Sialic Acids in the Immune Response during Sepsis

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Sialic acid-binding immunoglobulin-type lectins (Siglecs) are a group of cell surface transmembrane receptors expressed on immune cells, and regulate immune balance in inflammatory diseases. Sepsis is a life-threatened inflammatory syndrome induced by infection, and the pathogenesis of sepsis includes immune dysregulation, inflammation, and coagulation disorder. Here, we reviewed the various roles acted by Siglecs family in the pathogenesis of sepsis. Siglec-1, Siglec-5, and Siglec-14 play bidirectional roles through modulation of inflammation and immunity. Siglec-2 regulates the immune balance during infection by modulating B cell and T cell response. Siglec-9 helps endocytosis of toll-like receptor 4, regulates macrophages polarization, and inhibits the function of neutrophils during infection. Siglec-10 inhibits danger-associated molecular patterns induced inflammation, helps the initiation of antigen response by T cells, and decreases B-1a cell population to weaken inflammation. Regulating the Siglecs function in the different stages of sepsis holds great potential in the therapy of sepsis.

Keywords: sialic acid-binding immunoglobulin-type lectins, sepsis, infection, sialic acid, inflammation

INTRODUCTION

Sialic acid-binding immunoglobulin-type lectins (Siglecs), a broad range of cell surface transmembrane receptors that contain 2–17 extracellular Ig domains, are found on the surface of both innate and adaptive immune cells (1). Through recognition of their glycan ligands, they are involved in the regulation of immune balance in sepsis, autoimmune diseases, and cancer (2–5). Siglecs can be divided into two groups. Group 1 consists of sialoadhesin, CD22, Siglec-4, and Siglec-15, which are conserved across mammals. Group 2 consists of CD33-related Siglecs that vary from species to species, and humans express a much larger variety of CD33-related Siglecs than rodents due to the loss of Siglecs genes in rodents (6). The extracellular Ig domains includes an amino-terminal

Abbreviations: Arg-1, arginase1; BCR, B cell receptor; CCR7, chemokine receptor 7; CLP, cecal ligation and puncture; COPD, chronic obstructive pulmonary disease; DAMPs, danger-associated molecular patterns; dLNs, draining lymph nodes; ERK, extracellular signal-regulated kinase; GBS, group B Streptococcus; HIV, human immunodeficiency virus; HMGB1, high-mobility group box 1; Hsp, heat shock protein; IFN, interferon; IL, interleukin; IRF, interferon regulatory factor; ITIMs, immunoreceptor tyrosine-based inhibitory motifs; LCMV, lymphocytic choriomeningitis virus; LPS, lipopolysaccharides; MAPK, mitogen-activated protein kinase; MAPK/ERK, mitogen-activated protein kinase/extracellular signal-regulated kinase; MHC, major histocompatibility complex; MLV, murine leukemia viruses; MODS, multiple organ dysfunction syndrome; MyD, myeloid differentiation factor; NF-κB, nuclear factor-kappa B; PD-L1, programmed death-ligand 1; PET, positron emission tomography; PSGL1, P-selectin glycoprotein ligand 1; ROS, reactive oxygen species; Siglecs, sialic acid-binding immunoglobulin-type lectins; SOCS3, suppressor of cytokine signaling; STAT, signal transducer and activator of transcription; TGF-β, transforming growth factor-β; TLR, toll-like receptor; TRAM, TRIF-related adaptor molecule; IFN-β, interferon-β; TRIF, TRIF-domain-containing adapter-inducing interferon-β; VAP-1, vascular adhesion protein-1; VCCs, virus-containing compartments; WNV, West Nile virus.
V-set domain which contains the sialic acid-binding site, while the cytoplasmic domains have immunoreceptor tyrosine-based inhibitory motifs (ITIMs) which plays a key role in modulating function of immune cell via tyrosine phosphatases recruitment, such as the SH2 domain-containing protein tyrosine phosphatases SHP-1 and SHP-2 (2).

Sepsis is defined as life-threatening organ dysfunction induced by an uncontrolled host response to invading pathogens, which kills as many as one in four similar to acute myocardial infarction, stroke, or multiple injury, and is the leading cause of mortality of patients in ICU worldwide and (7–13). Some patients die rapidly from septic shock accompanied multiple organs dysfunction caused by the cytokines storm, while some patients survive the initial phase of sepsis but die from the secondary infection caused by immunosuppression state in the late time of sepsis (14–16). Thus, it can be seen that the dysregulation of immune function by immune cells contribute to the high mortality of sepsis. As important receptors in immune cells, Siglecs are involved in the pathogenesis and therapy of sepsis. Here, we present the recent developments at our understanding of the roles of some sepsis-related Siglecs family members (Siglec-1, Siglec-2, Siglec-5, Siglec-7, Siglec-9, Siglec-10, and Siglec-14) in immune regulation, and we also summarize current efforts to develop therapeutics targeting Siglecs for the treatment of sepsis (Table 1).

### SIGLEC-1

Siglec-1, also named sialoadhesin (CD169), a myeloid-cell receptor expressed on macrophages, recognizes viral membrane gangliosides and regulates the immune response of infection especially human immunodeficiency virus (HIV) infection (17, 18). On the one hand, Siglec-1 controls the severe immunopathology in infection. A recent study showed that the deletion of Siglec-1 in the plasmodium-infected mice increased the inflammation and vascular leakage, which increased the possibility of multiple organ dysfunction syndrome (19). Another recent study in the lymphocytic choriomeningitis virus infection, the interferon (IFN)-I production decreased, and mice exhibited severe immunopathology and died quickly after the deletion of Siglec-1 (20). Siglec-1 also promotes transforming growth factor-β (TGF-β) production in the in vitro macrophages, which suppresses the innate immunity and induces the endotoxin tolerance (21). On the other hand, Siglec-1 also promotes spread

### Table 1: Sialic acid-binding immunoglobulin-type lectins (Siglecs) related researches in sepsis.

| Siglecs | Research methods | Mechanisms underlying | Study type | Results | Reference |
|---------|------------------|-----------------------|------------|---------|-----------|
| Siglec-1 | Deletion of Siglec-1 | Inflammation↑ vascular leakage↑ | Plasmodium infected mice | MODS↑ Death↑ | Gupta et al. (19) |
| Siglec-1 | Deletion of Siglec-1 | IFN-I production↑ PD-L1↑; CD8+ T cell exhaustion ↓ | Mice with LCMV infection | Immunopathology↑ | Shaabani et al. (20) |
| Siglec-1 | by LPS-induced tolerant | TGF-β↑ | RAW264.7 macrophages | Innate immunity (endotoxin tolerant) | Wu et al. (21) |
|          |                  | Virus laden macrophages contacts to trans-infect B-1 cells and migrates into lymph nodes | MLV or HIV-1 infected mice | Spread of infection | Sewald et al. (22) |
| Siglec-2 | Soluble CD22 | Elevated in serum | Gram-negative bacterial septic patients | Correlated with severity of sepsis | Jiang et al. (26) |
|          | Deletion of Siglec-2 | Chemokine↑ | WNV infected mice | Accelerated infection | Ma et al. (28) |
| Siglec-5 | Human THP-1 cells, monocyte, neutrophils | Activated p38, MAPK, and Akt signaling pathways | GBS infection | Paired receptor to regulate immune response | All et al. (32) |
| Siglec-14 | Human tissue, THP-1 cells | Bind to Hsp70 | LPS stimulation | Paired receptor to regulate immune response | Fong et al. (35) |
| Siglec-7 | Ba/F3 cells | Bind to SOCS3 | Ba/F3 cells | Regulate cytokine-induced proliferation | Orr et al. (42) |
| Siglec-9 | BMDMs, 293T cells, TLR4-HEK cells | MyD88-specific manner | LPS stimulation | Negative regulation of TLR4 responses | Boyd et al. (47) |
|          | Siglec-E knockout mice | NF-κB and MAPK p38 signal pathway | Infected with Escherichia coli | Provide immune balance | Wu et al. (48) |
|          | RAW264.7 macrophages | MAPK/MEK/ERK pathways | IL-4 stimulation | Arg-1↑ | Higuchi et al. (52) |
|          | Deletion of Siglec-E | Akt activation | Aerosol of LPS | Neutrophil recruitment to lung↑; ROS↑ | McMillan et al. (53) |
|          | Human PBMC-derived macrophages | HS9-Fab03 bind to Siglec-9 antigen | LPS stimulation | Pro-inflammatory cytokines↑ | Chu et al. (57) |
| Siglec-10 | BMDMs, CHO cells, THP-1 cells | MyD88 and p38 MAPK signaling pathways | Campylobacter jejuni infection | Anti-inflammatory↑ | Stephenson et al. (59) |
|          | Deletion of Siglec-G | Bind with CD24 and DAMPs | AAP-induced liver injury in mice | Negative regulation of inflammation | Chen et al. (62) |
|          | Deletion of Siglec-G | Binds to the BCR of B-1a cells | Siglec-G−/− mice | Apoptosis↑ | Jellusova et al. (68) |
of infection and helps virus escape from neutralization. A recent study from murine leukemia virus or HIV-1-infected mice indicated that, after the capture of viruses by Siglec-1 on macrophages, the virus laden macrophages contacted to trans-infect B-1 cells, which subsequently migrated into the lymph node and contributed to the spread of infection (22). In an in vitro study, HIV-1 particles were inadequately accessed by anti-gp120 broadly neutralizing antibodies and thus were less susceptible to neutralization in deep virus-containing compartments in the help of Siglec-1 (23). It can been seen that Siglec-1 controls the severe immunopathology through increasing the production of IFN-1 and TGF-β, on the other side, it also promotes spread of virus infection at the same time. Therefore, Siglec-1 plays a bidirectional role in infection and acts as a potential target in the treatment of sepsis.

SIGLEC-2

Siglec-2 (CD22) is a cell surface receptor expressed mostly on B cells, and regulates B cells proliferation, survival, signaling, and antibody production (24). A previous study using Siglec-2−/− mice confirmed that the absence of Siglec-2 did not interfere with the severity of arthritis, survival, bacterial clearance, and the inflammatory response during *Staphylococcus aureus* infection (25). However, with the gradual progress of Siglecs research in sepsis, it seems that Siglec-2 is closely associated with the development of sepsis. First, serum soluble CD22, a fragment of Siglec-2, was significantly elevated in patients with gram-negative bacterial sepsis and was correlated with the severity of sepsis (26). Second, in septic patients, miR-19a in B cells was up-regulated, and it comprised a feedback loop with Siglec-2 for B cell response. That provided a potential therapeutic target to restore the immune homeostasis in sepsis (27). What is more, a recent Siglec-2−/− mice study confirmed that Siglec-2 helped to control West Nile virus infection through CD8 T cells response, promoted lymphocyte migration into the draining lymph nodes, and affected chemotaxis via controlling chemokine production (28). Siglec-2 specific immunotoxins have been used in clinical studies for hairy cell leukemia and autoimmune diseases (29, 30), however, studies on the sepsis is still lacking. To sum up, Siglec-2 is involved in the immune balance of sepsis through regulating B cell response and controlling chemokine production, and Siglec-2 targeting therapy holds a great potential for the treatment of sepsis.

SIGLEC-5 AND SIGLEC-14

Siglec-5 and Siglec-14, a paired receptor system in the Siglecs family expressed on monocytes and neutrophils, share almost identical ligand-binding domains but have opposing effects in the regulation of host immunity. This idea was discovered in the research of group B *Streptococcus* (GBS) infection (Figure 1). Early study showed that GBS β protein bound to Siglec-5 and promoted bacterial survival through impairing human leukocyte phagocytosis, oxidative burst, and extracellular trap production (31). Five years later, Ali et al. discovered that Siglec-14 also involved in the GBS infection as a paired receptor with Siglec-5. β protein of GBS bound to both Siglec-5 and Siglec-14 on neutrophils, and Siglec-14 counteracted pathogen-induced host immune suppression by activating p38 mitogen-activated protein kinase (MAPK) and Akt signaling pathways (32). As Siglec-14 is not expressed by all people, homozygous Siglec-14-null neutrophils are more susceptible to GBS immune subversion (32). This idea was also confirmed in
the research of chronic obstructive pulmonary disease (COPD). Loss of Siglec-14 reduces the risk of COPD exacerbation (33), and inhaled corticosteroids could exert two opposite effects depending on the patients’ phenotypes of Siglec-5 and Siglec-14 (34). What’s more, a recently study found that heat shock protein (Hsp) 70, a danger-associated molecular pattern (DAMP), could bind to both Siglec-5 and Siglec-14 and play a two-way role in the immune modulation (35). This may explain the contradictory conclusions on the function of extracellular Hsp70 in inflammation (36–38). In brief, the bidirectional action played by Siglec-5 and Siglec-14 involved both neutrophils function and Hsp70 modulation in infection.

**SIGLEC-7**

Siglec-7 (CD328) is constitutively expressed on natural killer (NK) cells, mast cells, basophils, and platelets. It has been proven as a very important regulator of the immune response through inhibiting NK cells activation, regulating apoptosis and death, and affecting IgE-mediated mast cells and basophils activation (39–41). In sepsis, Siglec-7 acts as a target of suppressor of cytokine signaling 3 (SOCS3) and amplify inflammation through activating monocytes (42). SOCS3 in the spleen, lung, and peritoneal leukocytes is up-regulated during sepsis (43). SOCS3 binds the phosphorylated ITIMs carried by Siglec-7 and blocks Siglec-7 mediated inhibition of cytokine-induced proliferation. This also contributes to the exaggerated inflammatory response induced by pro-inflammatory cytokines during infection (42).

Some pathogens escape host immune response through binding to Siglec-7 with sialic acids expressed on their surface. Varchetta et al. demonstrated that Siglec-7 activated a monocyte-mediated inflammatory and produced high level of pro-inflammatory cytokines and chemokines through phosphorylation of the extracellular signal-regulated kinase (ERK) pathway following pathogen recognition. What’s more, Siglec-7 also participated in generating a monocyte-mediated inflammatory when encountering pathogens not expressing sialylated glycans. This phenomenon may provide an alternative mechanism that Siglec-7 involved in sepsis (44).

**SIGLEC-9**

Siglec-9, Siglec-E in murine, the major CD33-related Siglec, is mainly expressed on neutrophils, monocytes, macrophages, and dendritic cells (45), and involves in the pathogenesis of sepsis through interacting with TLR4, regulating the polarization of macrophages, and inhibiting the stimulation of neutrophils (Figure 2).

Broad and direct interaction exist between TLR4 and Siglec-E (46). Murine Siglec-E is induced by TLR4 in a myeloid differentiation factor (MyD) 88-specific manner and negatively regulates TLR4 responses following lipopolysaccharides (LPS) stimulation (47). A recent study discovered that Siglec-E participated in the Escherichia coli-induced endocytosis of TLR4, and provided an immune balance in inflammation (48). Siglec-E deficient dendritic cells failed to internalize the TLR4 and resulted in high levels of pro-inflammatory cytokines through nuclear factor-kappa B (NF-κB) and MAPK p38 signal pathway when infected with E. coli (48). Taken together, Siglec-E plays a novel role in controlling the septic response with TLR4 and helps to maintain a healthy cytokine balance following infection.

![Figure 2](image-url) | Siglec-E/9 in the immune regulation of sepsis. (A) Siglec-E negatively regulates TLR4 responses in a MyD88-specific manner following LPS stimulation. (B) Siglec-E provides immune balance in inflammation when participating in the Escherichia coli-induced endocytosis of TLR4. (C) Siglec-9 enhances IL-4-induced Arg-1 and CD200R production through MAPK/ERK pathways. TLR, toll-like receptor; LPS, lipopolysaccharides; MyD, myeloid differentiation factor; NF-κB, nuclear factor-kappa B; TRIF, TIR-domain-containing adapter-inducing interferon-β; TRAM, TRIF-related adaptor molecule; IFN-β, interferon-β; IL-4, interleukin 4; MAPK/ERK, mitogen-activated protein kinase/extracellular signal-regulated kinase; STAT, signal transducer and activator of transcription; IRF, interferon regulatory factor; Arg-1, arginase1.
Macrophages polarization plays a pivotal role in the pathogenesis of sepsis, and regulating the phenotypes of macrophages in the different stages of sepsis holds a great potential in the treatment of sepsis (49, 50). Recent studies shown that Siglec-9 enhanced induction of Arg-1 through MAPK/ERK pathways in the stimulation of interleukin 4 (IL-4) (51). Siglec-9 enhanced IL-4-induced CD200R expression and inhibited LPS-induced CCR7 in human macrophages (52). However, the detailed mechanisms under Siglec-E and macrophages polarization in sepsis need to be further elucidated.

As an important regulator expressed on neutrophils, Siglec-E function as an inhibitory receptor on the neutrophils stimulated by LPS. McMillan et al. (53) demonstrated that Siglec-E inhibited the β2-integrin-dependent neutrophil recruitment to the lung and enhanced nicotinamide adenine dinucleotide phosphate-oxidase (NADPH) oxidase activation and reactive oxygen species production via Akt activation following exposure to LPS. What is interesting, the reason of neutrophils become much easier activated after separation from whole blood also involved in Siglec-9. A recent study discovered that the abundant sialoglycoprotein on erythrocytes engaged neutrophil Siglec-9 and dampened the innate immune cell activation (54).

What’s more, a recent study using mouse and Chinese hamster ovary cells discovered a new role for Siglec-E/Siglec-9 (55). Siglec-E/Siglec-9 could specifically bind to vascular adhesion protein-1 (VAP-1), an endothelial cell molecule involved in granulocyte migration to sites of inflammation. Using 68Gallium-labeled peptide of Siglec-9 to detect VAP-1 in vasculature as an imaging tool in inflammation in positron emission tomography will give great help in the treatment of inflammatory diseases.

Recently, great progress has been made in Siglec-E targeting therapy of sepsis. Spence et al. created nanoparticles decorated with sialic acid and developed a novel strategy to control inflammation. From human monocytes and macrophages in vitro model and human ex vivo model of lung injury, they revealed that those special nanoparticles blocked the production of inflammatory mediators induced by LPS in a Siglec-E-dependent manner through enhancing the oligomerization of Siglec-E receptor on macrophages (56). Another study from human peripheral blood mononuclear cell-derived macrophages showed that a human anti-Siglec-9 Fab fragment named hS9-Fab03, specially bound to Siglec-9 antigen with high affinity and attenuated LPS-induced pro-inflammatory cytokines production (57). Those discovery confirmed that Siglec-E/Siglec-9 as a druggable anti-inflammatory therapeutic target for sepsis.

SIGLEC-10

Siglec-10, Siglec-G in murine, is broadly expressed on B cells, dendritic cells, and macrophages subsets, which is also a member of the CD33-related Siglecs family (58). It involves in the process of innate and adaptive immune response, and plays an anti-inflammatory role in sepsis through increasing IL-10 expression, interacting with CD24, inhibiting dendritic cell cross presentation, and weakening B cell signaling (Figure 3).

Siglec-10 involved in the Campylobacter jejuni infection and promoted an anti-inflammatory function through binding to C. jejuni and purified flagellum and increasing IL-10 expression by MyD88 and p38 MAPK signaling pathways (59). Siglec-G

![Figure 3](image-url)
also mediated an immune evasion pathway in RNA virus infection. Chen et al. discovered that RNA virus specifically up-regulated Siglec-G expression in macrophages by RIG-I or NF-κB-dependent mechanisms. Siglec-G recruited SHP-2 and E3 ubiquitin ligase c-Cbl to RIG-I and induced RIG-I degradation via K48-linked ubiquitination at Lys813 by c-Cbl. The increased Siglec-G led to the persistence of RNA virus infection and severe immunopathology through the suppression of IFN-I production (60).

CD24 protects the host against the exaggerated inflammatory response in sepsis (61). CD24 is a small glycosyl-phosphoinositol-anchored protein that is able to provide costimulatory signals to T cells. In sepsis, CD24 associates with DAMPs, such as high-mobility group box 1, Hsp70, and Hsp90, negatively regulates their stimulatory activity and inhibits NF-κB activation through association with Siglec-G (62). What’s more, microbial sialidase targeting Siglec-G blocks the CD24-Siglec-G pathway and exacerbates inflammation. Using sialidase inhibitors to prevent disrupting sialic acid-based pattern recognition protected mice against cecal ligation and puncture (CLP) induced sepsis, and this process depended on the CD24 and Siglec-G interaction (63, 64). The pathogenesis of sepsis involves multiple inflammatory mediators and a lot of them are regulated by the interaction of CD24 and Siglec-G. Therefore, sialidase inhibitors targeting CD24-Siglec-G interaction has a great clinical potential in the treatment of sepsis.

Siglec-G expressed on dendritic cells also contributed to the initiation of antigen response by T cells. Siglec-G inhibits cross-presenting extracellular antigens with CD8 T cells by impairing major histocompatibility complex class I-peptide complexes formation. This process involves recruiting the phosphatase SHP-1 by Siglec-G, dephosphorylating the NADPH oxidase component p47phox, and inhibiting the activation of NOX2 on phagosomes (65). Soluble CD52 released by phospholipase C bound to Siglec-10 and impaired phosphorylation of the T cell receptor associated kinases Lck and Zap70 and T cell activation, which was distinct from regular T cells (66).

Siglec-G is also broadly expressed on B cells, and plays as a negative regulator of B cell receptor (BCR)-mediated signaling in inflammation. Siglec-G binds to the BCR on the B cell surface via interaction with sialic acid ligands, and controls B cell tolerance (67). Siglec-G+/− B-1a cells display an altered BCR repertoire and a higher expression levels of the transcription factor, and show a lower level of spontaneous apoptosis and a prolonged life span (68). Hence, Siglec-G negatively regulates the inflammation through decreasing B-1a cell population, weakening B-1 cell signaling, and shifting the immunoglobulin repertoire secreted by B-1 cells.

**SUMMARY**

Immune disorder contributes to the different stages of sepsis, while the Siglecs play significant roles in the immune regulation. There are heavy conjugations between Siglecs and the pathogenesis and therapy of sepsis. Siglec-1, Siglec-5, and Siglec-14 play bidirectional roles in sepsis through modulation of inflammation and immunity. Siglec-2 involves in B cell and T cell response during infection and regulates the immune balance. Siglec-9 helps endocytosis of TLR4, regulates macrophages polarization, and inhibits the function of neutrophils during infection. Siglec-10 inhibits DAMPs-induced inflammation, helps the initiation of antigen response by T cells, and decreases B-1a cell population to weaken inflammation.

However, our current knowledge of Siglecs in the pathogenesis and therapy of sepsis is in its infancy. Most research has focused on the pathogens-related sepsis, but the researches using CLP model, the golden standard of sepsis, are few and far between. Therefore, more Siglecs-related studies using CLP model are in urgent demand. In addition, more researches are also needed in the function of T cells and NK cells with the participation of Siglecs in sepsis. As the cytokines storms stage and the immunosuppression stage of sepsis are totally different immune state, investigating the different functions of Siglecs in the different stages of sepsis is also very meaningful. Collectively, investigating the roles played by Siglecs in the immune response will not only contribute to the therapy of sepsis, but also hold great potentials in the treatment of other inflammatory diseases.

**AUTHOR CONTRIBUTIONS**

Y-CL and M-MY drafted the manuscript and performed a literature review. S-TS and Y-FC were served as chief physicians. All authors read and approved the final manuscript.

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