



P2X7 receptor: A potential therapeutic target for autoimmune diseases

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ABSTRACT

P2X7 receptor (P2X7R), a distinct ligand-gated ion channel, is a member of purinergic type 2 receptor family with ubiquitous expression in human body. Previous studies have revealed a pivotal role of P2X7R in innate and adaptive immunity. Once activated, it will mediate some vital cascaded responses including the assembly of nucleotide-binding domain (NOD) like receptor protein 3 (NLRP3) inflammasome, non-classical secretion of IL-1 β , modulation of cytokine-independent pathways in inflammation such as P2X7R- transglutaminase-2 (TG2) and P2X7R-cathepsin pathway, activation and regulation of T cells, etc. In fact, above responses have been identified to be involved in the development of autoimmunity, specifically, the NLRP3 inflammasome could promote inflammation in massive autoimmune diseases and TG2, as well as cathepsin may contribute to joint destruction and degeneration in inflammatory arthritis. Recently, numerous evidences further suggested the significance of P2X7R in the pathogenesis of autoimmune diseases, including systemic lupus erythematosus (SLE), rheumatoid arthritis (RA), inflammatory bowel disease (IBD), multiple sclerosis (MS), etc. In this review, we will succinctly discuss the biological characteristics and summarize the recent progress of the involvement of P2X7R in the development and pathogenesis of autoimmune diseases, as well as its clinical implications and therapeutic potential.

1. Introduction

Purinergic signaling, a highly evolutionary conserved signaling system, which regards extracellular adenosine triphosphate (ATP), associated nucleotides and adenosine as transmitter molecules, acts a crucial role in considerable physiological processes and pathological reactions [1,2]. At present, two types and at least 24 subtypes of purinergic receptors have been described, among which P2X7 receptor (P2X7R) is the most extensively researched subtype from an immunological viewpoint [2,3]. Interestingly, the activation of P2X7R depends on the concentration and time of ATP stimulation, as well as the cell type [4]. Temporary treatment with low ATP concentration causes typical cation-selective channel opening that facilitates a prompt efflux of K⁺ and influx of Na⁺ and Ca²⁺ leading to changes in the ionic homeostasis of the cell, whereas prolonged challenge with higher concentration triggers formation of the large non-selective pore, which additionally allows the passage of molecules up to 900 Da via an uncertain mechanism [3,5].

The immune system comprises of innate and adaptive immunity, of which inappropriate or insufficient responses may result in visceral or systemic dysfunctions [6]. The first-line defense against challenge of pathogens is provided by the innate immune system, which

encompasses varied cell types including macrophages, dendritic cells (DCs), etc. [4] through pathogen-associated molecular patterns (PAMPs) recognized by pattern recognition receptors (PRRs) [7]. As a starting point for inflammation, the P2X7R exerts an indispensable role in sensing the purinergic danger that is called “alarmins” and considered as a distinct subset of endogenous danger signals and leads to autoimmune responses [4,8–11]. Autoimmune diseases manifest a large quantity of conditions characterized by errant, chronic auto-inflammation against self-tissues, whose the well-known representations are systemic lupus erythematosus (SLE), rheumatoid arthritis (RA), ankylosing spondylitis (AS), Sjögren's syndrome (SS), systemic sclerosis (SSc), etc. [12]. P2X7R, as a particular P2X receptor for its downstream signaling that is coupled with pro-inflammatory cascades such as inflammasome assembly and maturation and non-classical secretion of IL-1 β [5,13], plays a pivotal role in immunity and autoimmunity [3]. Current researches concerning P2X7R have partly revealed its essential mechanisms in inflammation-related disorders, which means that the P2X7R may open a brand-new door for anti-inflammatory therapy, naturally including autoimmune diseases [13–16].

In our review, we will succinctly summarize the immunological functions of P2X7R, as well as its role and significant therapeutic potential in autoimmune diseases.

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2. P2X7R and its immunological functions

P2X7R is a distinct member of P2X receptor subclass, which are ligand-gated ion channels belonging to the purinergic type 2 receptor family (P2) [13,17]. As a homomeric oligomer, it is formed by the assembly of the same three P2X7 subunit containing 595 amino acids, whose gene is located on human chromosome 12q24.31 with 53 kb spanning [3,18]. The structure of P2X7 subunit is two membrane spanning domains (TM1 and TM2) linked by an extracellular loop embracing ten conserved cysteine residues, an intracellular N termini, and a significantly longer intracellular C termini compared to that in the other P2X receptors [19]. TM1 and TM2 involve in the formation of the channel pore. The N and C termini determine the kinetics of receptor desensitization and pore expansion, as well as play a vital role in modulation of receptor functions by intracellular messengers, kinases, and reactive oxygen species (ROS), etc. [20]. Extracellular loop is thought to form disulfide bonds and serve as binding sites for transition metals that modulate P2X7R, as well as participate in the activation by ATP [19,21].

ATP, as the only known physiologic agonist for P2X receptors (P2XRs), can be released via non-physiological necrotic cell death or stress, plasma membrane injury and non-lytic pathways [3,4], such as pannexins [22], ATP-binding cassette (ABC) transporters [23], PRRs activation [24], secretory vesicles [25] and P2X7R itself [26]. Besides, several researches indicated that human cathelicidin-derived peptide LL-37 may be also a P2X7R agonist [13]. The P2X7R is widely expressed in numerous cells and nearly all tissues of the body including monocytes/macrophages [27,28], T cells [29], mast cells [30], fibroblasts [31], epithelial cells [32], microglia [33], endocrine and exocrine pancreas [34], etc., among which the distribution in the immune and inflammatory system cells of monocyte-macrophage lineage is the highest [28]. The activation of P2X7R requires three ATP molecules binding to its extracellular domain, reported by a series of electrophysiological data [35]. As mentioned above, compared to short-time treatment of low ATP concentration, prolonged stimulation with higher concentration leads to the channel-to-pore transition, which results in depolarization and permeability change of membrane, decreased cytosolic K⁺, increased cytosolic Ca²⁺ and Na⁺, as well as the flux of some other molecules such as bacterial products (PAMPs) entry and large-scale intracellular ATP release [5,36]. Though the exact molecular mechanism of channel-to-pore transition remains incompletely clear, several hypotheses have been convinced as following: a). intrinsic pore formation function via channel expansion. b). recruitment of membrane hemi-channels such as pannexin-1 [20,36–38]. Additionally, P2X7R activity is also regulated by cations and anions in extracellular medium [20,39]. Given that its ubiquitous expression on nearly all immune cell types [40], P2X7R is involved in both innate and adaptive immunity.

During innate immune responses, the key role of P2X7R is to activate the assembly of nucleotide-binding domain (NOD) like receptor protein 3 (NLRP3) inflammasome rapidly, which could consecutively facilitate caspase-1 mediated maturation and release of the pro-inflammatory cytokines interleukin-1 β (IL-1 β) and interleukin-18 (IL-18) [6,10]. The activation of NLRP3 inflammasome by P2X7R may result from decreased cytosolic K⁺ [13,41] and increased cytosolic Ca²⁺ [42,43], which could be caused by P2X7R activation resulting in channel opening or pore formation. Mitochondrial DNA (mtDNA) from damaged mitochondrial caused by cellular change in ionic homeostasis [42,44,45] and ROS related to P2X7R activation and mitochondrial damage [46,47] are also well-established. Meanwhile, P2X7R activation induced-pore allows entry of bacterial products (PAMPs) and ATP leading to production of ROS to drive inflammasome formation, as well as passage of hydrophilic solutes into cytosol, which is thought to cause cell death resulting in more ATP release [17,36]. Moreover, the direct interaction between P2X7R and NLRP3 inflammasome should not be neglected [11]. From another perspective, to increase the several key

genes products in inflammatory cascades including pro-IL-1 β , pro-IL-18 and inflammasome components such as NLRP3 and apoptosis-associated speck-like protein (ASC), P2X7R stimulation is regarded as the second signal by inducing K⁺ efflux, inflammasome assembly, and caspase-1 activation while PRRs such as Toll-like receptor (TLR) mediated nuclear factor kappa B (NF- κ B) pathway activation acts as the first signal [2]. Besides, IL-1 β and IL-18 lack a signal sequence for classical cellular secretory vesicles, the P2X7R also plays a key role in the mechanism of non-classical secretion of IL-1 β including the exosome pathway, microvesicle pathway and autophagy pathway, etc. [5,48]. Importantly, transglutaminase-2 (TG2) externalization in conjunction with thioredoxin-1 (Trx), which is mediated by P2X7R activation via member pore formation of P2X7R itself rather than ion channels or pannexin-1 channels, could modify immunogenic peptides to promote the activation and homing of T and B cells, thus contributes to several autoimmune conditions, such as inflammatory arthritis [49,50]. Additionally, P2X7R also modulates several intracellular signaling pathways such as MyD88/NF- κ B pathway and activation of mitogen-activated protein kinase (MAPK) pathway proteins [51,52].

During adaptive immune responses, intercellular interactions between T cells and antigen-presenting cells (APC) through the immune synapse induces intracellular ATP release, which serves as a local autocrine or paracrine signaling molecule and the initiation of T cell activation [53,54]. By responding to T cell receptor (TCR)-mediated released ATP, P2X7R participates in the activation and proliferation of T cell via triggering Ca²⁺ influx, which is required for activation of nuclear factor of activated T cells (NFAT) leading to the expression of interleukin-2 (IL-2) [54,55]. Furthermore, P2X7R modulates the balance between the helper T 17 (Th17) and regulatory T (Treg) cells [29,56]. Activation of ATP-P2X7R pathway impairs the suppressive activity and stability of Treg cells and inhibits the differentiation of naive CD4⁺ T cells into Treg cells, as well as promotes the conversion of Treg cells into Th17 cells [29].

On balance, IL-1 β and IL-18, as the main products of P2X7R activation during innate immunity, contributing to differentiation of Th17 and Th1 cells [57], decrease of Treg cells [58], enhanced expression of IL-2 receptor in lymphocytes and increased production of antibody by B cell amplification [59], are main pro-inflammatory agonists in inflammatory cascade [60]. Treg cells account for the maintenance of immunological self-tolerance that inhabits autoimmune responses, whereas Th17 cells, a distinct CD4⁺ T cell lineage, contribute to the induction and progression of inflammation by producing IL-17 [61,62]. IL-1 β and IL-6, especially IL-6, could mediate the conversion from Treg cells to Th17 cells [58]. The disequilibrium between Th17 cells and Treg cells favoring pro-inflammatory Th17 cells has been reported to be involved in numerous autoimmune disorders including systemic lupus erythematosus (SLE), rheumatoid arthritis (RA), inflammatory bowel disease (IBD), ankylosing spondylitis (AS), multiple sclerosis (MS), etc. [61]. Briefly speaking, P2X7R is implicated in recognizing endogenous danger signals and acting a vital role in inflammation-related disorders, as well as the immunity (Fig. 1).

3. P2X7R in autoimmune diseases

Autoimmune diseases comprise a cluster of at least 80 disorders with severe clinical syndromes, which share the common characteristic of insufficient self-tolerance contributing to an immune-mediated attack on the body's own cells, tissues and organs [61,63]. In spite of emerging target therapy progressing, the therapeutic effects and prognosis remains to be improved [64–66]. Accordingly, to explore an original and efficient therapeutic target for autoimmune patients is imperative. Based on the same hallmark of vitiated self-tolerance, the P2X7R has been reported to be implicated in the pathogenesis and progression of several autoimmune diseases, including SLE [67], RA [68], IBD [69], MS [70], and other disorders such as SS [71], psoriasis [72] and SSC [73] (Table 1).

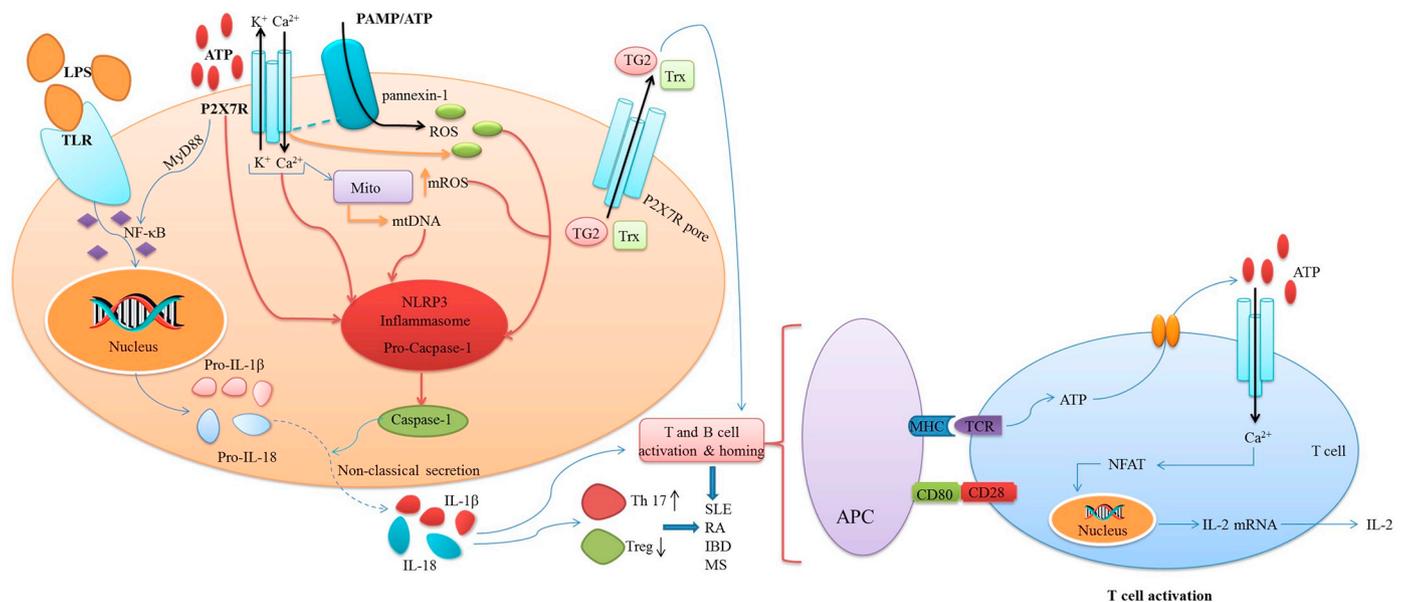


Fig. 1. Mechanisms of P2X7 receptor action.

Activation of PRR such as TLR mediated NF- κ B or MAPK pathway up-regulates the expression of several key genes products in inflammatory cascade including pro-IL-1 β , pro-IL-18 and inflammasome components.

Activated by ATP or LL-37, the P2X7R facilitates channel opening or pore formation resulting in efflux of K⁺ and influx of Ca²⁺ or additional flux of other molecules such as bacterial products (PAMPs) and ATP. Two hypotheses of mechanism of channel-to-pore transition are convinced: (1) intrinsic pore formation function via channel expansion. (2) recruitment of membrane hemi-channels such as pannexin-1. The models of NLRP3 inflammasome activated by P2X7R are as following: (1) Decreased cytosolic K⁺ and increased cytosolic Ca²⁺ trigger NLRP3 activation. (2) P2X7R stimulates the production of ROS, which contributes to the activation of NLRP3 inflammasome. (3) Changes of ionic homeostasis especially Ca²⁺ cause mitochondrial damage, including the release of mtDNA into the cytosol, increased production of mROS, and loss of membrane potential. Both mtDNA and mROS lead to NLRP3 activation. (4) The entry of bacterial products (PAMPs) and ATP via P2X7R activation-induced pore leads to ROS production to drive inflammasome formation. (5) P2X7R induces NLRP3 activation via direct mutual interaction. Activation of NLRP3 inflammasome converts pro-caspase-1 into mature caspase-1, which cleaves the pro-inflammatory cytokines such as IL-1 β and IL-18.

TG2 externalization in conjunction with Trx is mediated by P2X7R activation via member pore formation of the P2X7R itself rather than ion channels or pannexin-1 channels.

By responding to TCR-mediated released intracellular ATP, P2X7R participates in the activation and proliferation of T cell via triggering Ca²⁺ influx, which is required for activation of NFAT leading to the expression of IL-2.

PRR: sensor-pattern-recognition receptor; TLR: Toll-like receptor; NF- κ B: nuclear factor kappa B; MAPK: mitogen activated protein kinase; ATP: adenosine triphosphate; PAMPs: pathogen-associated molecular patterns; NLRP3: nucleotide-binding domain (NOD) like receptor protein 3; ROS: reactive oxygen species; mtDNA: mitochondrial DNA; mROS: mitochondrial ROS; TG2: transglutaminase-2; Trx: thioredoxin-1; TCR: T cell receptor; NFAT: nuclear factor of activated T cells; IL-2: interleukin-2; APC: antigen-presenting cells; SLE: systemic lupus erythematosus; RA: rheumatoid arthritis; IBD: inflammatory bowel disease; MS: multiple sclerosis.

3.1. Systemic lupus erythematosus (SLE)

Systemic lupus erythematosus (SLE) is a systemic autoimmune disorder featured with anti-nuclear antibodies production, complement consumption and immune complex activation, which leads to immune responses against self-antigens and causes multiple organs damage [74]. Along with researches deepening, though the specific etiology and pathogenic mechanisms of SLE is still incompletely elucidated, P2X7R is thought to have a potential involvement in the pathogenesis and development of the disease [75].

As a wide-used experimental mouse model of SLE, MLR/*lpr* strain manifests a closely similar autoimmune syndrome to human SLE and progressive lymphoproliferation due to the *lpr* mutation leading to impairment of the Fas receptor [76]. Zhao et al. [77] reported significantly increased expression of P2X7R-NLRP3 inflammasome signaling molecules including P2X7R, NLRP3 and ASC, as well as caspase-1-p20 subunit that represents the highly active forms of processed caspase-1 in the kidneys of MLR/*lpr* mice. Kahlenberg et al. suggested that caspase-1 might be involved in SLE pathogenesis via a pristane-induced lupus (PIL) model. Caspase-1^{-/-} mice showed decreased auto-antibodies and reduced vascular damage compared to caspase-1^{+/+} mice [78]. Meanwhile, in lupus nephritis (LN) patients, Tuener et al. [79] showed that up-regulated P2X7R expression in glomeruli and renal tubules contributes to autoimmune-related glomerulonephritis through apoptosis or the regulation of pro-inflammatory cytokines. From

another perspective, the phenomenon can be further confirmed by the fact that genetic deletion of P2X7R (P2X7R-KO) mice exhibited obvious protection against antibody-mediated glomerulonephritis [80]. Additionally, Kahlenberg et al. [81] also demonstrated that neutrophil extracellular traps (NETs) and LL-37-mediated activation of the inflammasome leading to autoimmune syndromes are increased in macrophages of lupus patients, and during above processes, P2X7R activation is required. At the gene level, human P2X7R gene locus at 12q24 called SLEB4 as a novel SLE susceptibility locus in Hispanic and European American families has been detected [74]. Besides, Chen et al. [82] also reported that single-nucleotide polymorphism (SNP) in the P2X7R gene (rs1718119) was associated with susceptibility to SLE in a Chinese population through a case-control study. However, the difference in the allele frequencies of 1513 AC SNP of P2X7R gene between sporadic SLE patients and controls was not found in Caucasian populations [83]. Interestingly, the 1513 AC SNP was also found to be associated with low expression of P2X7R in SLE patients, indicating an involvement of P2X7R in some other different mechanisms [75].

Indeed, Le Gall et al. [76] have found impaired P2X7R activity in T cells from autoimmune MRL/*lpr* mice. The converse situations of P2X7R expression are not mutually exclusive but contribute to similar autoimmune syndromes ultimately. As we know, autoimmune diseases could also be induced by defective homeostasis modulation of effector T cells [84]. In MRL/*lpr* mice, Le Gall et al. [76] identified that the loss of P2X7R expression on T cells was tremendously associated with the

Table 1
Linkages of several autoimmune diseases to P2X7R.

| Disease | Publication year | Author | Subjects | Linkages to P2X7R | Ref | |
|---------|------------------|---------------------------|---|---|--|-------|
| SLE | 2004 | Nath et al. | SLE patients | P2X7R gene locus at 12q24 (SLEB4) was a SLE susceptibility locus | [74] | |
| | 2007 | Turner et al. | LN patients | Increased expression of P2X7R in the renal biopsy tissue | [79] | |
| | 2008 | Forchap et al. | SLE patients | A loss-of-function of 1513 AC SNP of P2X7R gene was not a SLE susceptibility locus | [83] | |
| | 2009 | Taylor et al. | P2X7R-KO mice | P2X7R deficiency attenuated renal injury in GN | [80] | |
| | 2010 | Portales-Cervantes et al. | SLE patients | Low expression of P2X7R was associated with the 1513 AC SNP of P2X7R gene | [75] | |
| | 2012 | Le Gall et al. | MLR/lpr mice | Impaired P2X7R activation in T cells was involved in a T-cell homeostatic pathway | [76] | |
| | 2013 | Zhao et al. | MLR/lpr mice | Enhanced expression of P2X7R and related signaling pathway molecules in the kidneys | [77] | |
| | 2013 | Chen et al. | SLE patients | SNP in the P2X7R gene (rs1718119) associated with susceptibility to SLE | [82] | |
| | 2013 | Kahlenberg et al. | SLE patients and mice | Activation of P2X7R was required for the IL-37-mediated inflammasome activation | [81] | |
| | 2019 | Faliti et al. | SLE patients | Reduction of P2X7R mRNA and impaired response of Tfh cells to P2X7R stimulation | [85] | |
| | RA | 2002 | Dell'Antoni et al. | CFA-induced mouse model | P2X7R mediated the transmission of the inflammatory pain in arthritis | [102] |
| | | 2002 | Labasi et al. | Mice | P2X7R-deficient mice manifested reduced incidence and severity of arthritis | [135] |
| | | 2008 | Caporali et al. | RA patients | RA patients type B synoviocytes expressed functional P2X7R modulating IL-6 release | [93] |
| | | 2008 | Christenson et al. | RA patients | Increased neutrophils survival mediated by P2X7R promoted inflammation of RA | [101] |
| 2010 | | Portales-Cervantes et al. | Human neutrophils | Increased expression of P2X7R in PBMCs from RA patients as compared to controls | [75] | |
| 2010 | | Lopez-Castejon et al. | Human macrophages | P2X7R mediated cathepsins release as a cytokine-independent mechanism in pathogenesis of RA | [97] | |
| 2011 | | Al-Shukaili et al. | RA patients | The SNP of P2X7R gene at position 1068 and 1513 conferred susceptibility to RA | [88] | |
| 2012 | | Portales-Cervantes et al. | RA patients | The 489C > T SNP of P2X7R gene enhanced the function of P2X7R and may contribute to the pathogenesis of RA | [89] | |
| 2012 | | Keystone et al. | RA patients | P2X7R inhibitor AZD9056 was not efficient for the treatment of RA | [136] | |
| 2012 | | Stock et al. | RA patients | P2X7R antagonist CE-224,535 was not efficacious for the treatment of RA patients with an inadequate response to MTX. | [137] | |
| IBD | 2014 | McInnes et al. | Rat SCW model | P2X7R expression was detected in inflamed synovial tissue after onset of arthritis | [91] | |
| | 2016 | Fan et al. | RA patients | Inhibition of P2X7R reduced articular inflammation and erosive progression | [86] | |
| | 2018 | Chen et al. | RA patients | Increased expression of P2X7R in PBMCs from patients with RA | [92] | |
| | 2010 | Cesaro et al. | CD patients | Th17 cell differentiation induced by GII required P2X7R signaling in CIA mice | [112] | |
| | 2012 | Kurashima et al. | CD patients | Elevated P2X7R level in the synovial tissue with a great diagnostic value for RA | [30] | |
| | 2012 | Gulbransen et al. | Mice model colitis | Levels of IL-1 β and IL-6 were positively correlated with the levels of P2X7R in the synovial tissues | [111] | |
| | 2014 | Marques et al. | TNBS-induced mouse model | Significantly down-regulated expression of P2X7R in intestinal epithelial cells of CD patients | [108] | |
| | 2014 | Neves et al. | CD patients | P2X7R-mediated mast cell activation in both the initiation and exacerbation of intestinal inflammation and apoptosis | [106] | |
| | 2015 | Hofman et al. | CAC model | P2X7R-pannexin-1 mediated inflammation-induced neuron cell death and contributed to colonic motor dysfunction | [109] | |
| | 2015 | Eser et al. | CD patients | Prophylactic systemic P2X7R blockade prevented experimental colitis effectively | [114] | |
| | 2015 | Miller et al. | C57BL/6J mice | The up-regulation of P2X7R in CD inflamed mucosa was consistent with the involvement of purinoceptors in inflammation and apoptosis | [113] | |
| | 2016 | Wan et al. | P2X7R-KO mice | Evaluation of P2X7R inhibitor AZD9056 lacked potential anti-inflammatory effect but indicated a great value for the relief of chronic abdominal pain | [107] | |
| | 2005 | Narcisse et al. | MS patients | P2X7R inhibited inflammation in protozoan parasite Toxoplasma induced ileitis | [117] | |
| | 2007 | Mattute et al. | MS patients | P2X7R-NLRP3 inflammasome and P2X7R-NF κ B pathway were crucial in pathogenesis of RA | [119] | |
| | 2010 | Grygorowicz et al. | EAE model | Elevated P2X7R expression in normal-appearing axon tracts in MS patients | [121] | |
| | 2011 | Grygorowicz et al. | EAE model | ATP- P2X7R activation killed oligodendrocytes and contributed to the pathogenesis of EAE | [122] | |
| | 2011 | Oyanguren-Desez et al. | MS patients | Overexpression of P2X7R connecting to astrocytic pool of cells was detected at very early asymptomatic phase but relating to neurons emerged in the peak of neurological symptoms | [123] | |
| | 2015 | Gu et al. | MS patients | P2X7R overexpression related to synaptosomal fraction in the symptomatic phase and to the glial fraction in the recovery phase of EAE | [124] | |
| 2017 | Amadio et al. | MS patients | P2X7R-mediated signaling in the pathomechanisms of EAE with the possible relevance of astrocytic pool of cells. Higher frequency of a gain-of-function P2X7R SNP (rs17525809) in MS patients compared to controls | [118] | | |

(continued on next page)

Table 1 (continued)

| Disease | Publication year | Author | Subjects | Linkages to P2X7R | Ref |
|-----------|-------------------|---|--|--|-------|
| SS | 2012 | Woods et al. | Mice | Activation of the P2X7R with ATP or BzATP stimulated the cleavage and release of α -fodrin | [129] |
| | 2013 | Baldini et al. | SMG cell aggregates | ATP-P2X7R activation enhanced immune cell infiltration into the gland and initiated apoptosis of salivary epithelial cells | [128] |
| | | | pSS patients | Significantly higher P2X7R expression in salivary glands from pSS individuals | |
| | 2014 | Xie et al. | pSS patients | P2X7R-inflammasome-caspase-1-IL-18 axis involved in the development of pSS exocrinopathy. | [126] |
| | 2015 | Yu et al. | pSS patients | Substantially higher surface expression of P2X7R on PBMC in pSS patients than controls | [127] |
| 2017 | Khalafalla et al. | CD28 ^{-/-} , IFN γ ^{-/-} , NOD.H-2h4 mouse model | mRNA and protein levels of P2X7R on pSS PBMCs were significantly higher than in normal individuals | [125] | |
| Psoriasis | 2013 | Killeen et al. | Psoriasis patients | Treatment of the P2X7R antagonist A438079 in vivo ameliorated salivary gland inflammation and improved saliva secretion. | [72] |
| | | | | Increased expression of P2X7R in non-lesional and lesional psoriatic skin | |
| SSc | 2017 | Geraghty et al. Gentile et al. | IMQ-induced psoriasis mice SSc patients | mRNA level of P2X7R was significantly higher in non-lesional psoriatic samples | [130] |
| | | | | ATP-P2X7R-miR21 pathway served as a potential pathogenic mechanism for the initiation of psoriasis | |
| | | | | P2X7R was not essential for development of IMQ-induced psoriasis-like inflammation | |
| | | | | Increased P2X7R surface expression with enhanced Ca ²⁺ influx in SSc dermal fibroblasts | [73] |
| | | | | Activation of P2X7R led to pro-fibrotic effects | |

SLE: systemic lupus erythematosus; RA: rheumatoid arthritis; IBD: inflammatory bowel disease; CD: Crohn's disease; CAC: colitis-associated cancer; MS: multiple sclerosis; SS: sjögren's syndrome; pSS: primary sjögren's syndrome; SSc: systemic sclerosis; LN: lupus nephritis; GN: glomerulonephritis; CFA: complete Freund's adjuvant; CII: type II collagen; CIA: collagen-induced arthritis; BMDM: bone marrow-derived macrophages; SCW: streptococcal cell wall; TNBS: 2,4,6-trinitro-benzene-sulfonic acid; DSS: dextran sulphate sodium; EAE: experimental autoimmune encephalomyelitis; SMG: submandibular gland; IMQ: imiquimod; NF- κ B: nuclear factor kappa B; SNP: single-nucleotide polymorphism; PBMCs: peripheral blood mononuclear cells; MTX: methotrexate; Th 17 cell: T helper 17 cell; TGF- β : transforming growth factor- β ; NLRP3: nucleotide-binding domain (NOD) like receptor protein 3; BzATP: 3'-O-(4-benzoyl)benzoyl-ATP.

overexpression of B220, which could damage the T cells homeostatic regulation and cloak the response of normal B220⁺ T-cell subgroups to P2X7R stimulation resulting in drastically decreased sensitivity to ATP-mediated stimulation of P2X7R in T cells, thereby, they were protected from death arising from extracellular ATP. Along with the accumulation of the effector T cells, there will lead to autoimmune syndromes such as LN and T-cells mediated lymphoproliferation. In another study, Faliti et al. [85] revealed that down-regulation of P2X7R that possibly caused by observed reduction of P2X7R mRNA might contribute to a selective T follicular helper (Tfh) cell dysfunction associated with SLE. By using a PIL model characterized by peritoneal lipogranulomas, production of antinuclear antibodies (ANAs) and glomerulonephritis, Faliti et al. also demonstrated that P2X7R deletion significantly exacerbated immunopathology and worsened the disease by enhancing the secretion of autoantibodies. The over-production of these autoantibodies might result from down-regulation of immune checkpoint function during B cell differentiation [85].

3.2. Rheumatoid arthritis (RA)

Rheumatoid arthritis (RA) is a chronic autoimmune disease with elusive etiology that is characterized by chronic synovial inflammation, joint destruction and progressive articular damage, which leads to joint deformity, reduced quality of life and even impaired function of some other organ systems [68,86]. It has been shown that P2X7R acts as a potential site in the participation and regulation of the inflammation in RA [49,87]. The polymorphism at position 1068 and 1513 in the P2X7R gene may serve as a susceptibility gene locus for RA [88]. Another similar study has also demonstrated that 489C > T SNP of P2X7R gene enhanced the function of P2X7R in RA patients and may contribute to the pathogenesis of RA [89]. Interestingly, perhaps due to genetic polymorphism in the P2X7R, the sensitivity of RA patient mononuclear cells to ATP stimulation is higher than healthy controls [90].

Substantially increased P2X7R expression in peripheral blood mononuclear cells (PBMCs) from RA patients as compared to normal controls has been observed in several studies [75,86]. By using the rat streptococcal cell wall (SCW) arthritis model, P2X7R expression was detected in inflammatory synovial tissue after onset of arthritis [91]. In RA patients, Chen et al. [92] have also shown elevated P2X7R level in the synovial tissue with a great diagnostic value for RA. Specifically, functional P2X7R is expressed in human type B synoviocytes from rheumatoid joints, and it may regulate IL-6 release into synovial fluid [93]. Moreover, in the presence of LPS, obviously higher amounts of IL-1 β was detected in PBMCs from RA patients in response to ATP [90], which may arise from increased levels of P2X7R. In fact, Chen et al. [92] further reported that the expression levels of IL-1 β and IL-6 were positively correlated with the levels of P2X7R in the synovial tissues of RA patients.

In addition, accumulating evidence indicates that inflammatory T cells especially IL-17 producing Th17 cells are crucial in pathogenesis of RA [94,95]. Induced by immunization with an emulsion of complete Freund's adjuvant (CFA) and type II collagen (CII), collagen-induced arthritis (CIA) is the most extensively studied model of RA to explore the potential mechanisms of autoimmune disease [96]. Fan et al. [86] found that P2X7R signaling was required for CII-regulated the Th17 cell differentiation both in vivo and in vitro. Using the CIA mice, blocking the P2X7R signaling could reduce the production of IL-17A, ameliorate the hind paw swelling and attenuate CII-mediated joint damage [86].

During the inflammation process in RA, several other cytokine-independent pathways mediated by P2X7R also manifest a pivotal role, including P2X7R-TG2 and P2X7R-cathepsins pathway [49,97]. Accumulation of extracellular TG2 driven by P2X7R could mediate protein modification and crosslinking reactions that involve in the pathogenesis, indicating a potential pathogenic mechanism of RA [49]. Dzhabazov et al. [98] also reported that the presence of active TG2 significantly increases severity of disease in the CIA model. Instead of

inflammasome assembly, P2X7R-TG2 pathway depends on the membrane pore formation of the receptor, which is suggested to be more important than the ion channel activity in inflammation based on recent evidences, elucidating that selectively targeting this activity of the receptor is likely to be more therapeutic effectively with lower risk of side effects [49]. As for P2X7R-cathepsins pathway, cathepsins are lysosomal proteases, which could effectively degrade collagen extracellular matrix leading to joint destruction and involve in the inflammation and degeneration of arthritis [99]. Lopez-Castejon et al. [97] reported that P2X7R activation of human macrophages and mouse bone marrow-derived macrophages (BMDM) induced the release of cathepsin B, K, L, and S resulting in joint damage, which was independent of the presence of the cytokines IL-1 β and IL-18.

Of note, neutrophil apoptosis is important for the regulation of inflammation, P2X7R may also play an indirect role in promoting inflammation of RA via anti-apoptotic signaling in neutrophils [13]. Serum amyloid A (SAA) protein, an acute phase reactant correlating with active joint inflammation, was found to be elevated in circulation and tissues of RA patients [100]. In vitro experiment manifested that SAA purified from plasma of RA patients effectively inhibited α -CD95 (FAS) mAb-induced human neutrophil apoptosis [101]. Furthermore, administration of the P2X7R antagonist oxidised ATP (oATP) completely abrogated the anti-apoptotic effect of SAA, which indicated that SAA could increase neutrophil survival via a P2X7R-dependent mechanism [101].

Interestingly, experiment in CFA-induced arthritis mouse model revealed that P2X7R could mediate the transmission of inflammatory pain based on its expression in peripheral nerve endings and endothelial cells rather than the recruitment of immune cell [102].

3.3. Inflammatory bowel disease (IBD)

Inflammatory bowel disease (IBD), a complicated and multifactorial disorder characterized by chronic relapsing intestinal inflammation, which acts a critical risk factor for colitis-associated cancer (CAC), mainly includes Ulcerative Colitis (UC) and Crohn's Disease (CD) [103,104]. Although the exact etiology remains undetermined, accumulating studies have revealed that P2X7R participated in the pathogenesis of IBD via regulating various cellular behaviors [69] (Fig. 2). Furthermore, genetic factors contribute to IBD, however, the P2X7R polymorphisms His155Tyr, Arg307Gln and Glu496Ala appeared not to be susceptibility factors for CD [105].

Neves et al. [106] demonstrated up-regulated expression of P2X7R colocalizing more with macrophages and dendritic cells in inflamed CD epithelium and lamina propria, as well as higher IL-1 β level. Animal experiment in dextran sulphate sodium (DSS) induced mouse model of colitis showed significantly increased and consistent levels of caspase-1-p10 and IL-1 β . Importantly, subsequent research showed that blockade of P2X7R decreased levels of above factors, which indicated that P2X7R mediated NLRP3/Caspase-1 inflammasome pathway was involved in the initiation of IBD [107]. Additionally, based on the similar levels change of NF κ B and tumor necrosis factor (TNF), Wan et al. [107] also suggested that P2X7R-NF κ B pathway plays a pivotal role in the pathogenesis of IBD. Using the 2,4,6-trinitro-benzene-sulfonic acid (TNBS) induced model, the findings that decreased TNF level in concordance with lower activation of NF κ B of Marques et al. [108] also supported the role of P2X7R-NF κ B pathway.

In CD patients, besides elevated expression of P2X7R, higher level of IL-17 and lower level of IL-10 were detected compared to controls, which were mainly produced by Th17 and Treg cells respectively [106]. Another study in CAC model by Hofman et al. [109] found that P2X7R blockade promoted the accumulation of Treg cells within lesions of the intestine and increased the production of transforming growth factor- β (TGF- β). These all pointed to that the balance between Th17 and Treg cells regulated by P2X7R activation was a crucial pathogenic mechanism in IBD. Meanwhile, increased TGF- β in the case of P2X7R

blockade could stimulate the proliferation of intestinal epithelial cells [109], which suggested that P2X7R activation may aggravate inflammation of IBD through decreasing the level of TGF- β to block the repairment of intestinal epithelium.

As we know, the expression levels of P2X7R differ dramatically depending on the different tissues, among which colonic mast cells (MCs) highly express the P2X7R. Activation of MCs mediated by P2X7R could exacerbate intestinal inflammation via inducing inflammatory cytokines, chemokines and leukotrienes to recruit neutrophils, suggesting a potential new mechanism of IBD that is independent of inflammasome pathway [30]. Kurashima et al. [30] found significantly increased numbers of P2X7R positive MCs in colons of both CD patients and TNBS induced model. They also showed that treatment of P2X7R-specific antibody blocked mast cell activation and subsequent intestinal inflammation.

Additionally, IBD always accompanies with long-term gut dysfunction associated with the alterations to enteric neurons, as well as enteric nervous system, whose abnormalities in development could contribute to IBD [110,111]. Gulbransen et al. [111] reported that P2X7R-pannexin-1 mediated inflammation-induced neuron cell death and colonic motor dysfunction via in vivo models of experimental colitis, which indicated that P2X7R in enteric neurons are crucially involved in gut dysfunction of IBD.

However, some studies demonstrated that the expression of P2X7R was significantly down-regulated in intestinal epithelial cells of CD patients [30,112]. This may be regarded as the acute protective response in IBD based on the fact that P2X7R blockade could stimulate the proliferation of intestinal epithelial cells via enhanced production of TGF- β . Of note, over-proliferation of intestinal epithelial cells could also partly explain why P2X7R acts a critical risk factor for CAC. Conversely, Miller et al. [113] reported that P2X7R inhibit inflammation in protozoan parasite *Toxoplasma* induced ileitis, which is exactly opposite to that P2X7R promotes inflammation in classical models including DSS and TNBS induced model. In addition, another clinical evaluation on safety and efficacy of a P2X7R inhibitor did not get expected anti-inflammatory effect [114]. These all suggested that P2X7R could not be regarded as a simple promoting-inflammatory receptor and it may mediate some other different pathogenesis in different inflammation of IBD.

3.4. Multiple sclerosis (MS)

Multiple sclerosis (MS) is an autoimmune inflammatory disease of central nervous system (CNS), presenting with immune cell infiltration, glia activation, loss of oligodendrocytes and axonal depletion, resulting in extensive demyelination and neurodegeneration [115]. With unknown etiology, it has become evident that P2X7R exerts a pivotal role in pathogenesis of MS based on its widespread involvement in inflammation [116].

The diverse expression of P2X7R in different cell types related to MS is so fascinating. The autopsy of brain tissues by Narcisse et al. [117] detected a high expression of P2X7R on reactive astrocytes. Similarly, in brain samples from secondary progressive MS patients, Amadio et al. [118] also found incremental P2X7R on astrocytes in the parenchyma of frontal cortex. However, in spite of the occurrence of inflammation, Amadio et al. demonstrated that P2X7R was inhibited on peripheral monocytes during the acute phase, even absolutely absent from microglia/macrophages or oligodendrocytes during the secondary progressive phase. Furthermore, P2X7R was decreased in human monocytes after pro-inflammatory stimulation in vitro. In fact, this could be explained by the fact that down-regulation of P2X7R may increase their survival and invasion into the CNS tissue to further lead to the neuroinflammation [118]. Additionally, elevated P2X7R expression was found in normal appearing axon tracts in MS patients, indicating that disordered level of P2X7R likely to precede the emergence of clinical symptom and may be an early risk factor for the development of the

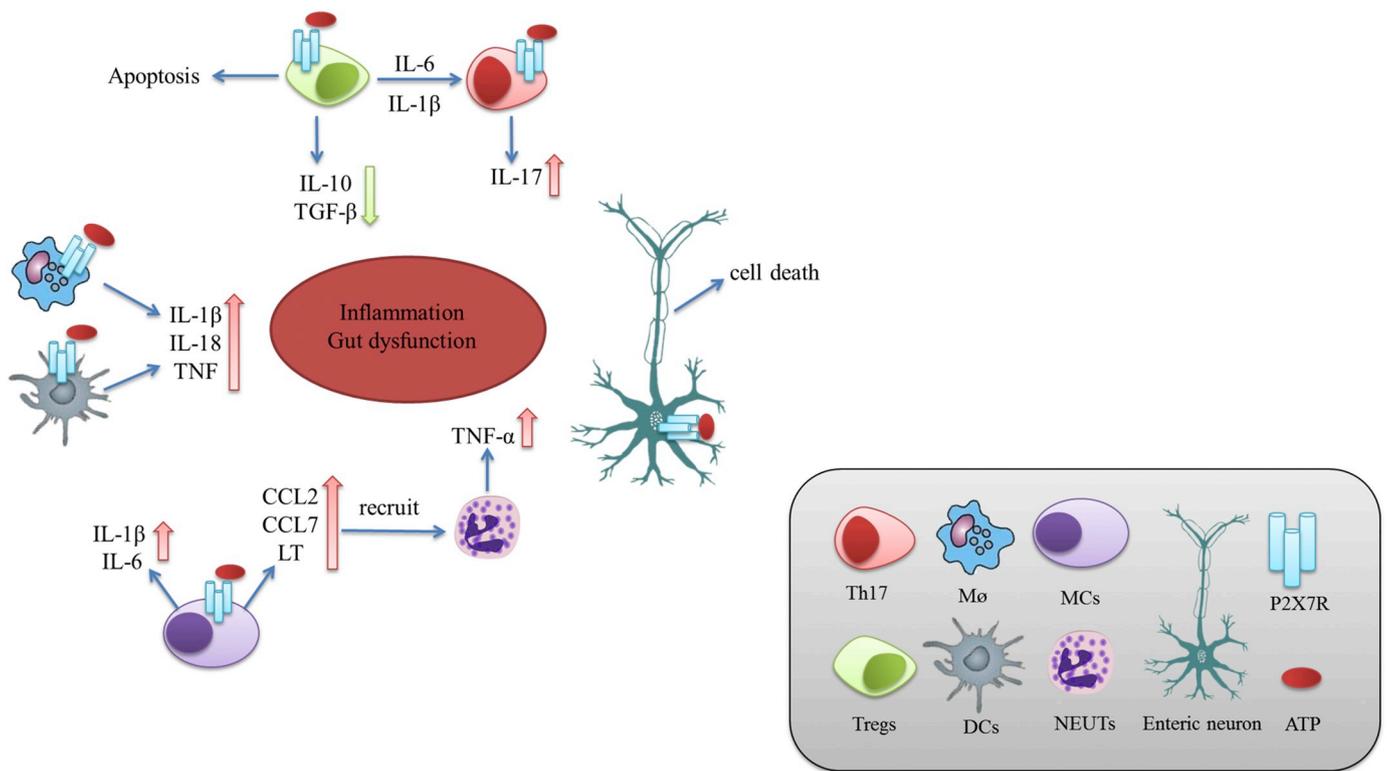


Fig. 2. ATP-P2X7 receptor signals mediated responses in IBD.

(1) In macrophages or dendritic cells, ATP-P2X7R signals lead to the release of IL-1 β and IL-18, as well as TNF through mediating inflammasome and NF κ B pathways. (2) ATP-P2X7R signals promote the release of IL-17 in Th17 cells whereas induce apoptosis and decrease the secretion of IL-10 and TGF- β in Tregs. IL-1 β and IL-6, especially IL-6, could mediate the conversion from Treg cells to Th17 cells. (3) Extracellular ATP induces P2X7R activation in mast cells to exacerbate intestinal inflammation via inducing the release of inflammatory cytokines (IL-1 β , IL-6,) chemokines (CCL2, CCL7, etc.) and leukotrienes (LT) to recruit neutrophils, which could produce massive TNF- α . (4) ATP- P2X7R- pannexin-1 meditates inflammation-induced neuron cell death contributing to gut dysfunction of IBD.

ATP: adenosine triphosphate; IBD: inflammatory bowel disease; TNF: tumor necrosis factor; NF- κ B: nuclear factor kappa B; Th17: T helper 17; TGF- β : transforming growth factor- β ; Tregs: regulatory T cells; M ϕ : macrophage; DCs: dendritic cells; MCs: mast cells; NEUTs: neutrophils.

disease [119].

Several studies have supported that ATP-P2X7R signaling is a crucial factor for the pathology of MS. As a rat model of MS, experimental autoimmune encephalomyelitis (EAE) is the most commonly used to explore the molecular mechanisms accountable for the inflammation and advanced demyelination features [120]. Matute et al. has demonstrated that enhanced ATP signaling via P2X7R could trigger oligodendrocyte excitotoxicity and lead to lesions that resembled the major characteristics of MS including oligodendrocyte loss, demyelination and axonal damage. Importantly, treatment with P2X7R antagonists to EAE rats substantially reduced demyelination and ameliorated the related neurological symptoms effectively [119].

Interestingly, the levels of P2X7R in different cell types are distinct depending on stages of the disease. Overexpression of P2X7R connecting to astrocytic pool of cells was detected at very early asymptomatic phase of EAE, suggesting that P2X7R might involve in the initiation and development of MS via certain pathway [121]. A potential mechanism is that P2X7R-induced cytokine release by glia likely lead to the inflammatory milieu establishment at early stage of the disease [2]. However, the increased expression of P2X7R relating to neurons was observed in the acute phase and peak of neurological symptoms, contributing to progressive inflammation in EAE [121]. Moreover, despite in the recovery phase of EAE, overexpression of P2X7R was still present and reported to be related to the glial fraction [122]. Notably, glial fibrillary acidic protein (GFAP), a marker of astrocyte activation, whose enhanced immunoreactivity in brain sections did not decline to control values in the recovery phase, similarly to P2X7R expression as mentioned above, indicating a role for P2X7R in sustaining the activation of astrocyte [122].

The gene variants of P2X7R may also act a key role in the pathogenesis of MS. A Spanish case-control study reported that the frequency of a gain-of-function P2X7R SNP (rs17525809) in MS patients was significantly higher compared to normal control [123]. Additionally, a meta-analysis of Australasian and European cohorts suggested that the Arg307Gln, which is encoded by a loss-of-function P2X7R SNP (rs28360457), conferred a protection against MS. The protective effect on the risk of MS may arise from dominant impaired pore formation and negative downstream pro-inflammatory responses [124].

3.5. Other autoimmune diseases

Besides above diseases, the P2X7R is also reported to be implicated in the complicated and elusive pathogenesis of several other autoimmune disorders, such as SS, psoriasis and SSc.

Sjögren's syndrome (SS) is a systemic autoimmune disease featured with salivary and lacrimal glands infiltrated by lymphocytes and monocytes leading to acini damage and exocrine secretion failure, predominantly in women [71,125]. Xie et al. [126] and Yu et al. [127] found that the expression of P2X7R on PBMCs in patients with primary SS (pSS) was substantially higher than normal controls. Meanwhile, significantly higher P2X7R expression in salivary glands from pSS individuals was showed by Baldini et al. [128]. Moreover, they also indicated an involvement of the P2X7R-inflammasome-caspase-1-IL-18 axis in the development of pSS pathology [128]. Several results from animal experiments further elucidated the role of P2X7R in SS. In wild-type mice, in vivo administration of P2X7R agonist 3'-O-(4-benzoyl) benzoyl-ATP (BzATP) to ligated submandibular gland (SMG) excretory ducts enhanced immune cell infiltration and triggered salivary

epithelial cells apoptosis, whereas P2X7R^{-/-} mice was absent from the phenomenon [129]. Importantly, treatment of the P2X7R antagonist A438079 to CD28^{-/-}, IFN γ ^{-/-}, NOD.H-2^{h4} mouse model of salivary gland exocrinopathy *in vivo* could reduce salivary gland inflammation and promote saliva secretion, representing a promising therapeutic strategy [125]. Of note, *in vitro* studies using SMG cell aggregates from wild-type and P2X7R^{-/-} mice suggested that activation of ATP/BzATP-P2X7R could stimulate the cleavage and release of α -fodrin, which is thought to serve as an auto-antigen in the development of SS, indicating a possible pathogenic mechanism [129].

Psoriasis is one of the most common chronic inflammatory human skin diseases, presenting with over-proliferation of keratinocytes and infiltration of leukocytes contributing to epidermal hyperplasia (acanthosis) and plaque- or pustular-like skin lesions [130,131]. Killeen et al. have shown increased presence of P2X7R in lesional and non-lesional psoriatic skin compared to healthy tissues. Furthermore, analysis of the P2X7R at the mRNA level indicated that non-lesional psoriatic samples displayed significantly higher P2X7R mRNA than that in lesional samples and healthy controls [72]. Meanwhile, the miR-150, which induces instability of P2X7R mRNA, was also reported to decrease in non-lesional tissues [72,132]. These findings prompted that non-lesional psoriatic tissues may be ready for the conversion to its lesion state via up-regulating the expression of P2X7R. Indeed, following treatment with a P2X7R agonist in non-lesional tissue, the psoriatic responses was initiated. Importantly, based on a series of data, the study also revealed that ATP-P2X7R-miR21 pathway leading to the expression of vascular endothelial growth factor (VEGF) and IL-6 may serve as a potential pathogenic mechanism for the initiation and development of psoriasis lesions [72]. However, another study indicated that P2X7R is not essential for the development of imiquimod (IMQ)-induced psoriasis-like inflammation, nevertheless, this result could not exclude a possible role for P2X7R in the development of human psoriasis or other animal models of this disease [130].

As for systemic sclerosis (SSc), it is a connective tissue disease (CTD) characterized by small vessel alterations and fibroblasts dysfunction, resulting in severe deposition of collagen and extracellular matrix (ECM), tissue fibrosis and even organ failure [133]. Compared to normal subjects, P2X7R surface expression was increased with enhanced Ca²⁺ influx in SSc dermal fibroblasts. In LPS-primed fibroblasts from SSc patients, activation of P2X7R could promote collagen production, α -smooth muscle actin (α SMA) expression, cell migration and connective tissue growth factor (CTGF) release to lead to pro-fibrotic effects. Additionally, the study has also shown that inhibition of extracellular signal-regulated kinases-1/2 (ERK-1/2) prevented P2X7R-induced collagen production and CTGF release, indicating a novel cytokine-independent pathway in P2X7R-induced fibrosis [73].

4. P2X7R as a potential therapeutic target for autoimmune disease

Based on its crucial roles in pathogenesis of inflammation cascade and autoimmunity, P2X7R may serve as a promising therapeutic target for autoimmune diseases. Understanding the molecular mechanisms of P2X7R in autoimmune diseases, together with research to develop more specific P2X7R inhibitor, may provide a novel therapeutic option.

Experimental blockade of P2X7R in animal models has gained auspicious results for several autoimmune diseases. A study was performed by application of P2X7R antagonist brilliant blue G (BBG) or specific small interfering RNA (siRNA) silencing P2X7R to MLR/*lpr* mice. At the macroscopic level, the lifespan of the treated mice was substantially prolonged and the severity of nephritis was ameliorated. At the microscopic level, blockade or absence of P2X7R activation strongly reduced renal immune complex deposition and the Th17:Treg cells ratio. At the molecular level, the inhibition or absence of P2X7R significantly decreased the NLRP3/ASC/caspase 1 assembly, serum levels of IL-1 β and IL-17, as well as the circulating anti-double-stranded DNA (anti-dsDNA) antibodies [77]. Due to the limitation that the

disorder in MLR/*lpr* mice is primarily induced by Fas deficiency, further experiment was carried out in NZM2328 mice, a new mouse model of SLE, in which the disease resembles immune complex-mediated glomerulonephritis [134]. The results manifested a significant reduction of IL-1 β and a twice-confirmation of the effectiveness of BBG treatment, which suggested that development of pharmacologic inhibitors blocking the P2X7R pathway may provide a novel therapeutic strategy for human SLE with general applicability [77].

Several other studies also evaluated the potential of targeting P2X7R in treatment of RA. Using animal models, the incidence and severity of arthritis including joint lesions and collagen degradation production induced by antibody + LPS in wild-type mice were markedly reduced in P2X7R-deficient mice [135]. Similarly, inhibition of P2X7R to SCW-induced model also ameliorated the inflammation and erosive progression of arthritis [91]. In cellular experiments, P2X7R knockdown via a siRNA in an immortalized RA synovial cell line (MH7A) obviously reduced the secretion of IL-1 β and IL-6 [92]. Blockade of P2X7R to two cytokine-independent pathway in RA including P2X7R-TG2 and P2X7R-cathepsin pathway has manifested expected effects: inhibition of P2X7R blocked acute release of a large quantity of TG2 by macrophages with no apparent effects on the level of cellular surface-related enzyme [50], meanwhile, the ATP-induced cathepsin release from macrophages was abolished by P2X7R antagonists (AZ10573295 and AZ11648720) and absent from P2X7R^{-/-} mouse [97]. Additionally, administration of P2X7R inhibitor oATP into inflamed paws of CFA-induced mouse model remarkably relieved the inflammatory pain [102]. However, some clinical evaluations have not achieved the expected efficacy of P2X7R antagonist to the treatment of RA and therein Keystone et al. even showed that P2X7R appeared not to be a therapeutically useful target [136,137]. This may be due to the wide variation of P2X7R gene resulting in the significantly high polymorphism in human population. Therefore, taking P2X7R genotype into consideration is requisite when evaluating the efficacy of P2X7R antagonists.

Besides positive effects of P2X7R blockade in SLE and RA, pharmacological blockade of P2X7R also effectively attenuated intestinal inflammation of TNBS and DSS induced mouse model, as well as CAC mouse model [107–109]. Importantly, P2X7R-KO mice failed to develop colitis while induced by TNBS or DSS [106]. Of note, despite the lack in anti-inflammatory potential of P2X7R inhibitor in a clinical evaluation, the results of the study indicated a great value of P2X7R antagonism for the relief of chronic abdominal pain [114].

Collectively, these evidences suggest that P2X7R may serve as a potential therapeutic target for autoimmune diseases.

5. Conclusion

The advanced researches have amplified our understanding of the pivotal role of P2X7R in inflammation and autoimmune diseases, as well as its intrinsic molecular mechanisms. Although there is much to be explored, accumulating evidences from *in vitro* and *in vivo* model in conjunction with development of P2X7R pharmacology and increased structural insights together support targeting P2X7R as a potential method for the treatment of autoimmune diseases [138,139]. However, several main unresolved aspects in mechanisms and therapeutic value of P2X7R should be noted. One of these is the difference of P2X7R between human and animal models. Firstly, P2X7R of mice could be also activated by NAD⁺, leading to similar downstream cascade responses as compared to that induced by ATP, which may contributed to ligand-associated signaling difference of P2X7R. Next, the affinity of ATP to P2X7R may differ dramatically among diverse species [140]. Meanwhile, significantly high polymorphism of P2X7R in human population may also result in functional distinction. There is another crucial resistance against the therapeutic value of P2X7R. Due to its extensive distribution in human body, the inhibitor relieving symptoms via regulating P2X7R in inflammatory tissues may also result in some

other side effects, which could be called as “Off-target” phenomenon. Therefore, on the one hand, further studies are seriously in need, especially in human systems to explicate the role of P2X7R in autoimmune diseases and evaluate the actual efficacy and safety of existing potential P2X7R inhibitor. On the other hand, to provide more details for new drug development and reduce the side effects, or even absolutely avoid the “Off-target” phenomenon, the different expression of P2X7R in different tissues or status, as well as other responses and physical or pathogenic effects that may be triggered by P2X7R signaling, should be also comprehensively explored. Undoubtedly, a further understanding of biological function and regulation of P2X7R will certainly be beneficial in the development of future therapeutics of autoimmune diseases.

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Declaration of Competing Interest

The authors report no conflicts of interest.

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