

REVIEW ARTICLE

PATHOPHYSIOLOGY OF SEPSIS

Andualem Mossie¹

¹Dr Andualem Mossie, PhD, Associate Professor of Medical Physiology
Department of Biomedical Sciences (Physiology)
College of Public Health and Medical Sciences
Jimma University

Telephone: 00251912051945

E-mail: andualemm2000@gmail.com

ABSTRACT

Background: Sepsis, a systemic inflammatory response to infection, resulting in organ failure and death. Pathogenic microbial agents trigger cascades of events in sepsis by stimulating the host's immune system. Macrophages require surface receptors to present pathogenic molecule to Toll-like receptors (TLR_s) to be activated and release proinflammatory cytokines such as tumour necrosis factor (TNF α) and interleukins (IL-1 β , IL-6 and IL-8) together with activation of the coagulation cascades. In the last three decade, several basic science and clinical researches have been conducted in order to improve treatment and survival of patients with sepsis. The main aim of this review is, thus, to highlight recent advances in the pathophysiology of sepsis for effective management of the problem.

Sources and Methods: More than 200 literatures of both experimental models of sepsis as well as clinical trials that were published from 1990 to 2011 were reviewed. Medline search was done based on the following key words; sepsis, systemic inflammatory response syndrome (SIRS), pathophysiology of sepsis, sepsis and coagulation, treatment of sepsis, mediator cytokines, multiple organ dysfunction syndrome (MODS) were performed. Both original publications and review articles were utilized.

Results: Both preclinical and clinical models reserches on sepsis have shown significant development in the identification of diagnostic molecules such as cytokines and cell surface receptors that are potential biomarkers of sepsis in the clinical settings. Certain advancements have also been made in the understanding of the pathophysiologic, immunologic and biochemical pathway of sepsis. Despite these significant advances in the pathophysiology of sepsis, the mortality rate is still high, 30% to 50%.

Conclusion: To win the war against sepsis, the pathophysiology of sepsis should be clearly elucidated. Early administration of broad spectrum antibiotics, effective source control, maintenance of normal glucose levels and goal-directed hemodynamic resuscitation are the cornerstone of successful management of sepsis.

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Key works: SIRS, Sepsis, Severe Sepsis, Septic Shock, Pathophysiology.

INTRODUCTION

Sepsis, severe sepsis, septic shock represents progressive stages of the same illness—a systemic response to infection mediated via macrophage derived cytokines (1). Severe sepsis and septic shock are major challenges in intensive care units (ICU) (2). Severe sepsis is associated with a mortality rate of 25 - 30% and mortality due to septic shock is 50-85% (3).

Pathogenic microbial agents trigger cascades of events in sepsis by stimulating the host's immune system. Severe sepsis is characterized by over activation of the host's immune system resulting in production of excessive inflammatory and anti-inflammatory cytokines that may result activation of the intravascular coagulation, systemic inflammation, systemic vasodilatation, increased capillary permeability and apoptosis that lead to the development of progressive multiple organ failure and ultimately death (4).

Endotoxin (lipopolysaccharide, LPS) from Gram-negative bacteria, peptidoglycan and flagellan from Gram-negative and Gram-positive bacteria, lipotechoic acid (LTA) from Gram-positive bacteria, mannan from fungi, and other antigens from infectious agents stimulate macrophages and monocytes to release tumour necrosis factor alpha (TNF- α), resulting in a cascade of cytokine release (5, 6). Following the release of TNF- α , other pro-inflammatory cytokines, including interleukins (IL-1 β , IL-6) are released into the circulation (7). These cytokines trigger numerous additional pro-inflammatory events from endothelial cells and leukocytes (5). TNF- α acts in conjunction with IL-1 β to produce the clinical signs of the systemic inflammatory response syndrome described in table 1, and their synergistic effects are probably responsible for the hypotension and resultant organ dysfunction seen early in the course of severe sepsis (8, 9, 10). Over the last 20 years, although there are promising therapeutic options supported by research evidences to improve survival of patients with sepsis, the overall result is without success, perhaps because of the complexity of pathogenesis of sepsis and basic pathophysiological differences between actual human sepsis and experimental animal models (11, 12).

Pathophysiology of sepsis is complex processes that encompass interaction of proinflammatory and anti-inflammatory cytokines, humoral, cellular, and circulatory involvement resulting from dysregulation of the immune response to infection and associated with hematological, hemodynamic and metabolic disturbances (9, 11).

Clear understanding of the pathophysiologic scope of the problem is the most essential aspect for researchers to look for effective therapies and for clinicians to manage it properly. Therefore, the main aim of this descriptive review is to elucidate the pathophysiology of sepsis for easy management of the problem.

SOURCES AND MATERIALS

More than 200 literatures of both experimental models of sepsis as well as clinical trials that have been published from 1990 up to 2011 were reviewed to organize this descriptive review. Medline search was made only in English language using the following key words; sepsis, SIRS, pathophysiology of sepsis, pathogenesis of sepsis, epidemiology of sepsis, sepsis and coagulation, treatment of sepsis, mediators cytokine, MODS were performed. Both original publications and review articles were utilized.

RESULTS

Flow chart

200 Literatures were reviewed, however,



Old articles published before 1990

Articles written in French and Deutsche

Articles that were not freely accessible were excluded

70 Articles were included

SEPSIS: TERMS AND DEFINITIONS

The American College of Chest Physicians/Society of Critical Care Medicine (ACCP/SCCM) consensus committee agreed on a set of definitions to create a universal understanding (13).

SIRS, sepsis, severe sepsis and septic shock represent the continuum of the same systemic response with increasing severity of the disease process (14).

By consensus, SIRS is defined as a widespread inflammatory response to a variety of severe clinical injuries. This syndrome is clinically indicated by the presence of 2 or more of the following: temperature $>38^{\circ}\text{C}$ or $<36^{\circ}\text{C}$; heart rate $>90/\text{min}$; respiratory rate $>20/\text{min}$ or $\text{PaCO}_2 <32 \text{ mm Hg}$; White blood cell count $>12 \times 10^9/\text{L}$ or $<4 \times 10^9/\text{L}$, or with $>10\%$ immature (band) forms (13, 15).

There are conditions that may produce systemic inflammatory response syndrome and organ dysfunction without any evidence of infection. These include tissue trauma, burn, and pancreatitis. (16).

As presented in table 1, sepsis is defined by the clinical signs of a systemic response to infection (17). In sepsis, the clinical signs of SIRS are presented together with definitive evidence of infection (18). Patients who have sepsis with acute organ dysfunction are considered to have severe sepsis (19). Sepsis is considered to be severe when it is associated with organ dysfunction, hypo-perfusion or hypotension. The manifestations of hypoperfusion may include, but are not limited to, lactic acidosis, oliguria and an acute alteration in mental status. (17,19). Septic shock is severe sepsis with hypotension, which represents a systolic blood pressure (SBP) $< 90 \text{ mmHg}$ or mean arterial pressure (MAP) $< 70 \text{ mmHg}$ or a SBP decrease by $>40 \text{ mmHg}$ below normal for age in the absence of other causes of hypotension (13, 15).

EPIDEMIOLOGY OF SEPSIS

Sepsis affects 1.8 million people worldwide every year and on average each case costs more than US\$22,000 to treat (20). The incidence of sepsis in the United States has been estimated to be 750,000/year (300 cases per 100,000 populations) with a mortality rate of 28% to 50%, resulting in an economic expense of nearly US\$17 billion annually (17). As shown in table 2, the incidence of sepsis is bigger than that of other diseases. It is the second leading cause of death among patients in non-coronary intensive care units and the 10th leading cause of death overall in the United States (21, 22). A greater number of sepsis cases are caused by infection with gram-positive organisms than gram-negative organisms, and fungal infections now account for 6% of cases (17). The annual incidence of sepsis is increasing by 8%. This upsurge has been attributed to such risk factors, including an increased number of aging population, emergence of antibiotic resistant microbial pathogens, the use of immunosuppressant drugs, an increased in invasive medical interventions, an increased in the rate of HIV AIDS infection (21).

ETIOLOGY OF SEPSIS: MICROBIAL CAUSATIVE AGENTS

The source of infection resulting in sepsis is most frequently bacterial, (23). Recent studies described that there has been a shift of causative agents of sepsis from the predominant Gram-negative bacterial infection in the late 1970s and 1980s to Gram-positive bacteria at present (17). Gram-positive and polymicrobial infection accounted for 30%-50% and 25% of cases respectively (Table 3) (24).

The most common sites of infection are the lungs (40%), abdomen (30%) and urinary tract (10%). The high rate of pulmonary infection may reflect frequent and prolonged use of mechanical ventilation and the rising occurrence of nosocomial pathogens encountered in intensive care facilities (25).

Gram-negative bacilli, mainly E.coli, Klebsiella species, Enterobacter, Proteus and Pseudomonas aeruginosa are the commonest microbes isolated from patients with sepsis and septic shock (26). Lipopolysaccharide (LPS, endotoxin) is an important component of the outer membrane of Gram-negative bacteria and has a pivotal role in inducing sepsis. (27). Gram-negative infection usually occurs in the lung, abdomen, and the urinary tract (23). Gram-positive cocci, mainly Staphylococci (Staphylococcus aureus, and Coagulase Negative Staphylococci) and Streptococci (Streptococcus pyogenes, Streptococcus pneumoniae, Viridians Streptococci) are the commonest cause of Gram-positive sepsis (28). They are usually responsible for infection of blood stream, skin, soft tissues and respiratory tract. Staphylococcus aureus produce Toxic Shock Syndrome Toxin (TSST) that results in septic shock and Streptococcus pyogenes produce Streptococcal pyrogenic exotoxin A (SEA) (29).

Gram-positive organisms cause sepsis by at least two mechanisms: by producing exotoxins that act as superantigens and by their cell wall components that can stimulate immune cells. (28). Superantigens are molecules that bind to MHC class II molecules of antigen presenting cells and T-cell receptors. In doing so, they activate large number of T-cells to produce massive amount of proinflammatory cytokines. Staphylococcal enterotoxins, toxic shock syndrome toxin-1, and Streptococcal pyrogenic exotoxin A are examples of bacterial superantigens (30, 31).

Gram-positive bacteria without exotoxin can also induce septic shock, by stimulating the innate immune system through similar mechanisms to those in Gram-negative sepsis. Indeed, toll-like receptors (TLR-2) has been shown to mediate cellular responses to heat killed Gram-positive bacteria and their cell wall structure (peptidoglycan, lipoproteins, lipoteichoic acid, and phenol soluble modulins) (32).

PATHOPHYSIOLOGY OF SEPSIS

Sepsis, severe sepsis, septic shock and multiple organ failure are complex processes that encompass pro-inflammatory, anti-inflammatory, humoral, cellular, and circulatory involvement resulting from dysregulation of the immune response to infection (28). The pathophysiologic process has been linked to the contribution of apoptosis or programmed cell death, which may have an important role in the organ dysfunction. (29).

The pathogenesis of sepsis involves a complex interaction between the host immune system and the infecting microorganisms (33). Bacteria and their products trigger cascades of cellular response in the host's body that involve several cell types (leukocyte, mast cells, endothelial cells and platelets), and several cellular pathways (pro-inflammatory, anti-inflammatory, coagulation cascades, complement activation, adhesion and apoptosis) (34).

The normal immune response to bacterial infection is a complex inflammatory process that attempts to localize and limit the spread of the infection and repair the tissue (27). This response involves the activation of phagocytes and endothelial cells to produce both pro-inflammatory and anti-inflammatory mediators. The balance between these groups of mediators helps to protect the host against the invading pathogens and to facilitate tissue healing. Sepsis comes when the balance is lost, in which the inflammatory response to infection produced in excess and extends beyond the infected site causing generalized functional alterations (35). Due to exaggerated immune response to the invading pathogen, widespread inflammatory mediators and vasodilators are released (36). Inflammatory cytokines such as $\text{TNF}\alpha$, $\text{IL-1}\beta$, and IL-6 , are secreted by the monocytes and macrophages. Tissue factor (TF) is expressed by monocytes and the damaged vascular endothelium. Coagulation is activated with release of thrombin and the formation of the fibrin clot in order to wall off the spread of the infection (35).

Circulatory problem arises from the combination of vasodilatation, capillary leakage, and reduced myocardial contractility. Organ failure may be the first clinical sign of severe sepsis, as the consequences of the excessive products of inflammatory mediators. Mortality rates increase with the increase of failed organs (21).

The clinical manifestation of sepsis is the consequence of complex groups of mediators secreted by the immune cells in response to pathogens such as LPS (endotoxin) produced by bacteria (33, 37, 38). Free LPS that is formed from degradation of Gram-negative bacteria cell wall, once in the circulation binds with LPS binding protein (LBP), to form a complex (39). This complex then binds to monocyte/macrophage cell surface receptor, CD_{14} and TLR_4 that result in cellular activation and secretion of inflammatory cytokines (40). Prominent among the central mediators of sepsis produced in response to LPS activation of receptors are $\text{TNF}\alpha$, $\text{IL-1}\beta$, IL-6 , eicosanoids, platelet activating factor (PAF), oxygen free radicals and nitric oxide (41).

These primary cytokines and many other intermediate and distant mediators interact with each other, setting into motion of complex cascades of reactions and series of local and systemic inflammatory response (39, 40). Secretion of $\text{TNF-}\alpha$ mainly from macrophages is particularly strongly stimulated by LPS and it is believed that this mediator, but not LPS directly, is one primary agent, which sets in motion a pronounced cellular, metabolic and vascular changes occurred during sepsis (41, 42). Serum $\text{TNF-}\alpha$ and $\text{IL-1}\beta$ reach toxic level in mice and human volunteers within 1-2 hours after LPS infusion (43).

Patients with severe sepsis exhibited biphasic hemodynamic and immunological response. (a) The early, hyperdynamic phase is characterized by increased cardiac output, tissue perfusion, and decreased vascular resistance. The hallmark of this early phase is the pro-inflammatory state that is mediated primarily by neutrophils, macrophages, and monocytes that have been stimulated by microbes (44); followed by (b) Hyporeponsive phase (cold phase, vasoconstricted phase): a period of immunodepression in which anti-inflammatory cytokines are produced in order to counterbalance inflammatory mediators resulting in immunoparalysis (45). It is characterized by cold extremities, hypotension, small pulse pressure, low cardiac output, high SVR and low cardiac contractility. It is associated with loss of phagocytic functions, and decreased Th_1 cytokine release (44, 46). Persistence of this hypo responsiveness is associated with increased risk of nosocomial infection and death (47, 48). Immunoparalysis: CD_4 lymphocytes play a key role in the inflammatory response during sepsis. Early in the process of sepsis, these cells assume a T-helper-1 (Th_1) phenotype, where they produce large amounts of the pro-inflammatory mediators, including $\text{IF-}\gamma$, $\text{TNF-}\alpha$, and IL-2 (49). T-lymphocytes may evolve over time to a T-helper-2 (Th_2) phenotype, whereby the CD_4 lymphocytes produce anti-inflammatory cytokines, including IL-10 , IL-4 , and IL-13 (50). These cytokines dampen the immune response and can lead to the deactivation of monocytes. Additionally, $\text{TNF-}\alpha$ released early can cause apoptosis of lymphocytes in the gut, leading to further immune suppression (51).

Sepsis-triggered immunological cascade is multimodal: initial systemic inflammatory response syndrome (SIRS; excessive pro-inflammatory, but no/low anti-inflammatory mediators), intermediate homeostasis with a mixed anti-inflammatory response syndrome (MARS; both pro-inflammatory and anti-inflammatory mediators)

and final compensatory anti-inflammatory response syndrome (CARS; excessive anti-inflammatory, but no/low pro-inflammatory mediators) (52, 53, 54).

The balance between pro-inflammatory and anti-inflammatory mediators derived from the innate immune system defines the progression and severity of infection. If unbalanced, an overproduction of endogenous pro-inflammatory mediators, including cytokines, platelet activating factor, oxygen radicals and nitric oxide, synergistically interact to mediate hypotension, multiple organ failure and death (53). Progression from sepsis to septic shock coincides with an increase in circulating levels of pro-inflammatory cytokines such as TNF- α , IFN- γ , IL-1 β , and IL-6 (39). Administration of TNF α in animal models mediates sepsis typical to endotoxin and removal of TNF- α from animals challenged with LPS using pharmacological strategies significantly improves survival (41).

The pathophysiology of sepsis can be initiated by the outer membrane component of Gram-negative organisms (e.g., lipopolysaccharide [LPS], lipid A, endotoxin) or Gram-positive organisms (e.g. lipoteichoic acid, peptidoglycan), as well as fungal, viral, and parasitic components (55, 56). Signalling by these mediators occurs via a family of transmembrane receptors known as toll-like receptors (TLRs) (57). Within the monocytes, nuclear factor- κ B (NF- κ B), is activated, which leads to the production of pro-inflammatory cytokines, TNF- α , and IL-1 β . TNF- α and IL-1 β lead to the production of toxic downstream mediators, including prostaglandins, leukotrienes, platelet-activating factor, and phospholipase A₂ (56). These mediators damage the endothelial lining, leading to increased capillary leakage. Finally, activated macrophages and neutrophils release nitric oxide, a potent vasodilator that leads to septic shock (58).

Pathogenic molecules derived from microbial infectious agents that can bind with specific receptors on immune cells to activate production of cytokines and chemokines are collectively called pathogen associated molecular pattern (PAMP) (57). The common PAMPs are surface molecules such as, lipopolysaccharide (LPS), peptidoglycan (PGN), lipoteichoic acid (LTA), unmethylated bacterial DNA (CpG DNA), lipoarabinomannan (LAM) of mycobacterium, viral ds RNA, flagellin, lipopeptides, mannans and zymosan of yeast and choline-containing phosphoglycolipids and internal motifs released during bacterial lysis such as heat shock proteins and DNA fragments (59).

Recognition of PAMPs by PRRs results in the activation of different intercellular signalling cascades that in turn lead to the expression of various effector molecules (57). Immune cells recognize microbes through pattern recognition receptors (PRRs) on the cells (59). Toll-like receptors (TLRs) are among PRRs and activate immune cells to produce pro-inflammatory cytokines and chemokines (60). Toll-like receptors regulate antimicrobial host defense mechanisms and play a central role in the activation of innate immunity (61). Toll-like receptors are a family of cellular surface protein receptors that recognize pathogenic molecules of various microorganisms. Bacterial components including lipopolysaccharide, lipoteichoic acid, flagellin and other cell wall components interact with Toll-like receptors. Different microbial pathogenic products bind to different receptors. TLR2 and TLR6 have been shown to react with lipoteichoic acid, TLR4 specifically recognizes lipopolysaccharide, and TLR5 with flagellin (62). These findings implicate that the innate immune response is tailored in a pathogen specific manner (61). In the initial phase of infection, Toll-like receptors activate the innate immune system; as a result invading pathogens are destroyed by macrophages, natural killer cells and complement system. In the second phase, Toll-like receptors form an important link between innate immunity and adaptive immunity, by activating T and B lymphocytes to produce cytokines (63, 64, 65, 66).

TREATMENT OF SEPSIS

Current treatment of severe sepsis involves early appropriate antibiotic therapy, surgical drainage of infected site, mechanical ventilation, optimizing oxygen delivery to patients with lactic acidosis, adequate fluid resuscitation (colloids, crystalloids and other plasma expanders), administration of vasopressors (dopamine, noradrenaline) to improve cardiovascular parameters, tight glycemic control, blood transfusion in case of low haemoglobin (< 7 g/dl), use of low dose corticosteroid, and supportive therapy. Despite all these therapeutic applications, mortality remains high; therefore, the search endeavour in sepsis should be scaled up to find appropriate immune-modulatory therapies with remarkable survival records (68, 69, 70).

CONCLUSION

Although there are promising therapeutic options supported by research evidences to improve survival of patients with sepsis, the overall result is without success, perhaps for two major reasons: the complexity of pathogenesis of sepsis and basic pathophysiological differences between actual human sepsis and experimental animal models. The pathophysiology of sepsis involves a complex interaction of inflammatory mediators, activation of host cells to release inflammatory and counter-inflammatory mediators. Multiple derangements exist in sepsis involving several different organs and systems. Having a clear understanding of the

pathophysiology of sepsis is a precondition for an effective management of the problem. The early administration of empirical antibiotic therapy, effective source control, and goal-directed hemodynamic resuscitation are the cornerstone of successful management. Further large scale study on sepsis is recommended.

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Table 1:- Standard definition of sepsis and related terms (13, 15).

Terms	Definitions
1. SIRS	<i>The systemic inflammatory response to a wide variety of severe clinical insults (infection, pancreatitis, ischemia, trauma, autoimmunity and others), manifested by 2 or more of the following</i> Core Temperature: <36 °C or >38 °C Heart Rate: >90 beats/min Respiratory Rate: >20 breaths/min or PaCO ₂ <32 mm Hg WBC count: <4,000/mm ³ or >12,000/mm ³ or >10% Immature neutrophils (band) forms.
2. Sepsis	<i>The systemic inflammatory response to infection. It is SIRS plus documented infection.</i>
3. Severe Sepsis	(Sepsis + ≥ 1 organ failure) <i>Sepsis associated with organ dysfunction, hypoperfusion or hypotension. Hypoperfusion may include, but not limited to, lactic acidosis, oliguria, or an acute alteration in mental status.</i>
4. Septic Shock	<i>A subset of severe sepsis and defined as sepsis induced hypotension despite adequate fluid resuscitation along with perfusion abnormalities that may include but not limited to, lactic acidosis, oliguria, or an acute alteration in mental status. Hypotension responds to inotropic or vasopressor agents.</i>
5. MODS	<i>Presence of altered organ functions in an acutely ill patients such that homeostasis cannot be maintained without intervention.</i>
6. Sepsis-induced Hypotension	<i>A systolic BP < 90 mm Hg or a reduction of ≥ 40 mm Hg from baseline in the absence of other causes for hypotension</i>

Table 2:- Incidence of sepsis in comparison with other diseases worldwide.

HIV AIDS	17/100,000 inhabitants
Colon carcinoma	50/100,000 inhabitants
Breast cancer	110/100,000 inhabitants
Cardiac insufficiency	196/100,000 inhabitants
Sepsis	300/100,000 inhabitants

Table 3: Estimated frequency of pathogens in sepsis (24)

Pathogens	Frequency
Gram-positive bacteria	30 –50%
Staphylococcus aureus	20-35%
Other Staphylococcus spp	1-3%
Streptococcus pneumoniae	9-12%
Other Streptococcus spp	6-11%
Enterococcus spp	3-13%
Other Gram positive bacteria	2-7%
Gram-negative bacteria	25-30%
E.coli	9-17%
Pseudomonas aeruginosa	8-15%
Klesiella pneumoniae	2-7%
Other enterobacter spp	6-16%
Other Gram negative bacteria	8-29%
Fungus	3-5%
Protozoa (parasites)	1-3%
Virus	2-4%

Corresponding Author

Dr Andualem Mossie

Department of Biomedical Sciences (Physiology)

College of Public Health and Medical Sciences

Jimma University, Jimma, Ethiopia

Telephone: 00251912051945

E-mail: andualemm2000@gmail.com