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Enhancing acute inflammatory and sepsis treatment: superiority of membrane receptor blockade

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Conditions such as acute pancreatitis, ulcerative colitis, delayed graft function and infections caused by a variety of microorganisms, including gram-positive and gram-negative organisms, increase the risk of sepsis and therefore mortality. Immune dysfunction is a characterization of sepsis, so timely and effective treatment strategies are needed. The conventional approaches, such as antibiotic-based treatments, face challenges such as antibiotic resistance, and cytokine-based treatments have shown limited efficacy. To address these limitations, a novel approach focusing on membrane receptors, the initiators of the inflammatory cascade, is proposed. Membrane receptors such as Toll-like receptors, interleukin-1 receptor, endothelial protein C receptor, µ-opioid receptor, triggering receptor expressed on myeloid cells 1, and G-protein coupled receptors play pivotal roles in the inflammatory response, offering opportunities for rapid regulation. Various membrane receptor blockade strategies have demonstrated efficacy in both preclinical and clinical studies. These membrane receptor blockades act as early stage inflammation modulators, providing faster responses compared to conventional therapies. Importantly, these blockers exhibit immunomodulatory capabilities without inducing complete immunosuppression. Finally, this review underscores the critical need for early intervention in acute inflammatory and infectious diseases, particularly those posing a risk of progressing to sepsis. And, exploring membrane receptor blockade as an adjunctive treatment for acute inflammatory and infectious diseases presents a promising avenue. These novel approaches, when combined with antibiotics, have the potential to enhance patient outcomes, particularly in conditions prone to sepsis, while minimizing risks associated with antibiotic resistance and immune suppression.

KEYWORDS

sepsis, membrane receptor blockade, damage-associated molecular patterns, pathogen-associated molecular patterns, inflammation



1 Introduction

The intricate interplay between acute inflammatory diseases, infectious diseases, and the development of sepsis poses a serious challenge in modern healthcare (1). The timely intervention in these diseases after their inception is of paramount importance (2-4). Without initial treatment, these diseases progress and a substantial number of patients develop sepsis, which is a life-threatening condition due to a dysregulated host response to infection (5). Sepsis has a high mortality rate (6). Sepsis affects nearly 49 million people globally each year, resulting in approximately 11 million deaths. These figures highlight sepsis as one of the major causes of mortality worldwide, accounting for about 20% of all deaths (7). The economic impact is substantial, with global healthcare systems incurring billions of dollars annually due to long hospital stays and intensive medical interventions (8). Particularly in regions with limited healthcare infrastructure, the burden of sepsis is high, and the prevalence of antibiotic-resistant infections further exacerbates these challenges (9). Antimicrobial resistance (AMR) significantly impedes effective treatment options, extending the duration of illness and increasing both mortality rates and healthcare costs (10). Understanding the biological mechanisms that lead from initial infection to sepsis is crucial. This involves delving into the molecular intricacies that govern the immune response during the early stages of these diseases.

Intricate signaling pathways are involved in the early stages of these diseases. Intracellular signaling cascades are initiated by the recognition of endogenous damage-associated molecular patterns (DAMPs) or exogenous pathogen-associated molecular patterns (PAMPs) by pattern recognition receptors (PRRs) (11, 12). This cascade activates immune responses involving cytokines, alarmins, and other mediators, aimed at eliminating pathogens and maintaining cellular homeostasis (13, 14). However, within the context of inflammatory diseases, certain signaling pathways are dysregulated, leading to the overproduction of proinflammatory cytokines and alarmins (15). The resulting excessive systemic inflammation and immune suppression contribute to disease severity (16). Given the complexities of these pathways, it becomes evident why traditional treatment strategies often fall short. This segues

into a discussion of how conventional treatments typically approach these challenges, and why there's a pressing need for innovative therapeutic strategies.

The conventional treatments for sepsis, which includes acute inflammatory and infectious diseases, often involve the use of antibiotics such as cefepime, piperacillin-tazobactam, and ceftobiprole as the primary intervention (17, 18). These antibiotics work by targeting and killing the bacteria responsible for the infection, thus reducing the bacterial load and aiding in infection control. Additionally, standard sepsis treatments include supportive measures like fluid resuscitation, vasopressors, mechanical ventilation, and renal replacement therapy to stabilize patients and maintain vital organ function, but these do not directly address the underlying infection (19, 20). These supportive measures are essential for maintaining patient stability and managing the immediate lifethreatening symptoms of sepsis. However, they do not directly target the underlying infection or the immune dysregulation associated with sepsis. In addition to these challenges, the efficiency of antibiotic treatments may be impaired due to the emergence of antibiotic resistance and delayed drug response. For instance, methicillinresistant Staphylococcus aureus (MRSA) and carbapenem-resistant Enterobacteriaceae (CRE) are examples of antibiotic-resistant bacteria that complicate sepsis treatment, leading to higher mortality rates (8, 21). These resistant strains are not effectively killed by standard antibiotics, necessitating the use of alternative or combination therapies, which may not always be readily available or as effective. Moreover, delays in administering appropriate antibiotics, often due to the time required for microbial culture and sensitivity testing, can result in worsening patient outcomes. Early and appropriate antibiotic administration is critical for improving sepsis survival rates; however, the need for precise identification of the causative pathogen can lead to significant delays in treatment initiation (22-24). To address these issues, recent approaches have focused on the coadministration of antibiotics with other therapeutic agents that can modulate the immune response. Inflammatory regulators are a representative example of such combination therapies. These agents target various components of the inflammatory signaling pathway, including membrane receptors and intermediate substances (25-28). This evolution in treatment underscores the growing importance of targeting specific components of the inflammatory cascade, which leads us to examine innovative methods for inflammation regulation that address broader issues such as sepsis and antibiotic resistance.

Among the new methods for regulating inflammation, membrane receptor blockade can be used to address the issues related to sepsis, antibiotic resistance, and the use of conventional treatments (29). Therapeutic candidates targeting membrane receptors show the ability to rapidly control inflammation (30–32), so could be effectively applied for the treatment of acute inflammatory diseases, infectious diseases, and sepsis, for which the initial treatment is crucial. Membrane-receptor-targeting substances have immunomodulatory abilities to restore immune function rather than just inducing complete immunosuppression (33, 34). It is important to distinguish between immunosuppressants and immunomodulators. Immunosuppressants are characterized by low therapeutic indexes, meaning there are close windows between therapeutic ranges and toxic

zones. They also exhibit significant intra- and interindividual drug kinetic variations. These shortcomings are mitigated by correct drug doses, which are calculated based on ideal body weight (a calculated weight considered optimal for health) or lighter body weight (used to avoid overdosing in underweight patients). Additionally, end-organ toxicity testing and, in some cases, close monitoring of plasma drug levels (parent or metabolite peak and depth levels) are used. In contrast, immunomodulators have a broader therapeutic index, a higher safety margin, more predictable drug behavior properties and fewer intra-individual variability. In addition, although immunosuppressants typically affect host immune reactions globally, immunomodulators can act selectively on specific parts of the immune system and therefore reduce the risk of complications related to immune failure (35). In this context, the drugs investigated in the referenced papers did not induce cytotoxicity or cell death.

This review presents new therapeutic approaches for acute inflammatory and infectious diseases that increase the risk of sepsis, as well as complementary methods to enhance current sepsis treatments.

2 Methodology

For this review, we utilized PubMed and ClinicalTrials.gov to search the literature using specific terms, including 'sepsis,' 'membrane receptor blockade,' 'damage-associated molecular patterns,' 'pathogen-associated molecular patterns,' and 'inflammation.' Our focus was on English-language articles that present original research, prioritizing significant and relevant studies in the domain. Although our search was not restricted by publication date, we made a concerted effort to cite the most recent studies, generally avoiding references older than ten years.

3 Acute inflammatory and infectious diseases, and their correlation with sepsis

Acute pancreatitis (AP) often progresses to severe acute pancreatitis (SAP), where systemic inflammation frequently leads to multiple organ dysfunction syndrome (MODS) and, in some instances, sepsis, significantly increasing mortality in about 20% of these cases (36, 37). Moreover, AP results in the breakdown of bacterial translocation and intestinal integrity, and an increased risk of infection (38). Ulcerative colitis (UC) is a chronic inflammatory condition that can lead to acute kidney injury, which is associated with septic shock (39). In addition, delayed graft function (DGF), which occurs in one in five patients receiving organ transplant, provides a favorable environment for sepsis because repeated biopsies of the transplanted organs are required (40). Patients with acute inflammatory and infectious diseases and those who have undergone DGF frequently develop sepsis, a systemic inflammatory disease with a remarkably high mortality rate (41). Furthermore, infections caused by injury can lead to sepsis. Sepsis can occur due to other infections, such as viruses and fungi, but most infections are caused by bacteria. Bacteria are classified into Gram-negative and

Gram-positive types and are both capable of causing sepsis; however, they have distinct differences. The most commonly isolated bacteria in sepsis are Staphylococcus aureus (S. aureus), Streptococcus pyogenes (S. pyogenes), Klebsiella species (Klebsiella spp.), Escherichia coli (E. coli), and Pseudomonas aeruginosa (P. aeruginosa). To cause disease, pathogens must use a number of factors called virulence factors to protect themselves from host innate immune system, to cross mucous membrane barriers, to spread, and to replicate to distant organs (42, 43). To understand the factors causing bacterial pathogenicity, it is crucial to study how pathogenic agents escape host immune systems, cross mucous membrane barriers, spread, and replicate in distant organs. For example, S. aureus and S. pyogenes infect primarily the skin and lung and activate toll-like receptor (TLR) 2. Some alleles of the TLR2 pathway are associated with an increase in sensitivity and severity to sepsis caused by large Gram-positive pathogens such as S. aureus (44-47). Klebsiella spp. infects a variety of sites including the lungs, urinary tracts, blood vessels, wounds, surgery, and the brain. TLR2 recognizes the OmpA of Klebsiella pneumoniae (K. pneumoniae) and activates the NF-KB signal, while TLR4 recognizes the LPS and K1-CPS of K. pneumoniae (48-50). E. coli, commonly linked to intestinal diseases, also significantly contributes to systemic infections like sepsis. Additionally, pathogenic E. coli, especially strains like uropathogenic E. coli (UPEC), can cause severe conditions such as bacteremia and sepsis. These extraintestinal infections caused by pathogenic E. coli are the leading cause of sepsis in healthcare environments and communities (51-53). P. aeruginosa is a common pathogen found in severe burn injuries and is associated with various nosocomial infections such as pneumonia, surgical wounds, urinary tract infections, and bacterial infections. Signals induced by TLR4 and flagellin mediate the acute inflammatory response of pseudomonas, while TLR2 plays a counterregulatory role. The myeloid differentiation primary response 88 (MyD88)-dependent pathways, as well as the pathways downstream of TLR2, TLR4 and TLR5, are necessary for lung defense against P. aeruginosa (54, 55). Sepsis is an inflammatory response in the host to serious infections that pose a threat to life and accompany organ dysfunctions (56) and initiates a complex interaction of host proinflammatory and anti-inflammatory processes. Sepsis involves dynamic interaction between the host immune system and pathogens (57), the result of which depends on a delicate balance between anti-inflammatory and pro-inflammatory pathways. Septic shocks are a subgroup of sepsis with profound circulatory, cellular and metabolic abnormalities that are associated with a higher mortality risk than sepsis alone. In the current definition of sepsis, the term "abnormal regulation and host response" is not explicitly defined, but is conceptualized as a maladaptive response within the immune and non-immune systems that leads to organ failure and death (58). In the sepsis context, adherence to an initial treatment response timeframe is crucial for controlling mortality rates (59). As such, there is an increasing need for a novel approach that includes the use of inflammatory regulators as adjunctive therapy with antibiotics, not only to manage the excessive inflammation caused by sepsis but also to prevent septic shock and organ dysfunction, thereby effectively controlling and preventing the progression of sepsis within the required timeframe.

4 Signaling pathways and limitations of acute inflammatory and infectious disease treatment

When inflammatory and infectious diseases develop, a cascade of events is set in motion, including the release of DAMPs and PAMPs, leading to cellular damage (Figure 1). At the onset of these diseases, the host recognizes endogenous DAMPs or PAMPs, which interact with membrane proteins such as a series of PRRs located on the cell membrane or in the intracellular space (60, 61). Intracellular signaling pathways are activated, culminating in the production of diverse end-products with the potential to serve as DAMPs. DAMPs and PAMPs interact and directly bind to membrane receptors such as TLRs, interleukin-1 receptor (IL-1R), C-type lectin receptors (dectin 1 and dectin 2), and receptor for advanced glycation end products (RAGE) (62-64). As a result of these interactions, interferon regulatory factor (IRF) (which is responsible for the production of type I interferon (IFN)) (65), nuclear factor-kB (NF-kB), and the activator protein 1 signaling pathway (which is involved in the early activation of genes encoding inflammatory and endothelial cell surface molecules) are activated (66). DAMPs and PAMPs are recognized by various membrane receptors, initiating intracellular signaling cascades. PRRs encompass diverse membrane receptors, which are activated in a cell-specific manner. To identify which membrane receptor activation is responsible for initiating the intracellular signaling cascades in specific cells, we referred to Human Protein Atlas data. Among endothelial, epithelial, adaptive immune cells, and innate immune cells, innate immune cells, including Macrophages, Dendritic cells, and neutrophils, expressed TLR2, TLR4, and Dectin 1 most abundantly. endothelial protein C receptor (EPCR), Dectin 2, IL1R, and RAGE were predominantly expressed in epithelial cells. Notably, TLR2 was highly expressed in epithelial cells following innate immune cells, while TLR4 showed a sequence of expression in innate immune cells, followed by endothelial cells, and then epithelial cells. EPCR exhibited abundant expression in epithelial cells, followed by endothelial cells, adaptive immune cells, and innate immune cells, while Dectin 2 was prominently expressed in epithelial cells followed by innate immune cells. IL1R demonstrated significant expression in epithelial cells, followed by immune cells and endothelial cells, whereas RAGE showed prominent expression in epithelial cells followed by immune cells. We posit that such differential expression of these membrane receptors contributes to cell-specific membrane receptor activation. Immune cells can engage and interact with complex intracellular signaling systems, leading to the activation of the innate immune responses aimed at eliminating invading pathogens and maintaining cellular homeostasis (67). However, under the conditions in inflammatory and infectious diseases, specific host signaling pathways are drastically upregulated, resulting in the robust release of cytokines (such as tumor necrosis factor (TNF)-a, interleukin-1 (IL-1) and interlukin-6 (IL-6)) and alarmins proteins (such as high mobility group box 1 (HMGB1) and S100 family proteins) (68-70). These cytokines and alarmins act as potent mediators that contribute to immunosuppression and excessive systemic inflammation.



Mechanism of inflammatory and infectious diseases develop. (A) Variations in pattern recognition receptor (PRR) expression are cell-specific. The receptors named for each cell type reflect their high level of expression, as determined by the Human Protein Atlas data. (B) In the early stages of inflammatory and infectious diseases, the host recognizes DAMPs or PAMPs. DAMPs and PAMPs directly bind to membrane receptors such as TLRs, IL-1R, C-type lectin receptors, RAGE, TREM-1, and GPCR. This interaction triggers the activation of key signaling pathways, including IRF, responsible for IFN production, NF- κ B signaling pathway. Signaling pathways are significantly upregulated, resulting in the robust release of cytokines such as TNF- α , IL-1, and IL-6, as well as DAMPs like HMGB1 and S100 family proteins.

Numerous factors operating at the endogenous level are central to the regulation of the inflammatory response; therapeutic strategies aimed at controlling these factors are under development. Furthermore, drugs such as pentoxifylline, simvastatin, N-acetyl cysteine (NAC), tofacitinib, hydrocortisone, cobitolimod, and ulinastatin (UTI), which target endogenously positioned molecules such as phosphodiesterase, 3-hydroxy-3-methylglutaryl coenzyme A (HMG-CoA) reductase, glutathione synthetase, inhibitor of nuclear factor kappa B (I κ B) kinases, Janus kinase (JAK), glucocorticoid receptor, TLR9, and serine protease, are currently the focus of clinical trials (Table 1) (71–77). Given the complex interplay of signaling pathways and mediators involved in inflammation, targeted therapeutic interventions are necessary to modulate these processes effectively. The following section describes how specific drugs in clinical trials aim to manipulate these mechanisms to mitigate the inflammatory response.

4.1 Phosphodiesterase

Pentoxifylline - Pentoxifylline activates the adenosine receptor 2 by non-selective inhibition of phosphodiesterase enzymes and anti-inflammatory effects, leading to an increase in cyclic adenosine monophosphate (cAMP). The activation of protein kinase A (PKA)

TABLE 1 Inflammation modulators targeting endogenous factors and clinical trials.

Compound	Target	Condition	Status	Clinical trial ID
Pentoxifylline	Phosphodiesterase	Pancreatitis	Phase 3	NCT02487225
Simvastatin	HMG-CoA reductase	Sepsis	Phase 2	NCT00528580
N-acetylcysteine	Glutathione synthetase, IκB kinases	Systemic inflammatory response syndrome	Phase 4	NCT03589495
Tofacitinib	Janus kinase	Ulcerative colitis	Phase 3	NCT01465763
Hydrocortisone	Glucocorticoid receptor	Septic shock	Phase 4	NCT02768740
Cobitolimod	Toll-like receptor 9 (TLR9)	Ulcerative colitis	Phase 3	NCT04985968
Ulinastatin	Serine protease	Septic shock	Phase 4	NCT05895240

via cAMP suppresses the nuclear transfer of NF-κB and suppresses the transcription of inflammatory cytokines (78).

4.2 HMG-CoA reductase

Simvastatin - HMG-CoA reductase inhibitors (statins) may have a beneficial effect through various mechanisms of sepsis syndrome. Statin, inhibition of HMG-CoA reductase, not only reduces cholesterol levels but also decreases cholesterol-synthesis intermediates that affect intracellular signaling, cytokine expression, and chemokine regulation. This leads to reduced expression of adhesion molecules in leukocytes and endothelial cells. Additionally, statins have immune modulation and anti-coagulant effects, and evidence suggests they have a direct antimicrobial effect on bacteria (79).

4.3 Glutathione synthetase and IkB kinases

N-acetyl cysteine - NAC stimulates the synthesis of the main cell reduced Glutathione (GSH), regulates the redox state of the cell, and inhibits apoptosis caused by oxidative stress. In addition, NAC affects the NF-κB signal pathway, which targets IκB kinases (IKKα and IKKβ) that play a key role in this pathway. NAC regulation of IKKα and IKKβ is associated with the suppression of NF-κB activation induced by external stimuli such as TNF. Through these mechanisms, NAC improves the regulation of cell redox and interferes in inflammatory reactions and apoptosis processes and provides a cell protective effect (80).

4.4 Janus kinase

Tofacitinib - Tofacitinib regulates inflammation by inhibiting JAKs, which play an essential role in signaling inflammation mediators. Tofacitinib inhibits JAK1, JAK3, and to a lesser extent, JAK2. This inhibition blocks STAT protein phosphorylation, preventing cell gene expression modification. Consequently, tofacitinib regulates cytokine signaling pathways, reducing immunogenicity and inflammation in diseases like UC (81).

4.5 Glucocorticoid receptor

Hydrocortisone - Hydrocortisone binds to glucocorticoid receptors and has downstream effects such as inhibiting phospholipase A2, NF-κB, other inflammatory transcription factors, and promotion of anti-inflammatory genes. Glucocorticoids inhibit neutrophil apoptosis and decomposition, suppress phospholipase A2, reduce the formation of acoustic acid derivatives, suppress NF-κB and other inflammatory transcription factors, and promote anti-inflammatory genes such as interleukin-10 (IL-10) (82).

4.6 TLR9

Cobitolimod - Cobitolimod is a locally administered deoxyribonucleic acid (DNA)-based oligonucleotide that interacts with TLR9. Its clinical efficacy in patients with moderate-to-severe UC was demonstrated (76). Moreover, cobitolimod induces regulatory T cells in lymphocytes and antigen-presenting cells, leading to IL-10 production and the inhibition of TH-17 cells (83).

4.7 Serine protease

Ulinastatin - UTI, also known as a urinary trypsin inhibitor, inhibits serine proteases and plays a key role in reducing systemic inflammation and preventing cell apoptosis. The protease inhibitor inhibits the activation of NF- κ B by reducing p38-Mitogen-activated protein kinase (MAPK) phosphorylation and promoting antiinflammatory effects (84).

However, drugs targeting endogenous factors have the drawback of relatively slower response times (85–88). This limitation is particularly significant in the context of intracellular drug delivery, where the efficiency of drug transport across cell membranes is crucial. Furthermore, if exogenous molecules cannot pass through cell membranes, it becomes difficult to reach the cytosol (89). Unlike membrane receptors, which are located on the cell surface and interact directly with external drugs, intracellular proteins require drugs to undergo specific steps to cross the cell membrane and reach their target sites within the cell. In addition, in

the c-induced sepsis animal model, the expression time of MyD88 and its downstream factors appears to be different. Experimental animal studies have shown a sequence expression of MyD88 between the cecal ligation and puncture (CLP) surgery group and the sham control group. In addition, CLP surgery significantly increased the concentration of MyD88 messenger ribonucleic acid (mRNA) in tissue in 2 hours after Sepsis induction. When MyD88dependent TLR signaling pathways are activated, NF- κ B is activated and serum cytokines, including TNF- α , are significantly increased (90). Therefore, when considering the time required to express downstream factors, drugs that block the initial signal may have a faster response time than those that affect downstream factors.

5 Targeting membrane receptors for rapid regulation in acute inflammatory and infectious diseases

To improve the limitations of drug response time, it is necessary to focus on the regulation of membrane receptors that initiate the inflammatory cascade. Acute inflammatory and infectious diseases are initiated by receptors located on the cellular membrane. The intracellular signaling cascades induced by membrane receptors such as PRRs drive the transcription of the inflammatory mediators that regulate the elimination of infected cells and pathogens (91). However, abnormal activation of this inflammatory system triggers septic shock or immunodeficiency (92). For example, cells rich in specific PRRs, such as TLRs, IL1R, EPCR, and others, can recognize PAMPs and DAMPs, leading to excessive inflammation that could potentially result in septic shocks. On the contrary, if endotoxin tolerance is established, even though activation is required, PRRs can remain inactive, leading to an immunosuppressive state (93, 94). Thus, the efficient regulation of membrane receptor proteins that act as PRRs might be effective as an early treatment for acute inflammatory and infectious diseases.

TLRs are the most well-known PRRs due to their crucial role in the host defense system and their involvement in various pathological processes such as sepsis (95). The activation of TLRs enables pathogen elimination through the promotion of the antibacterial activity of immune cells and the maturation of antigen-presenting cells, which involves the development of adaptive immunity (96). However, dysregulation of TLRs can lead to severe disease such as sepsis, AP, acute UC, and acute lung inflammation (27, 97, 98). TLR4 is a signaltransducing component of the lipopolysaccharide (LPS) receptor complex, which includes Myeloid differentiation factor 2 (MD-2) and cluster of differentiation 14 (CD14). Furthermore, the progression of LPS-triggered proinflammatory reactions relies solely on the endocytosis rate of TLR4 and its trafficking via the endolysosomal compartment (99). Therefore, TLR4 is considered a key therapeutic target for treating inflammatory and infectious diseases (100, 101), potentially extending to the treatment of sepsis (102). TLR2 is expressed on diverse cell surfaces, including immune cells such as macrophages, dendritic cells, and lymphocytes. The factors recognized by TLR2 include PAMPs and DAMPs (103). TLR2 is associated with the development of infectious and inflammatory diseases. TLR2 promotes the inflammatory response via activation of the MyD88/ NF- κ B signaling pathway (104). Furthermore, TLR2 is a therapeutic target for AP treatment (105). TLR9 was originally discovered as a sensor of bacterial DNA, which is rich in unmethylated CpG dinucleotides. However, it can also recognize the DNA released from damaged cells as a DAMPs, triggering sterile inflammation (106).

IL-1R orchestrates the signaling initiated by important IL-1 pathways involved in various pathogenic processes. Typically, IL-1R promotes the transcription of inflammatory cytokines via the canonical NF-kB pathway. However, in certain specialized cell types such as neurons, it induces inflammation independent of the NF-kB pathway (107). In addition to these membrane PRRs, EPCR negatively regulates inflammatory responses and coagulopathy in sepsis (108). Moreover, the µ-opioid receptor (MOR), which is responsible for opioid recognition, is implicated in exacerbating the immunopathology of bacterial infections (109). The triggering receptor expressed on myeloid cells-1 (TREM-1), stimulated by specific ligands such as DAMPs and PAMPs during infections or tissue damage, activates the DAP12associated Syk signaling pathway. This activation leads to NF-KB translocation into the nucleus, which in turn upregulates the production of pro-inflammatory cytokines and chemokines. This cascade is pivotal in amplifying inflammatory responses, particularly in sepsis, where such dysregulation can exacerbate the condition (110). G-protein coupled receptors (GPCRs), a large and diverse family of transmembrane receptors, play a critical role in a wide range of physiological processes, including immune responses. These receptors are activated by various ligands, leading to conformational changes that facilitate the interaction with and activation of G-proteins. This interaction results in the dissociation of $G\alpha$ and $G\beta\gamma$ subunits, which then modulate downstream signaling pathways, influencing cellular responses. Dysregulation of GPCR signaling has been implicated in the pathogenesis of numerous diseases, including sepsis, highlighting their potential as therapeutic targets. For example, The kinin B1 receptor is involved in sepsis-induced vascular hyperpermeability, demonstrating the nuanced roles GPCRs play in immune responses and their potential as therapeutic targets (111, 112). Also, the C5a-C5a receptor (C5aR) axis plays a critical role in inflammatory responses, especially in the context of sepsis. Upon activation by pathogens, the complement system generates C5a, a potent inflammatory peptide, which interacts with its receptor to attract immune cells like neutrophils and macrophages, leading to oxidative bursts and release of pro-inflammatory cytokines. This cascade contributes to vasodilation, tissue damage, and multiple organ failure (MOF) in acute inflammation (1, 113).

These membrane receptors are sufficient targets for drugs administered at the exogenous level and offer the potential for rapid regulatory effects owing to their high accessibility.

6 Membrane receptor blockade as treatment for acute inflammatory and infectious diseases

In the past, exposure to acute inflammation-related diseases primarily led to the administration of antibiotics to prevent initial

infections (114). Recognizing the limitations, such as antibiotic resistance and the emergence of 'super bugs,' ongoing research is exploring the potential of adjunct therapies. Despite some partial effects of using biological factors in adjunct therapies, a substantial increase in survival rates was not achieved (115).. Therefore, since significant effects were not observed with biological factors alone, various experimental studies are investigating counteractive drugs targeting PAMPs and DAMPs, which are being explored for their potential to enhance the efficacy of antibiotics in treating inflammatory conditions and possibly sepsis (116-118). However, these approaches are still under investigation and have not vet been established in medical practice. Although these findings are preliminary and require further validation, the observed beneficial effects of co-administering immunomodulators with antibiotics in treating sepsis and other inflammatory infectious diseases suggest that continuing research in this direction could be promising. For example, C10-LRR shows promise as a potential new treatment option for managing overactive pro-inflammatory cytokine release by macrophages and could be effectively used to treat severe inflammatory conditions such as sepsis when combined with other therapies, including antibiotics or anti-TNFa antibodies (119). siTACE (TNF- α converting enzyme) reduces proinflammatory cytokines and the presence of inflammatory macrophages. TKPR-9R peptides, targeted at macrophages, enhance the cell-permeability of siRNA both in vitro and in vivo. When used together in combination therapy, siTACE/TKPR-9R complexes and antibiotics deliver simultaneous anti-inflammatory and antibacterial benefits (120). The use of a β -lactam and a macrolide together was linked to lower mortality rates in patients suffering from pneumococcal Community-acquired pneumonia (CAP) and those experiencing intense systemic inflammation. In cases where both conditions were present, the combination of β lactam and macrolide demonstrated a protective effect against mortality in the multivariate analysis (121). Such approaches have shown potential in enhancing treatment efficacy, which supports the need for well-designed, large-scale clinical trials to confirm these benefits and establish safe and effective treatment protocols (122-125).

Acute inflammatory and infectious diseases produce a notable shift in the patterns of inflammatory mediators with disease progression, characterized by a downregulation of the expression of proinflammatory cytokines. Concurrently, DAMPs, including HMGB1, S100 family, and heat shock proteins, are upregulated (126–128). Notably, the peak TNF- α concentration occurs in the blood approximately 2 hours after endotoxin injection, whereas the concentrations of HMGB1 and S100 family protein peak around 12–18 hours after endotoxin injection (129). Moreover, in experimental studies, after TNF- α reached its peak concentration in the bloodstream, the administration of specific antibodies to inhibit HMGB1 resulted in decreased mortality rates (130).

The inhibition of the products of the inflammatory signaling pathway is effective for disease treatment. The released products, such as HMGB1, interact extracellularly with cell surface receptors such as TLRs, triggering signal transduction, which activates dendritic cells. This activation prompts the secretion of TNF- α and other proinflammatory cytokines, thereby amplifying the

inflammatory response (131). Furthermore, these released products bind to the RAGE on the endothelial cell surface and induce the expression of ICAM-1 and VCAM-1, facilitating leukocyte extravasation and promoting tissue damage (132, 133). Consequently, the release of inflammatory products due to damaged endothelial or epithelial cells, along with sustained inflammation mediated by monocytes and dendritic cells, plays a critical role in the progression of sepsis toward MOF.

Taken together, these findings highlight the potential value of targeting DAMPs regulation to attenuate the detrimental effects of inflammatory responses during the later stages of acute inflammatory and infectious diseases. This suggests that in the treatment of diseases where the regulation of early inflammation is crucial, targeting the upstream inflammatory signaling pathway, rather than solely eliminating the products of the inflammatory response, may provide high-value drug candidates that overcome the challenges associated with antibiotics.

Until recently, the majority of acute inflammatory and infection diseases and DGF research had primarily centered around blocking the initial hyperinflammatory phase of conditions, which is mediated by cytokines (134–136). The following sections introduce exogenous blockers that are the most advanced membrane receptor blockers, working as early stage inflammation modulators developed for the treatment of various acute inflammatory and infectious diseases.

7 Membrane receptor blockade as an inflammatory regulator

7.1 TLR4

Resatorvid - A widely recognized inhibitor of TLR4 is TAK-242, which is also known as resatorvid or ethyl-(6R)-[N-(2-chloro-4-fluorophenyl) sulfamoyl], developed by Takeda Pharmaceutical Company in Osaka, Japan (137). TAK-242 interacts with cysteine 747 within the intracellular domain of TLR4. This interaction leads to the inhibition of both MyD88-dependent and MyD88-independent pathways, which are activated by LPS (138).

Eritoran - Eritoran acts as an antagonist of MD2-TLR4 and effectively blocks LPS-induced hyperinflammation in *in vitro* and *in vivo* experimental animal models (139). Furthermore, it reduces phosphorylated p38-MAPK and NF-κB expression levels (140).

NI-0101 - NI-0101 forms an immune complex with the citrullinated proteins that bind to TLR4, effectively preventing cytokine release, both *in vitro* and *in vivo*. Furthermore, it successfully suppresses the anticipated increase in C-reactive protein (CRP) levels following LPS administration *in vivo* (141, 142).

7.2 TLR2

OPN-305 - OPN-305 blocks TLR2/1- and TLR2/6-mediated signaling, thereby reducing the production of proinflammatory cytokines through TLR2, inducing high TLR2 occupancy, and reducing IL-6 secretion (143). In addition, its role involves the

inhibition of TLR2-mediated ischemia-reperfusion injury, a pivotal factor in the pathogenesis of DGF and its subsequent complications (144).

7.3 IL-1R

Anakinra - Anakinra, an interleukin receptor antagonist, effectively inhibits the biological function of IL-1 by competitively inhibiting its binding to IL-1R (145). It is effective in diseases with severe activation of the NLRP3 inflammasome complex, as evidenced by rapid improvements in clinical symptoms and inflammatory markers, and substantial reductions in major organ-related symptoms (146).

7.4 EPCR

rhAPC - In addition to its role as an anticoagulant that downregulates thrombin generation, rhAPC binds to EPCR, activating protease-activated receptor 1 in endothelial cells, thereby inducing a multifactorial cytoprotective signaling pathway (147). Furthermore, rhAPC administration decreases the concentration of proinflammatory cytokines in the bloodstream (148).

7.5 MOR

Naldemedine - Naldemedine interacts with MOR and dosedependently inhibits cAMP level reductions and β -arrestin recruitment increases. Moreover, naldemedine administration substantially attenuates the upregulation of genes related to immune checkpoints (149).

As mentioned in this session, compounds capable of interacting with membrane proteins under various exogenous conditions are being investigated as therapeutic agents for the treatment of acute inflammatory and infectious diseases. Furthermore, their efficacy has been substantiated in clinical trials, as described in Table 2.

Examining the outcomes of completed clinical trials, phase 3 trials revealed that eritoran was not effective in reducing mortality rates in severe sepsis patients, a finding that contradicts earlier phase 1 results and preclinical studies. TAK-242, another investigational drug, did not decrease cytokine levels or enhance organ function in patients with severe sepsis-induced shock or respiratory failure, although some mortality benefits were noted in a specific subgroup. Post hoc analysis of anakinra suggested a possible 30% reduction in 28-day mortality among patients with liver issues and disseminated intravascular coagulation (DIC), despite the main trial showing no overall survival benefits. Additionally, the trial involving recombinant human activated protein C indicated no advantages in treating severe septic shock. Finally, the PAMORA-RAP trial is evaluating the efficacy of naldemedine in preventing recurrent AP in a randomized controlled setup (150-154). Although initial clinical phases have shown success, there have been numerous subsequent failures. However, this does not mean that immunomodulators lack potential in treating sepsis. One possible reason for clinical failures could be that blocking the TLR4 receptor does not completely halt inflammation, as other PRRs can still recognize various PAMPs or DAMPs and activate inflammatory gene transcription independently of TLR4 signaling. Therefore, it could be beneficial to explore research modifying membrane receptor blockades to interact with a variety of PRRs, which might enable more rapid and effective control of inflammation at the start of the signaling pathway.

Although clinical trials have not yet been conducted, efforts to develop drugs targeting TREM-1 for the treatment of sepsis are underway. Peptide-based therapies such as LP17, M3, and N1 bind to TREM-1, reducing inflammatory responses and demonstrating therapeutic effects in various sepsis models induced through different methods (155–157). Various types of GPCRs are also under extensive research as targets for sepsis treatment. BI113823 is an orally active nonpeptide kinin B1 receptor antagonist of small molecule and is a potent anti-inflammatory agent with a good cardiovascular profile. BI113823 reduces systemic and tissue inflammation, prevents hemodynamic derangement, reduces multiorgan injury, and improves overall survival in the rat model of

TABLE 2 Membrane receptor blockades act as inflammation modulators and clinical trials.

Compound	Target	Condition	Status	Clinical trial ID
Eritoran	Toll-like receptor 4 (TLR4)	Sepsis	Phase 3	NCT00334828
Resatorvid		Sepsis	Phase 3	NCT00633477
NI-0101		Acute respiratory distress syndrome	Phase 2/3	NCT04401475
OPN-305	Toll-like receptor 2 (TLR2)	Delayed graft function	Phase 2	NCT01794663
Anakinra	IL-1 receptor (IL-1R)	Sepsis	Phase 2	NCT04990232
rhAPC	Endothelial protein C receptor (EPCR)	Sepsis	Phase 3	NCT00625209
Naldemedine	μ-Opioid receptor (MOR)	Pancreatitis	Phase 3	NCT04966559

polymicrobial sepsis caused by CLP (158). The function of C5aR in sepsis was also explored using a C5aR antagonist, C5aRa. C5aRa, a cyclic peptide, competes with C5a to bind to C5aR. In cases of sepsis, C5aRa treatment hindered the chemotactic responses of neutrophils to C5a, thereby preventing C5a/C5aR-induced impairment of innate immunity, resulting in enhanced survival during a 9-day investigation. These findings additionally underscore C5aR as a prospective therapeutic target in sepsis (159, 160).

Many sepsis patients may experience a phase of excessive inflammation for a relatively brief duration. Hyperinflammation and the onset of MOF manifest over time, with early MOFs posing a risk of fatality (161). Therefore, it is crucial to promptly diagnose and initiate appropriate treatment during the early stages of sepsis. Membrane receptor blockades serve as early inflammation modulators designed for various acute inflammatory and infectious diseases. Consequently, they can swiftly counter the initial inflammatory responses in sepsis, potentially diminishing inflammation and enhancing treatment outcomes and survival rates.

8 Membrane receptor blockade: novel approach to address the challenge of antibiotic resistance in sepsis treatment

Conditions such as organ transplantation, UC, and AP exhibit high mortality rates due to sepsis. A swift initial intervention plays a crucial role in mitigating these mortality rates. In the context of sepsis, diagnosis and suitable management within the initial hours after onset result in notably improvements in patient outcomes (10). In the event of the onset of sepsis, antibiotic treatment must be quickly initiated, ideally within the first hour (162). Although conducting relevant culture tests before antibiotic administration is important, such diagnostic procedures should not delay antibiotic administration. The administration of an antibiotic combination effective against the presumed causative bacteria should be completed within a 3-hour timeframe (163). However, the use of antibiotic-based combination therapy is advised for a duration not exceeding 3-5 days, and transitioning to monotherapy based on the susceptibility results should be promptly considered. Generally, the antibiotic treatment period spans approximately 7-10 days (164). The constraint on antibiotic dosing regimens is that as the number of antimicrobial-resistant bacteria acquired in both community and healthcare settings increases worldwide, effective antibacterial therapy becomes increasingly difficult, especially regarding empirical antimicrobial selection (165).

In the management of acute infections and immune-related diseases, initial treatment often relies on the essential and unavoidable administration of excessive doses of antibiotics, which, over time, can create a formidable barrier known as antibiotic resistance (24, 166). Inflammatory regulators used for the treatment of acute inflammation and infectious diseases, which are applied to solve the problem of antibiotic resistance, can be categorized into drugs that regulate the entire signaling pathway and those that regulate factors at the intracellular level (167). However, in diseases such as those mentioned above where timing is crucial, medications focused on rapidly blocking signal transduction pathways may be more efficient.

Numerous obstacles exist in relying solely on antibiotic-based treatment for sepsis, which is common practice (168). Consequently, several researchers have explored the use of neutralizing-antibody-based inflammation control as adjunct therapy. However, obtaining meaningful results has been challenging owing to their diminishing efficacy over time. Furthermore, the compounds targeting endogenous factors introduced in Table 1 have a slower response time at the *in vitro* level, making it difficult to alleviate the initial symptoms within the recommended 3-hour timeframe as stipulated by the 3-Hour Surviving Sepsis Campaign Guideline (59).

Exogenous inflammatory regulators, as described in Table 2, have relatively prompt inflammation control capabilities in vitro. In particular, regulatory substances related to TLRs, such as resatorvid and eritoran, can control inflammation within 1-2 hours of treatment (169-171). Sepsis is a frequent complication arising from combat injuries and trauma and is characterized by lifethreatening organ dysfunction resulting from a dysregulated host response. Although the pathophysiology remains unclear, immunosuppression is currently acknowledged as a major cause of the high mortality rate associated with sepsis (172). From this perspective, inflammatory regulators must not permanently inhibit the inflammatory mechanism but revert to a normal state once the initial disease is alleviated. The restoration of cellular mechanisms to a normal state is evident 24 h after treatment with exogenous inflammation regulators. Additionally, the expression of TLR4, a crucial component in the recognition of secondary infectious agents following sepsis, returns to its preinhibitory state. Furthermore, the translocation of HMGB1, a significant factor in exacerbating lesions, can be inhibited at a rate ranging from 60% to 70% (30, 34, 170, 173-175). In other words, since these therapeutic candidates are not irreversible, they have the ability to modulate immunity rather than suppress it, thereby facilitating the restoration of immunity to its natural state.

In summary, patients with various acute inflammatory and infectious diseases are highly susceptible to sepsis, typically requiring the initial use of antibiotics. The frequent reliance on broad-spectrum antibiotics in high doses often leads to antibiotic resistance, a significant challenge in treatment. To address this, adjunctive therapies like immunomodulatory agents are explored, designed to be used alongside antibiotics to enhance treatment effectiveness. These agents help manage inflammation, providing valuable time to accurately identify the appropriate antibiotic treatment, thereby reducing tissue damage and mitigating antibiotic resistance. Importantly, while antibiotics target the pathogens, immunomodulatory agents control the excessive inflammatory responses that could lead to further complications. This synergistic approach allows for more precise and conservative use of antibiotics, avoiding high doses or broad-spectrum use, and protects the body from collateral damage caused by an overactive immune response. Furthermore, it is important to avoid inducing immune suppression with inflammatory regulators to prevent secondary infections and ensure the natural restoration of immune function. The novel immunomodulatory agents discussed in this paper, which target membrane receptors, present a valuable adjunct approach that enhances the efficacy of antibiotic therapy. These adjunctive therapies are not intended to replace antibiotics but to be used concurrently, enhancing their efficacy while managing inflammation effectively. It is crucial to note that in clinical trials exploring the co-administration of biological factors and antibiotics in the treatment of sepsis, antibiotics were never discontinued (176–178). Instead, these therapies are designed to complement each other, with the goal of improving outcomes by mitigating the effects of inflammation and reducing the risk of resistance development. These immunomodulatory agents, which target membrane receptors, modulate immune responses, decrease inflammation, and mitigate organ harm more rapidly than agents targeting exogenous factors. Thus, potentially reducing the need for prolonged antibiotic usage and aiding in patient recovery.

9 Concluding remarks

This review emphasizes the importance of early intervention in acute inflammatory and infectious diseases, particularly those associated with the risk of sepsis. AP, UC, and DGF can escalate to sepsis, resulting in high mortality rates. Timely treatment of sepsis, which is characterized by a complex interplay of proinflammatory and anti-inflammatory processes, is crucial to



FIGURE 2

Efficacy of membrane receptor blockade. Membrane receptors blockade holds promise for enhancing patient outcomes, especially in patients prone to sepsis, all while mitigating the risks associated with antibiotic resistance and immunosuppression.

effectively control mortality rates. However, antibiotic resistance is a major problem with antibiotic-based therapy, which the most common sepsis treatment method. However, the efficiency of cytokine-based therapy, which was developed to address this problem, is low.

To improve these treatment methods, new methods of controlling inflammation must be developed. Current therapeutic strategies targeting endogenous molecules show promise but suffer from slow response times. To address this limitation, a novel approach focusing on membrane receptors, which are initiators of the inflammatory cascade, was proposed. Membrane receptors such as TLRs, IL-1R, EPCR, and MOR play crucial roles in the inflammatory response and provide opportunities for rapid regulation.

Various membrane receptor blockade targeting receptors such as TLR4, TLR2, IL-1R, EPCR, and MOR have shown efficacy in preclinical and clinical studies. Despite the lack of evident effects in clinical trials of TLR inhibitors for the treatment of sepsis, these failures have provided crucial information for future research. Specifically, TLR4 inhibitors such as Eritoran and Resatorvid have been safely permitted in clinical trials, laying the groundwork for subsequent studies. The failures in clinical trials have been attributed to various factors, including patient diversity, inflammation severity, and differences in infecting pathogens. Therefore, there is a suggestion for the need to explore diverse treatments and combinations in future clinical studies. And Membrane receptor blockers act as early stage inflammation modulators and produce a faster response than traditional therapies. Furthermore, the reviewed membrane-targeting substances demonstrate immunomodulatory capabilities without inducing complete immunosuppression, allowing the restoration of natural immune functioning.

In summary, the exploration of membrane receptor blockade as an adjunct treatment for acute inflammatory and infectious diseases is a promising research avenue. When combined with antibiotics, these novel approaches have the potential to improve patient outcomes, especially in patients prone to sepsis, while minimizing the risks associated with antibiotic resistance and immunosuppression (Figure 2).

Author contributions

S-JM: Conceptualization, Investigation, Writing – original draft. EC: Conceptualization, Investigation, Writing – original draft. HK: Conceptualization, Investigation, Writing – original draft. WG: Investigation, Software, Writing – original draft. C-SY: Writing – original draft, Writing – review & editing.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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