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**Cellular Hibernation and Immunoparalysis in the Context of Sepsis Induced Organ
Dysfunction**

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Dedicated to my beloved father.

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1. Abstract

Sepsis remains the leading cause of death in the intensive care unit. The limited effectivity of current therapies suggests that more research targeting the complex pathophysiologic processes of sepsis is essential to establish a comprehensive treatment with significant outcome improvement. This thesis is aimed to describe the counterintuitive pathophysiology of sepsis while focusing on the immunologic course of sepsis, the effects of cytokine storm, the cellular hibernation response to stress and their implication in organ dysfunction, the state of immune cell paralysis and potential treatment options.

2. Keywords

Sepsis, MODS, Cytokine Storm, Cellular Hibernation, Immunoparalysis, Immunotherapy

3. Introduction

Sepsis is defined as a dysregulated host immune response to infection, which can lead to life-threatening organ dysfunction (1).

The epidemiologic relevance was presented in 2020 by the World Health Organization in the “Global Report on the Epidemiology and Burden of Sepsis (2020)”, stating that Sepsis accounts for approximately 20% of overall deaths worldwide and the mortality rate of septic patients is approximately 20%, while for those requiring treatment in the Intensive Care Unit it is up to 42%.

Research of the last decades changed the concept of systemic infection from pathogen centered to a mixed pathophysiology, in which the host response to infection plays a major role and lead to the first consensual definition of sepsis as “Systemic response to an infection” in 1992, diagnosed by the presence of a Systemic Inflammatory Response Syndrome (SIRS) plus proven or suspected infection (2). In 2016, a new definition was conceptualized, defining sepsis as “Life-threatening organ dysfunction due to a dysregulated host immune response to infection”, along with new diagnostic criteria composed of a minimum 2 point elevation of the SOFA (Sepsis-related Organ Failure Assessment) Score plus proven or suspected infection (1), which allows the early detection of sepsis and timely

administration of bundle care, which has essential influence on the clinical course and mortality (1).

The mortality of septic patients is based on the clinical course and increases with the development of complications, such as septic shock and/or MODS (multi-organ dysfunction syndrome). The former is defined clinically as “persisting hypotension requiring vasopressors to maintain MAP >56 mmHg (Mean Arterial Pressure) plus a serum lactate concentration of >2 mmol/L despite adequate volume resuscitation” and increases mortality to 40% (1). The latter is assessed with the SOFA score (Sequential Organ Failure Assessment Score), while the mortality increases proportionally to the number of failing organs (1).

Despite intensive scientific research in the field of septic pathology, the clinical translation of experimental findings is often unsuccessful. This thesis is aimed to describe the current understanding of cellular hibernation and immunoparalysis which are important pathophysiologic factors of sepsis-induced organ dysfunction.

4. Literature Search Strategy

A Systematic literature search was conducted using the Database “Medline” with the search engine “PubMed”, which is maintained by the “National Center for Biotechnology and Information (NCBI)” at the “National Library of Medicine”. The selected time range of articles spans from 1986 to 2021. Main key words used: Sepsis, Septic Shock, Septic Organ Dysfunction, Cytokine Storm, Inflammatory Cytokines, Inflammatory Mediators, Disease Tolerance, Disease Resistance, Immunity, Immunoparalysis, Cellular Hibernation, Mortality in Sepsis, Immunotherapy, Blood purification techniques.

5. Sepsis: Course and Mortality

While many factors, such as pathogen (type, quantity) and patient factors (immune status, age, co-morbidities) influence the host response to an infection (3–5), special attention can be drawn to the epigenetic control of gene transcription and genetic polymorphism of sepsis-associated genes, of which many have been and continue to be identified leading to the understanding of the role of genotypic clusters involved in the pathway of host response (6–9). Especially cytokine receptor SNP’s (single nucleotide polymorphisms) and transcription

factor SNP's have been studied, outcome-correlated (6,10–12) and may be used as biomarkers of sepsis susceptibility and in the prognosis of the course of the disease (13).

In contrast to the normal response to an acute infection, which is proportional to the pathogen burden and terminated after elimination thereof, sepsis causes a specific pattern of an initial hyper inflammatory phase called SIRS (systemic inflammatory response syndrome) followed by a late hypo inflammatory phase called CARS (compensatory anti-inflammatory response syndrome) (4,5). The former is caused by PAMP (pathogen-associated molecular pattern) and DAMP (damage-associated molecular pattern) induced massive and uncontrolled activation of PRR's (pathogen recognition receptors), which are expressed on the surface of cells of the innate immune system, for instance TLR's (toll-like receptors), leading to the excessive release of pro-inflammatory cytokines (cytokine storm) (3,14), inducing pro-inflammatory cascades, with IL1 β , IL-6, IL-12, IL-17 (Interleukins) being the most prominent (6). The latter is caused by the down-regulation of the primary effector cells (granulocytes, macrophages, monocytes, mast cells and natural killer cells) via anti-inflammatory cytokines, mainly IL-1 RA (receptor antagonist), IL-4 and IL-10 (6) with apoptotic and autophagic loss, as well as the formation of tolerance mechanisms by lymphocytes (15), resulting in a state of immunoparalysis.

The shift in the immune homeostasis correlates with distinct mortality phases, that can be divided into an early phase (day 1-5), late phase I (day 6-15) and late phase II (day 16-150) (16). Early phase coincides with the hyperinflammatory phase, where death is mainly caused by hyperinflammatory syndromes, such as toxic shock syndrome, refractory shock and severe metabolic disturbances (4) but may be anticipated by early recognition using the SOFA score with timely bundle treatment initiation (17–20) amidst a cumulative mortality of 10% (15). Late phase I and II are divided by different cumulative mortalities of 20-40% and 50-70% respectively (15) and marked by immunosuppression making the host vulnerable for a flare up of a not sufficiently cleared primary infection, reactivation of latent viruses or superinfection by opportunistic pathogens and hospital organisms (4,16). Thus, sepsis-induced immunosuppression is the major cause of high mortality and organ dysfunction in sepsis (13,21).

It is important to note that certain patient groups exhibit a different pattern of sepsis course. Patients with intrinsic or extrinsic immunosuppression may have an absent or diminished early pro-inflammatory phase (22,23) and elderly patients are, due to immunosenescence,

more likely to develop an early late phase with a pronounced immunoparalysis (22,24), which may alter the mortality pattern described above.

6. Disease Resistance vs. Disease Tolerance

Disease resistance describes the classical view of infectious response, defined as the ability of the immune system to recognize and eliminate pathogens, despite the possible causation of collateral damage caused by this process (25). In contrast, disease tolerance is the ability to limit tissue injury caused by the invading pathogen itself, or the inflammatory/immune response to them by utilizing tissue damage control mechanisms as the cellular stress and/or damage response (25–27).

Due to the collateral damage caused by the immune cascades activated in sepsis, the severity of the inflammatory responses has been associated with the development of multiple organ dysfunction (28). This led to the concept of disease tolerance in septic pathology, as an induction of tolerance may regulate or prevent the course of sepsis development, with the protective effect being presumable disease- and pathogen-specific (25,27,28). Multiple mechanisms of disease tolerance induction are under investigation, for example the low-dose treatment with cytostatic drugs (29).

7. Cytokine Storm

Cytokines are small signaling proteins (< 40 kDa) that bind to specific receptors acting in an autocrine, paracrine and endocrine manner to induce target cell activation, proliferation and migration, thereby encompassing an important immunomodulating function (6,30). They are produced predominantly by macrophages and T cells (31) and are classified into Interleukins, Chemokines, Interferons, Tumor Necrosis Factors and Growth Factors with Interleukins having the major impact on immune cells in the context of infectious disease (6). The excessive release of cytokines during the early stage of sepsis pathogenesis is termed “Cytokine Storm” in which cytokine cascades build a positive feedback loop with immune cells resulting in an overwhelming inflammatory response to infection (31).

Important potentiators and potentially therapeutic targets of the cytokine storm are transcription factors. Especially NFκB (nuclear transcription factor κ B), SP-1 (specific

Protein 1), hPNPase(oid-35) (polynucleotide phosphorylase), AP-1 (activator protein 1) and PPARs (peroxisome proliferator-activated receptors) have been identified to play a role in upregulating genes coding for cytokines (32–34).

In this context NFκB is a transcription factor of major importance. It is activated by more than 400 stimuli, for example bacterial toxins, bacterial products and pro-inflammatory cytokines, by binding to specific subtypes of TLR's (toll-like receptors) and, in turn, upregulates the transcription of many pro-inflammatory genes encoding cytokines, cellular adhesion molecules and enzymes, such as COX-2 and iNOS (cyclooxygenase 2, inducible nitric oxide synthase) (31,35–37). Mycobacteria are identified to activate TLR1, gram-positive bacteria predominantly activate TLR 2 via PG (peptidoglycan) and gram-negative bacteria primarily bind to TLR-4 via LPS (lipopolysaccharide) (37–39).

Studies have shown an elevation of NFκB in all organs upon exposure of bacterial toxins, as well as in PMN's (peripheral mononuclear cells) of septic patients, indicating a role in septic pathology (37,40–43). Further the NFκB level positively correlates with sepsis mortality and the APACHE Score (Acute Physiology and Chronic Health Evaluation Score) predicting hospital mortality in critically-ill patients (37,44,45). The resulting signaling cascades of NFκB activation eventually leads to tissue injury and contributes to the development of tissue injury and MODS (37).

Various NFκB inhibitors have been identified and tested clinically or in animal models to identify their therapeutic potential and are found to have a positive effect on septic mortality. An improved survival of patients with severe sepsis was observed upon treatment with Human Recombinant Active Protein C (37,38), which has been approved by the EMA (European Medical Association) in 2002 after the PROWESS trial results indicated a significant improvement in the 28 day mortality rate (46) but was taken off-market in 2011 after further trials could not reproduce the positive effect of the drug. The outcome of endotoxemia in LPS-induced Shock is positively influenced by IL-10 and a deletion of the gene encoding for poly(ADP-ribose)polymerase-1 made mice resistant to endotoxic shock (37,47–49). Furthermore an endotoxin tolerance mechanism is observed with increased IκBα, p50 or p52 expression manifesting with a diminished NFκB activity and expression of NFκB-dependent genes (37,50–52).

Until today, trials targeting specific cytokines with monoclonal antibodies or the use of apheresis to remove cytokines from the circulation have not shown a clear benefit in the

treatment of sepsis-associated cytokine storm, so that organ-supportive therapy and antibiotic control of pathogen burden remain the mainstay of treatment (53). Although certain patient groups may benefit from targeted immunotherapy, the lack of research on inclusion criteria for certain drugs does not allow a clear matching, making it necessary to focus on precision diagnostics in future research (53).

8. Mechanisms of MODS

Multiple organ dysfunction is the most fatal complication of sepsis. Causative molecular pathways lead to microcirculatory and mitochondrial dysfunction resulting in organ damage.

The six most relevant organ systems in septic organ pathology are the cardiovascular, respiratory, renal, hepatic, hematologic, nervous and gastrointestinal systems, in order of their sequential failure (13).

The release of pro-inflammatory cytokines, such as TNF- α (tumor necrosis factor alpha), IL-1, IL-6, IL-12, IL-18 and INF- γ (interferon gamma) trigger an acute phase response leading to symptoms such as pyrexia, tachypnea, tachycardia, metabolic alterations, specifically increased gluconeogenesis, muscle catabolism and altered lipid metabolism, activation of the coagulation and complement pathway, downregulation of natural anticoagulants, leukocytosis with neutrophilic predominance, and the production of acute phase proteins in the liver, namely C-reactive protein and serum amyloid A, induced by IL-1 and TNF- α plus fibrinogen and α -2 macroglobulin, induced by IL-6 (54,55,37,56). In turn they upregulate the NF κ B pathway leading to a positive feedback loop of pro-inflammatory mediator production (37,53).

Increased expression of cellular adhesion molecules such as ICAM-1 (intracellular adhesion molecule 1) and VCAM-1 (vascular adhesion molecule 1) and chemokines such as IL-8, MIP-1/2 (macrophage inflammatory protein 1/2), MCP 1/2 (monocyte chemoattractant protein 1/2) and CINC (cytokine-induced neutrophil chemoattractant) are a milestone for neutrophilic infiltration into organs, their activation, and the subsequent release of ROS (reactive oxygen species), RNS (reactive nitrogen species) and proteolytic enzymes (13,37,57–61). This leads to the development of endotheliopathy, which is describing endothelial activation and injury followed by increased vascular permeability leading to

leakage of molecules and cells into tissue with resulting edema, hemorrhage, infection or ischemia (13,37,57–61).

A upregulated expression of COX-2 (cyclo-oxygenase 2), 5-LO (5 lipo-oxygenase) and FLAP (5-LO-activating protein) lead to the production of PG's (prostaglandins), LT's (leukotrienes) and TBXA2 (thromboxane A2), which have an effect on the cardiovascular system, contributing by vasodilation to systemic hypotension and may cause direct tissue injury by various mechanisms (37,38,62).

Furthermore increased iNOS (inducible nitric oxide synthase) expression leads to elevated levels of NO (nitric oxide), which acts, along with PG's on VSMC's (vascular smooth muscle cells) to cause vasodilation and consequent systemic hypotension and vascular hyporeactivity (37,38,63–67). This effect may be anticipated by utilization of iNOS inhibitors (37,65,66).

Increased levels of TF (tissue factor), PAI-1 (plasminogen activator inhibitor -1) and factor VIII (coagulation factor 8) caused by pro-inflammatory cytokines activate the coagulation system and impair fibrinolysis, which may be causative for the development of the DIC (disseminated intravascular coagulation) syndrome, characterized by an initial hypercoagulable state with the formation of organ-threatening microthrombi, which may result in microvascular occlusion and/or may embolize, followed by an hypocoagulable state characterized by thrombocytopenia caused by the consumption of platelets, fibrinogen and coagulation factors, eventually leading to severe internal hemorrhage (68–72,35,37). The treatment for the DIC syndrome is based on general control of the causative agent for sepsis plus anticoagulant therapy in the hypercoagulable state and plasma or coagulation factor concentrates and/or platelet transfusion according to the PT (prothrombin time), fibrinogen concentration and platelet count, for bleeding patients (68,73–77).

Cumulation of a diminished vascular tone caused by NO and PG's and the formation of microthrombi or hemorrhages caused by DIC, lead to microcirculatory dysfunction, tissue hypoperfusion and hypoxia, resulting, along with increased vascular permeability caused by endotheliopathy, in tissue injury, which may precipitate organ failure (visualized in Fig.1).

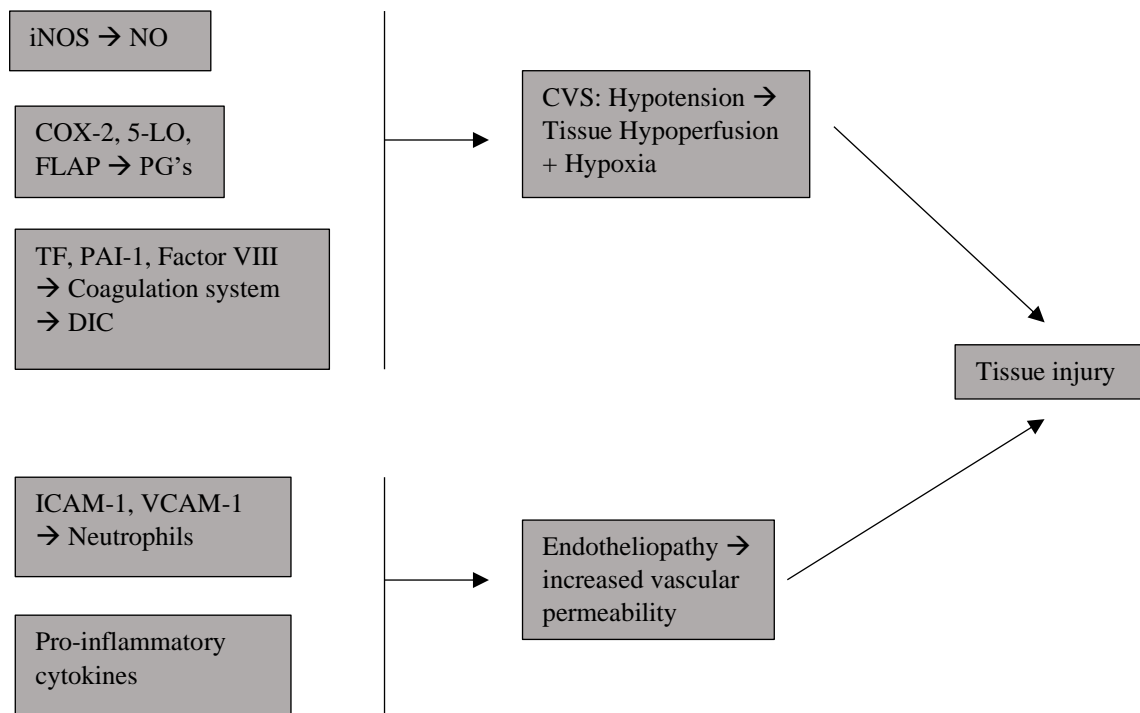


Fig. 1: Mechanisms of tissue injury in septic patients

Graphic adapted from: Liu SF, Malik AB. NF-kappa B activation as a pathological mechanism of septic shock and inflammation. *Am J Physiol Lung Cell Mol Physiol.* 2006 Apr;290(4):L622-L645.

9. Cellular Hibernation

Contrary to the assumption that septic organ dysfunction is a result of direct tissue injury followed by cellular apoptosis or necrosis, post-mortem histological examination of cells taken from patients who died from sepsis or septic shock showed a quantitative lack of apoptotic and necrotic cells sufficient to impair organ dysfunction (78). This finding led to the theory of cellular hibernation describing the rearrangement of cellular metabolism in response to hypoxia and oxidative stress to a low-energy consumption state to preserve long-term viability (79–81).

The energy required for all cellular processes is ATP (adenosine triphosphate), which can be cleaved to ADP (adenosine diphosphate) and Pi (inorganic phosphate), thereby releasing free energy. The production of ATP occurs in the mitochondria by either glycolysis or the OXPHOS (oxidative phosphorylation) process, that couples the tricarboxylic acid cycle with the electron transport chain to produce ATP. In the tricarboxylic acid cycle, proteins, fatty

acids and carbohydrates are oxidized and electrons are transferred by NADH (nicotinamide adenine dinucleotide) and FADH₂ (flavin adenine dinucleotide) to the electron transport chain, where the oxidation of the coenzymes yields ATP in a series of reactions by creation of a proton gradient across the inner mitochondrial membrane. Oxygen is essential for this process of energy production. Other factors such as the bio-availability of NADH and FADH₂ and the integrity of the numerous enzymes involved in the process of OXPHOS may also influence the efficiency (80).

Tissue injury caused by a change in mitochondrial respiration can be described in the concept of cytopathic hypoxia - a diminished oxygen utilization by cells due to mitochondrial dysfunction (82). Multiple theories about the mechanism behind the development of cytopathic hypoxia in septic pathology have been established – the inhibition of pyruvate dehydrogenase, the nitric-oxide-mediated inhibition of cytochrome a₃, the peroxynitrite-mediated inhibition of mitochondrial enzymes and the poly(ADP-ribose) polymerase hypothesis (82).

Pyruvate dehydrogenase is an enzyme catalyzing the reaction from pyruvate, the end product of glycolysis, to acetyl CoA (Acetyl Coenzyme A), which subsequently enters the tricarboxylic acid cycle whose activity is regulated by negative feedback and enzyme phosphorylation mediated by the pyruvate dehydrogenase kinase family. Studies have shown increased activity of the pyruvate dehydrogenase kinases (82–84) and an increased ratio of the inactive to active form of the pyruvate dehydrogenase complex has been found in animal models of chronic sepsis (82,85,86), limiting the utilization of the tricarboxylic acid cycle coupled to the electron transport chain to produce ATP, leading to increased lactate production and the possibility of metabolic acidosis.

As described earlier, sepsis leads to upregulation of iNOS enzymes, and subsequent increased production of NO. Cytochrome a₃ is the terminal complex of the electron transport chain and subject to competition between NO, which acts as an inhibitor, and oxygen (O₂), which acts as an activator. In conditions of a raised NO concentration (> ~1 mcM) or in a state of low pO₂ (partial oxygen pressure), NO reversibly inhibits cytochrome a₃ and thereby prevents ATP production (82,87–92).

Furthermore, NO forms ONOO⁻ (peroxynitrite) by a reaction with O₂⁻. Increased production of O₂⁻ by mitochondria was found in hypoxic conditions or in a state of cytochrome a₃ inhibition (82,93,94). Combined with raised NO levels this may lead to a substantial increase

in ONOO- production , which causes irreversible inhibition of mitochondrial energy production in in-vitro conditions by inhibition of mitochondrial F0F1ATPase responsible for ATP production from ADP (95), inhibition of complexes of the electron transport chain (95) and inhibition of the aconitase enzyme of the tricarboxylic acid cycle (96).

Poly(ADP-ribose) polymerase 1 (PARP-1) is a nuclear enzyme activated by single-strand breaks in nuclear DNA, that catalyzes the production of ADP-ribose from NAD+ and polymerization thereof (82,97,98). ROS like ONOO- produced in sepsis may induce PARP-1 by DNA damage and lead to a depletion of cellular NAD+/NADH content which is essential for the OXPHOS process to generate ATP (82,99).

In total, the altered macro- and microcirculation in early sepsis prior resuscitation results in a decreased PO₂ in tissues (100) with changes in the microcirculation remaining even post macrocirculatory resuscitation, which leads to sustained tissue hypoxia causing cellular hibernation in response to low PO₂, and mechanisms of cytopathic hypoxia, and other direct insults caused by inflammatory mediators with resulting organ dysfunction (101). Due to the diminished O₂ utilization by mitochondria, a decreased tissue VO₂ stands in contrast to the rising tissue PO₂, which is sending a negative feedback to the microcirculation causing vasoconstriction, thereby further potentiating the process of cellular hibernation (102) (visualized in figure 2).

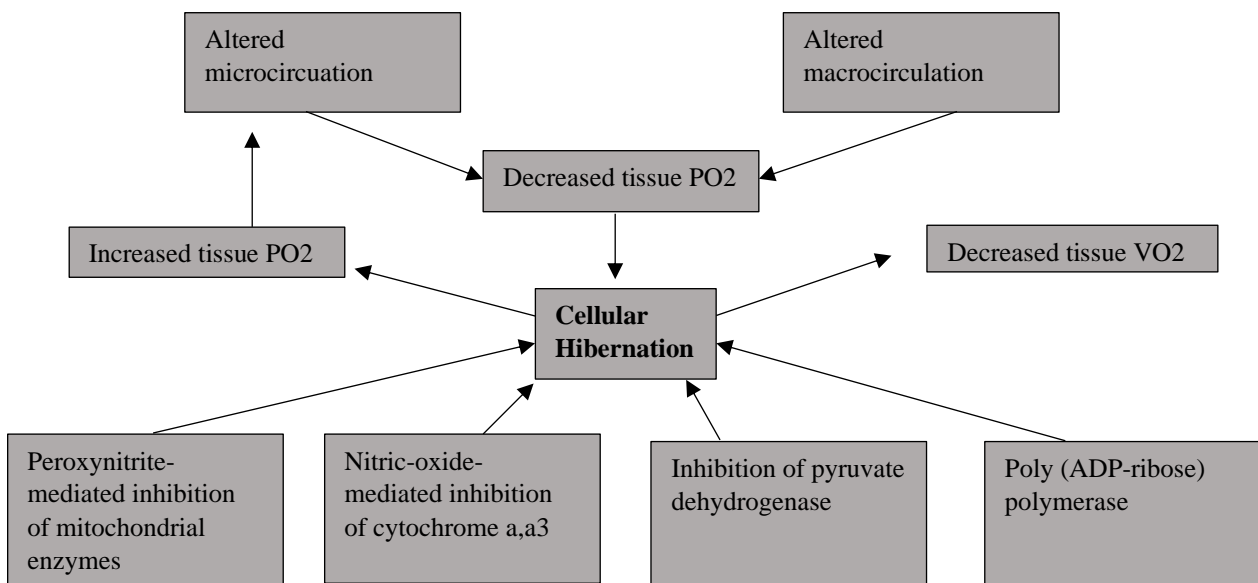


Fig. 2: Mechanisms of Cellular Hibernation

10. Immunoparalysis

Beside the quantitative loss of immune cells (15) during late phase of sepsis, namely T- cells, B- cells, macrophages and dendritic cells, as shown in post-mortem analysis (78,103–106), several mechanisms are suggested to cause a qualitative loss in innate and adaptive immune cell function, termed immunoparalysis, resulting in insufficient clearance of septic foci and the potential for superinfections and reactivation of latent viruses (22,107,108). Although it is thought to occur in the late phase of sepsis during CARS, some evidence also suggests that a certain degree of immunoparalysis may exist from the onset of sepsis (22,16,109,110). The mortality associated with the immunoparalytic state (15) suggests the importance of biologic markers to assess the degree of immunoparalysis and potentially aid in prediction of the disease course and outcome (22,111,4,112). Moreover they could be decisive for the therapeutic induction of immunocompetence by the use of immunomodulatory medications (22,111,4,112).

Monocytes have been shown to exhibit an impairment in antigen presentation and pro-inflammatory mediator production due to decreased expression of mHLA-DR (monocyte human leukocyte antigen DR isotype), which is the most researched biomarker for immunoparalysis to date and has been positively correlated with mortality (22,113–116).

Further, several receptors that initiate pro-inflammatory cascades upon stimulation have been shown to be downregulated in immunoparalytic conditions (117). Together with upregulation of negative costimulatory molecules located on lymphocytes and monocytes, PD-1 (programmed cell death protein 1) and PD-L1 (programmed cell death protein ligand 1), whose binding induces a IL-10 mediated down-regulation of inflammatory functions, this leads to a diminished immune functional capacity (22,107,118).

An upregulation of Treg's (T regulatory lymphocytes) is associated with immunoparalysis and actively down-regulate the adaptive immune response (22,119–121) by TGF- β (transforming growth factor β) and IL-10 secretion and attenuation of pro-inflammatory cytokines, as well, as induction of cytotoxic T-cells and Monocyte down-regulation (22,121–124).

Epigenetic regulation plays a major role in the development of immunoparalysis by modulation of pro- and anti-inflammatory gene expression (125) through multiple mechanisms such as histone deacetylation or methylation (125–127). An example for epigenetic modification is the shift of macrophages to the M2 phenotype (128).

Macrophages are thought to be the key mediator of shifting the cytokine balance to the anti-inflammatory site by polarizing into M2 phenotype, which is secreting high levels of diverse anti-inflammatory molecules, such as IL-10 and IL-1-RA (IL-1 receptor antagonist), thereby down-regulating pro-inflammatory cytokine release (129). In vitro experiments also suggest a generalized diminished capacity of pro-inflammatory cytokine secretion by monocytes without changes in anti-inflammatory cytokine secretion capability (130).

Further pathophysiologic mechanisms causative for sepsis-induced immunoparalysis are under investigation and provide scientists with several potential therapeutic targets of sepsis.

11. Clinical implications

Despite the establishment of timely bundle care with focus on infection source control and organ-supportive therapy, sepsis remains the leading cause of death in the ICU.

To anticipate deterioration of a septic patient, the pathophysiological processes described in the previous chapters need to be taken into consideration when searching for further treatment options. Until today many scientists are focused on finding ways to target the immunopathology of sepsis, with only a limited number of successful translations of in-vitro to in-vivo conditions.

Regarding the cytokine storm as initiator of the septic sequelae, either a decreased cytokine and inflammatory mediator production or an increased elimination can be targeted. The former implies the investigation of immunotherapeutic drugs, with many concepts from oncology being utilized for research, while for the latter, the effectivity of extracorporeal blood purification techniques in the context of sepsis is investigated.

Extracorporeal blood purification therapies implicate convection, adsorption, combination and further therapies, as listed in table 1 (131). Although evidence for their potential in extracting inflammatory mediators and bacterial toxins from blood has been established, there is a lack of high-quality trials in septic patients (132).

Convection techniques include HVFT (high volume hemofiltration therapy), CCRT (continuous renal replacement therapy) and HCO (high-cut-off) membrane therapy. In HVFT, an increased convective target dose of > 35 ml/kg/h is used, compared to CCRT, leading to mediator elimination and the effectivity of different convective target dosages and

application modes (continuous vs. intermittent) for certain septic patient groups have been investigated (133,134), but a comprehensive conclusion cannot be drawn for all sepsis patient groups. Similar to HVFT, the utilization of HCO membrane therapy resulted in increased removal of mediators in a septic patient subgroup, but was associated with concomitant loss of albumin (135), while further trials were not completed (132).

TPE (therapeutic plasma exchange) is thought to be beneficial not only by clearing the body from mediators and toxins, but also by providing instant replenishment of plasma molecules and the utilization is currently under investigation (131,136)

The combination of filtration and adsorption, CPFA (coupled filtration and adsorption) did not show any benefit in clinical trials (131,137,138).

Adsorption techniques include hemoperfusion through PMX (polymyxin B immobilized) fiber columns, for which the data is contradictory until today (131,139–141), and Hemadsorption using CytoSorb, which is able to absorb many cytokines and mediators, but also free Hb (Hemoglobin), Myoglobin, Bilirubin, Bile acids, bacterial toxins, activated components of the complement system and drugs, which might be especially beneficial for septic patients when utilized in early phase (131,142–144). Recent studies using CytoSorb in patients with sepsis and/or septic shock suggest a decreased vasopressor requirement, increased lactate clearance and the resolution of septic shock (145,146) and an improved 28-day mortality rate (147), making CytoSorb the most promising blood purification technique up to this date.

| | |
|-----------------------|---|
| Convection Therapies | CCRT – Continuous Renal Replacement Therapy HVFT – High Volume Hemofiltration Therapy HCO – High-cut-off Membrane Therapy |
| Adsorption Therapies | PMX – Immobilized Polymyxin B CytoSorb - Hemoadsorption |
| Combination Therapies | CPFA – Coupled Plasma Filtration Adsorption CFA - Combined Filtration and Adsorption |
| Further Therapies | TPE- Therapeutic Plasma Exchange RAD - Renal Assist Device SCD – Selective Cytopheretic Device |

Table 1 – Overview of blood purification techniques

Graphic adapted from: Jarczak D, Kluge S, Nierhaus A. Sepsis-Pathophysiology and Therapeutic Concepts. Front Med (Lausanne). 2021;8:628302

Immunomodulatory treatment targeting innate and adaptive immune cell function and cytokine production may aid in restoration of immune homeostasis in a septic context and is currently highly investigated. Up until today there is not much clear evidence of beneficence, mainly due to the lack of clinical translation of in-vivo or animal models, uncompleted trials, or negative trial results.

The previously described down-regulation of immune cell function by upregulation of PD-1 and PD-L1, representing a “checkpoint” in the negative costimulatory pathway of a normal T-cell response, lead to the investigation of monoclonal antibodies blocking either PD-1 or PD-L1, thus inhibiting the binding ability, which lead to an unwanted increase in cytokine production and secretion by T cells and monocytes in ex-vivo studies (148). A phase I trial using Nivolumab, a monoclonal antibody targeting PD-1, in 38 sepsis and/or septic shock patients resulted in no increase in inflammatory cytokines and did not result in unexpected safety findings and stimulated INF- γ , an anti-inflammatory cytokine, production by T cells (131,149), making Nivolumab a potential candidate to target immune cell function without increasing cytokine production, yet further trials on larger patient groups are needed.

IL-7, also called the “maestro of the immune system”, has an important role in the development, maturation, expansion and homeostasis of B and T cells (150) with studies demonstrating an increase in absolute lymphocyte count and the number of circulating CD4+ and CD8+ T cells without increased pro-inflammatory response or organ function deterioration and increased INF- γ production by T-cells upon treatment with recombinant human IL-7 in patient with septic shock and severe lymphopenia (131,151), making recombinant human IL-7 also a potential therapeutic candidate.

Polyvalent immunoglobins have been shown to be able to neutralize toxins, interact with complement factors and increase opsonization of pathogens leading to increased phagocytosis (131,152–155). Until today no immunoglobulin therapy is recommended for treatment in septic patients due to lack of definitive data due to inconsistency of study protocols, heterogeneity of patients and alterations of laboratory parameter spectrums, but more RCT’s (randomized controlled trials) are on the way to further clarify utility (131,156).

12. Conclusion

Many advances in the understanding of septic pathophysiology have been made and are continued to be made, hence allowing the identification of the main therapeutic targets in sepsis treatment. Beside source-control and organ support, the immunologic nature of sepsis needs to be taken into consideration, namely regulation of the cytokine production and secretion, as well, as immune cell function. Many molecular pathways are targeted in research in vitro and in vivo, with some having promising results. Nevertheless, more research is needed to provide a comprehensive therapy for patients in sepsis and septic shock.

13. List of Abbreviations

List of Abbreviations

SIRS – systemic inflammatory response syndrome
SOFA – sepsis-related organ failure assessment
MODS – multiple organ dysfunction syndrome
MAP – mean arterial pressure
NCBI – national center for biotechnology and information
SNP – single nucleotide polymorphisms
CARS – compensatory anti-inflammatory response syndrome
PAMP – pathogen-associated molecular pattern
DAMP – damage-associated molecular pattern
PRR – pathogen recognition receptor
TLR – toll-like receptor
IL– interleukin
NK-cell - natural killer cell
RA- receptor antagonist
DNA – deoxyribonucleic acid
NFkB – nuclear transcription factor kappa B
AP – activator protein
PPAR – peroxisome proliferator-activated receptors
COX – cyclooxygenase
iNOS – inducible nitric oxide synthase
PG – peptidoglycan
LPS – lipopolysaccharide
PMN – peripheral mononuclear cells
APACHE – acute physiology and chronic health evaluation
EMA – European medical association
P – protein
INF – interferon
TNF – tumor necrosis factor
ICAM – intracellular adhesion molecule
VCAM – vascular adhesion molecule

MIP – macrophage inflammatory protein
MCP – monocyte chemoattractant protein
CINC - cytokine-induced neutrophil chemoattractant
ROS – reactive oxygen species
RNS – reactive nitrogen species
LO – lipoxygenase
PG – prostaglandin
FLAP – 5-LO-activating protein
LT – leukotrienes
TBX – thromboxane
NO – nitric oxide
VSMC – vascular smooth muscle
TF – tissue factor
PAI – platelet activator inhibitor
DIC – disseminated intravascular coagulation
PT – prothrombin time
ATP – adenosine triphosphate
ADP – adenosine diphosphate
Pi – inorganic phosphate
OXPHOS – oxidative phosphorylation
NADH/NAD⁺ – nicotinamide adenine dinucleotide
FADH – flavin adenine dinucleotide
CoA – coenzyme A
O₂ – oxygen
pO₂ – partial oxygen pressure
ONOO⁻ – peroxy nitrite
PARP – poly (ADP-ribose) polymerase
VO₂ – Volume of oxygen
mHLA-DR – monocyte human leukocyte antigen DR isotype
PD – programmed cell death protein
PD-L – programmed cell death protein ligand
Treg – T regulatory lymphocytes

TGF- β – transforming growth factor beta

IL-RA- interleukin receptor antagonist

HVFT – high volume hemofiltration therapy

CCRT – continuous renal replacement therapy

HCO – high-cut-off

TPE – therapeutic plasma exchange

CPFA – coupled filtration and adsorption

PMX – polymyxin B immobilized

Hb – hemoglobin

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