly by inducing central negative selection of DERAA-binding T cells, which bind not only HLA-DERAA-derived peptides on dendritic cells but also DERAA-containing foreign and autoantigens such as vinculin. Accordingly, HLA-DERAA-negative women who are pregnant with a HLA-DERAA-positive foetus have increased risk for developing RA.³²⁰ This may be due to foetal HLA-DERAA-containing cell micro-chimerism in the mother's body which activates DERAA-binding T cells and subsequently stimulates ACPA-expressing B cells. The selection of HLA molecules as well as citrullinated peptides to form pHLA-NP for RA therapy can therefore be challenging. Citrullinated-vinculin-derived DERAA-containing peptides with HLA-SE nanoparticles may be an option to induce tolerance towards citrullinated autoantigens in ACPA-positive healthy individuals as a preventative vaccine against ACPA-positive RA.

9.3 Autoantigen-drug conjugate

Along with being able to decoy autoantibodies from their anatomical targets and to induce tolerance, autoantigens can also serve to deliver drugs specifically to autoreactive B cells. The conjugated autoantigen can deliver drugs both on the surface and the inside of autoantigen-specific B cells. BCR-mediated internalization of autoantigen-drug conjugate can promote delivery of the drug intracellularly. Depending on the mode of action of the drugs, a suitable linker between the autoantigen and drug can be added. While cleavable linkers can benefit the release of intracellular drug upon internalization by antigen-specific B cells, non-cleavable linkers may sustain the inhibitory capacity of drugs targeting surface protein.

The risk of this approach is that autoantigen-drug conjugates bind to circulating autoantibodies which may induce toxicity against FcR-expressing cells. To avoid adverse toxicity, drugs which target B cell receptor signalling pathways such as BTK inhibitors may be favourable choices. Citrullinated peptide-BTK inhibitor conjugates may not only inhibit ACPA-expressing B cell activation but also macrophages which have active BTK signalling and are able to bind to the drug-immune complex.³²¹ For that reason, citrullinated peptide-BTK inhibitor conjugates may be able to target two pathogenic cells in RA, namely ACPA-expressing B cells and activated macrophages.

Besides BTK, B cell-specific inhibitory receptors may also be targeted using the autoantigen-drug conjugate strategy described. Some inhibitory receptors, such as CD22 and CD32b, are indispensable for the induction of peripheral B cell tolerance. Knocking-out of the genes expressing these receptors leads to autoimmunity in mice.^{322,323} CD22 binds to alpha2,6-linked sialic acids in cis to regulate B cell receptor-induced proliferation.³²⁴ On the other hand, CD32b binds with low affinity to the Fc tail of antibodies in the form of immune complexes to inhibit antigen-specific B cell activation.³²⁵

Inhibitory receptors generally require co-localization with the BCR to enact their inhibitory capacity. Co-ligation of the BCR with CD22 or CD32b ligands inhibits the development arthritis in mouse models.^{326,327} Therefore, co-ligation of ACPA BCRs with an inhibitory receptor through citrullinated antigen-inhibitory ligand conjugates may be effective in inhibiting ACPA-expressing B cells (Figure 4a). It is hypothesized that ACPA-expressing B cells may survive from immunological tolerance checkpoints by downregulating their inhibitory receptors. Therefore, the usefulness of this strategy depends on the expression and function of the inhibitory receptors on these cells.

9.4 Sequential pro-drug strategy: targeting B cells and plasma cells

The autoantigen-drug conjugate has two possible shortcomings. First, circulating autoantibodies may have neutralizing effects on the autoantigen-drug conjugates which limits the availability of the conjugates for autoreactive B cells. Second, there are limited options of drugs that can be used to avoid toxicity. To circumvent these issues, a sequential pro-drug strategy can be an option. First, a cytotoxic drug is conjugated with an inactive form of an autoantigen using a cleavable linker. Subsequently, an activating drug which binds to the surface of B cells is given to activate the autoantigen-drug conjugate in the vicinity of BCR. Only autoreactive BCRs will bind to the active form of the autoantigen-drug conjugate which is then internalized through antigen-induced BCR endocytosis. Upon linker cleavage in the endolysosome, the cytotoxic drug is released inside the cell (Figure 4b). We have shown that this "Trojan horse" strategy can be feasible in targeting ACPA-expressing B cells *in vitro*.³²⁸

Some autoreactive B cells exert their pathogenicity through the production of pathogenic autoantibodies that activate a receptor aberrantly or disturb essential molecular interactions. These autoantibodies are produced by plasma cells which cease to express BCR on their surface. To target these cells in an antigen-specific



Figure 4. Autoantigen-specific B cell inhibition or depletion. (A) Autoantigen-drug conjugate can deliver immunomodulating drugs only on autoantigen-specific B cells. (B) Sequential prodrug strategy can circumvent neutralization by circulating autoantibodies and deliver cytotoxic drug to autoantigen-specific memory B cells. Moreover, this strategy can also specifically target autoantibodies contain idiotopes which can induce immune response. Idiotype of ACPA, alongside with adjuvants, can be used as vaccine against ACPA-positive RA through development of anti-ACPA idiotypic immune response. (D) Adoptive therapy using geneticall modified CD8⁺ T cells which express chimeric autoantigen or autoantibody idiotype receptor are effective in depleting pathogenic autoreactive B cells.

manner, an "affinity matrix" technology was develop by conjugating an antigen to an antibody fragment against CD138, a marker of plasma cells.³²⁹ The specific antibodies will then bind to the antigen and induce complement-dependent cytotoxicity.

The possible caveat of this strategy is that autoantibodies may bind to the autoantigen on the affinity matrix before binding to plasma cells. This can lead to depletion of plasma cells regardless of antigen specificity. To circumvent this potential issue, a sequential pro-drug approach could be employed. In this approach, an inactive form of autoantigen is conjugated to an antibody fragment against CD138. A few hours after the injection of the pro-drug, an activating drug can be administered to activate the autoantigen. The period between the injection of the pro-drug and the activating drug would need to reflect the time needed by the pro-drug to bind to plasma cells. The activated autoantigen then bind autoantibodies and induce complement-dependent cytotoxicity (Figure 4b). This approach could increase the prospect of success in targeting antigen-specific plasma cells.

9.5 Anti-idiotype immunotherapy

During B and T cell development, BCR and TCR undergo extensive recombination and hypermutation leading to highly diverse variable domains (VD). The novel VD BCR and TCR sequences and structures may not be tolerated by the immune system and therefore can serve as immune target determinants or epitopes. The parts of VD BCR and TCR which serve as epitopes are referred to as idiotype. In this concept, an antibody idiotype represents a new antigen to which immune responses can develop. Anti-idiotypic immune responses are hypothesized to regulate antigenspecific immune responses through the formation of idiotype-reactive cytotoxic T cells and anti-idiotypic antibodies.^{330,331} In fact, anti-idiotypic antibodies binding to autoantibodies are detectable in healthy individuals.³³² These anti-idiotypic antibodies may induce autoantibody clearance and inhibit autoantibody binding to their anatomical targets.³³³

Moreover, anti-idiotype responses can also be induced. Mice immunized with murine autoreactive antibodies against *N*-glycolylated gangliosides develop anti-idiotype B cell responses which can stimulate cytotoxic T cells in an antigen-specific manner.³³⁴ Furthermore, serum transfer from autoantibody-immunized rabbits protects guinea pigs from developing autoimmune nephritis.³³⁵ These findings suggest that it may be possible to induce anti-idiotype responses also against ACPA. ACPA are diverse antibodies which may have various idiotypes. It is therefore

interesting to gain knowledge on dominant idiotypes of ACPA which then can be used to make either active or passive immunization which targets ACPA-expressing B cells (Figure 4c).

9.6 Cell-based therapy

Cell-based therapy can ameliorate RA by direct killing of autoreactive B cells or by inducing autoantigen-specific tolerance. Depletion of autoreactive B cells can be achieved by adoptive cell therapy using CAAR T cells. Engineered human cytotoxic T cells which express chimeric autoantibody receptors (CAAR) consisting of an autoantigen fused with T cell signalling domains selectively eliminate desmoglein 3-specific B cells in a mouse model of pemphigus vulgaris (Figure 4d).³³⁶ The use of CAAR technology circumvents the possible restriction mediated by the HLA genotype of the patients. As citrullination is a posttranslational modification, methods to conjugate a citrullinated peptide to the CAAR signalling domain construct will be needed to enable this technology for RA treatment. Alternatively, the antigen binding domain of anti-ACPA idiotype antibody can replace the need of a citrullinated peptide. However, as discussed previously, the ACPA idiotypes and the antibodies that recognize these are still unknown.

Other cell-based RA therapies aim to induce tolerance in either antigen-specific or non-specific manners. Antigen-specific tolerance can be induced by autologous transfer of modified, tolerogenic dendritic cells (DCs) which are exposed to citrullinated peptides *ex vivo* (Figure 3c). In a phase 1 trial of this DC-based tolerogenic vaccine, the ratio of Treg to effector T cells was increased, T cell responses against a citrullinated peptide were inhibited and the disease activity score within one month was decreased.³³⁷ This trial provides a rationale to develop DC-based vaccine against ACPA-positive RA.

Besides autologous DC, regulatory T cell (Treg) and mesenchymal stem cell (MSC) transplantation are also promising strategies to induce immunological tolerance in autoimmune diseases (Figure 2b). Tregs promote peripheral immunological tolerance through competition with effector T cells for activating signals, production of immunosuppressive molecules and cytolysis of DC.³³⁸ In RA synovial fluid, Tregs are enriched.³³⁹⁻³⁴¹ Their immunoregulatory capacity is, however, impaired in the inflammatory milieu.^{172,342} Moreover, the activated phenotype of T and B cells in the synovial tissue confers relative resistance towards immunoregulation by Tregs.³⁴¹ To enhance the number and functionality in the inflammatory arthritic milieu, Tregs can be treated with a combination of anti-CD28 and IL-

2, as well as all-trans retinoic acids *ex vivo*.^{343,344} Moreover, inhibition of IL-6 and TNF alpha also recovers Treg functions.^{172,342}

Tregs require TCR activation by antigen-MHC complexes to induce their immunoregulatory capacity. Once activated, they can exert their suppressive properties in an antigen-independent manner.³⁴⁵ In experimental autoimmune encephalitis, a model of MS, adoptive transfer of myelin basic protein-reactive Treg improves symptom recovery and prevents relapse after disease onset.³⁴⁶ Polyclonal Tregs are, however, unable to exert similar protection. To gain sufficient numbers of autoantigen-specific Treg, autoantigen-specific TCR gene transfer can be employed before transferring the cells into the patient. Indeed, adoptive therapy using antigen-specific *TCR* and *FOXP3* gene-transferred CD4⁺CD25⁺ T cells to a mouse model of arthritis suppresses arthritic bone destruction (Figure 3d).³⁴⁷

Antigen-nonspecific cell therapy using polyclonal Tregs can also be beneficial to induce immunological tolerance without suppressing systemic immunity. Adoptive transfer of CD4⁺CD25⁺ T cells reduces disease progression of arthritis and reverses type 1 diabetes in mouse models of these respective diseases.^{343,348} Another antigen-nonspecific cell therapy using mesenchymal stem cell (MSCs) transplantation has also been trialled for therapy against autoimmune diseases. MSCs are non-hematopoietic progenitor cells which reside in many tissues to replace injured cells. These cells possess immunosuppressive and anti-inflammatory properties, which contribute to tissue homeostasis.³⁴⁹

Autologous MSCs transplantation is efficacious in treating refractory autoimmune diseases by restoring the frequency of Tregs and inducing anti-inflammatory phenotypes of macrophages as well as autoreactive T cells (Figure 2b).³⁵⁰ Moreover, MSCs inhibit pro-inflammatory responses by RA synovial cells.³⁵¹ Umbilical cord-derived MSC transplantation in RA patients decreases serum levels of IL-6 and TNF-alpha, increases the frequency of circulating Tregs and reduces disease activity.³⁵² In a phase 1b/2a clinical trial, adipose-derived MSC transplantations showed a trend for clinical efficacy in patients with active refractory RA.³⁵³ In addition, allogeneic MSC transplantation ameliorated disease activity in refractory lupus patients in a pilot clinical study.³⁵⁴ These data suggest that despite being antigen-nonspecific, polyclonal Tregs and MSCs may be able to induce immunological tolerance in autoimmune diseases.

In conclusion, the breach of tolerance towards citrullinated autoantigens and activated ACPA-expressing B cells have central pathogenic roles in the



development and chronicity of ACPA-positive RA. There are several strategies to induce immunological tolerance in these patients. Patient's data suggest that immune regulatory cells are present in the RA synovial fluid and tissue. However, these cells fail to exert their immunomodulatory properties, most likely due to the inflammatory arthritic milieu.^{172,342,355} Therefore, combinatorial therapeutic strategies to deplete existing inflammation-inducing ACPA-expressing B cells and to tolerize citrullinated antigens may provide both short- and long-term protection against ACPA-positive RA.

10 Scope of this thesis

This thesis is divided in two parts, namely the characterization of ACPA-expressing B cells which is described in Chapters 2 and 3; and the development of strategies to specifically target ACPA-expressing B cells which is described in Chapters 4, 5 and 6.

10.1 The characterization of ACPA-expressing B cells

Epidemiological studies on ACPA and clinical studies using rituximab for the treatment of RA give strong indications that ACPA-expressing B cells may play major roles in the development of RA. However, the characteristics of these rare cells were largely unknown. Using previously developed staining method to identify these cells for flow cytometry, direct characterization of ACPAexpressing B cells was conducted. Chapter 2 describes the phenotypic and functional characterization of ACPA-expressing B cells. First, the percentage of ACPA-expressing B cells and their development states in both blood and synovial fluid of RA patients are described. Then the expression of activation markers such as CD19, HLA-DR, CD80, CD86 and Ki-67 on these cells from ACPA-positive patients with arthralgia (pre-disease state), early immunosuppressor-naïve RA and established immunosuppressor-treated RA is investigated. These characteristics are subsequently compared to those of tetanus toxoid (TT)-specific B cells in both quiescent and activated states. Next, the production of proinflammatory cytokines by ACPA-expressing B cells in the blood and synovial fluid is explored. This chapter highlights that in RA, ACPAexpressing memory B cells are proliferative and active which are not observed in the arthralgia state. These cells also highly express co-stimulatory molecules, produce abundant pro-inflammatory cytokines, and differentiate into plasma cells at the site of inflammation. These findings argue for direct roles of ACPA-expressing B cells in the pathogenesis of RA. To understand how these cells sustain their activated state, their expression of an essential immune checkpoint

receptor CD32B is also studied. This chapter describes that the expression of CD32 on ACPA-expressing memory B cells (MBC) from RA patients is downregulated. **Chapter 3** further investigates the expression of immune checkpoint receptors on ACPA-expressing MBC including CD5, PECAM-1, CD200R, LAIR-1, FcRL4, CD22 and Fas. Dysregulation of any of these immune checkpoint receptors leads to aberrant B cell function and/or the formation of autoantibodies in murine models. This chapter highlights the fact that the expression of immune checkpoint receptors on ACPA-expressing MBC was generally similar to that on quiescent TT-specific MBC with an exception of Fas which is highly expressed by ACPA-expressing MBC. The high expression of Fas and the downregulation of CD32 on these cells are mimicked by recently boosted TT-specific MBC. Importantly, more than 90% of ACPA-expressing MBC express CD22 which will be targeted in **Chapter 5**.

10.2 Development of strategies to specifically target ACPA-expressing B cells

Due to their pathologic characteristics, ACPA-expressing MBCs are good targets for RA treatment. Therefore, three strategies are described to inhibit or eliminate these pathologic cells specifically while conserving protective MBCs. Chapter 4 investigates the feasibility of a sequential prodrug strategy to selectively eliminate these pathologic cells using autoantigen-drug conjugate. In this chapter, we test a non-B cell specific cytotoxic drug saporin which is a ribosome inactivating protein. It is hypothesized that a cyclic citrullinated peptide (CCP)-saporin conjugate will be captured by the BCR of ACPA-expressing B cell, get internalized into endolysosome compartment and saporin will be released into the cytosol of the cells to exert its cytotoxicity. A potential problem of this strategy is that the CCP can bind to circulating ACPA. Subsequently, the resulting immune complexdrug conjugate will eventually bind and eliminate Fc receptor-expressing cells. To prevent the circulating ACPA from binding the CCP-saporin conjugate, we aim to block the citrullinated part of the peptide so that it cannot bind to circulating ACPA. ACPA recognition should then be restored upon introduction of an activating drug only when the prodrug is in the vicinity of B cells. This chapter not only elaborates this concept but also shows *in vitro* that the blocking of citrulline residue using carboxy-p-nitrobenzyl (CNBz) group and its activation using nitroreductase occur with 100% efficiency. Moreover, this chapter shows that both CCP-saporin and nitroreductase-treated CNBz-containing CCP-saporin conjugates selectively eliminate ACPA-expressing B cell line in vitro. Chapter 5 explores the second strategy which aims to activate the immune checkpoint receptor CD22 specifically on ACPA-expressing B cells using polyisocyanopeptides (PIC) multivalent scaffold.



As discussed in chapter 3, ACPA-expressing MBC highly express CD22 which can be used as a target. A PIC construct containing both CD22 ligand and CCP can inhibit IL-8 production by ACPA-expressing B cell line as compared to those treated with separate PIC-CCP and PIC-CD22 ligand at the same concentrations. **Chapter 6** investigates the feasibility of another autoantigen-drug conjugate. Using CCP covalently conjugated to a specific inhibitor of Bruton's tyrosine kinase, acalabrutinib, we aim to selectively inhibit the activation and survival of ACPA-expressing MBC by blocking the BCR signalling in antigen-specific manner. CCP-acalabrutinib conjugate induce partial cytotoxicity towards ACPA-expressing B cell line but not to tetanus-specific B cell line. This selective cytotoxicity, however, is not mediated by Btk inhibition. Therefore an off-target effect may play role in the selective toxicity of CCP-acalabrutinib. Finally, **chapter 7** provides a summary and general discussion of the findings in the thesis.

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