

From amino acids polymers, antimicrobial peptides, and histones, to their possible role in the pathogenesis of septic shock: a historical perspective

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Abstract: This paper describes the evolution of our understanding of the biological role played by synthetic and natural antimicrobial cationic peptides and by the highly basic nuclear histones as modulators of infection, postinfectious sequelae, trauma, and coagulation phenomena. The authors discuss the effects of the synthetic polymers of basic poly α amino acids, poly L-lysine, and poly L-arginine on blood coagulation, fibrinolysis, bacterial killing, and blood vessels; the properties of natural and synthetic antimicrobial cationic peptides as potential replacements or adjuncts to antibiotics; polycations as opsonizing agents promoting endocytosis/phagocytosis; polycations and muramidases as activators of autolytic wall enzymes in bacteria, causing bacteriolysis and tissue damage; and polycations and nuclear histones as potential virulence factors and as markers of sepsis, septic shock, disseminated intravascular coagulopathy, acute lung injury, pancreatitis, trauma, and other additional clinical disorders

Keywords: histones, sepsis, septic shock

Introduction

The pioneering work of Anton van Leeuwenhoek, Edward Jenner, Ignaz Semmelweis, Louis Pasteur, Robert Koch, Elia Metchnikoff, and many others linked microbes with infectious diseases and helped establish the germ theory of disease.¹ Robert Koch's postulates and Metchnikoff's phagocytosis theory described various functions of macrophages and their ability to kill microorganisms. This formed the basis for numerous studies on the biochemical properties and role in infection and immunity of phagocytic cells such as neutrophils and macrophages. These phagocytic cells are rich in antimicrobial peptides, lysosomal hydrolases, and oxidants. These early theories also lead to the discovery of proinflammatory cytokines and their role in microbial infections. However, the early discovery of lysozyme in 1923, and the further discovery of penicillin by Fleming, Florey, and Chane in 1928, was a major step in attempting to control severe microbial infections that may result in septic shock and organ failure; these conditions have a very high mortality even today. The recent concern regarding acquisition of antibiotic resistance by the microbe and the ongoing risk to life of the postinfectious sequelae prompted a very intense search for alternative antimicrobial cationic peptides to hopefully cope with severe microbial infections and their aftermath.

The basic concepts of microbial virulence were skillfully reviewed by Casadevall and Pirofski.² They stated that "while the importance of a host's susceptibility for

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a microbe's virulence was often recognized, the existing definitions did not account for the contributions of both pathogen and host." As we know today, sepsis is indeed the host response (organ dysfunction) to invasion by pathogenic microorganisms (infection), while septic shock is the more extreme form of this host reaction where vasodilation and translocation of fluid from the vascular space to the interstitial space causes hypotension and other cellular and metabolic abnormalities. Casadevall and Pirofski² reviewed historical concepts of microbial pathogenicity and virulence, proposed new definitions, and suggested a classification system for microbial pathogens based on their ability to cause damage as a function of the host's immune response. One typical example is septic shock (see section on "Can histone released from neutrophils" section).

This review will address the nature and role of histones as well as other modulators of infection and the host response to infection. Histones may be defined as highly basic proteins around which DNA coils to form chromatin. Their role in infection is described further.

The effect of the synthetic polymers of basic of poly α amino acids poly L-lysine and poly L-arginine on blood coagulation, fibrinolysis, bacterial killing, and blood vessels

Even as early as 1952–1956, teams of investigators at the Weitzman Institute of Science in Rehovot, Israel, headed by Ephraim Katchalski, were the first to synthesize and investigate the role of the linear polymers of basic amino acids such as poly L-lysine, poly L-arginine, and poly L-ornithine in the retardation of blood coagulation and fibrinolysis, killing of bacteria and mammalian cells, promotion of phagocytosis, and toxicity to blood vessels.^{3–9} It is of note that synthetic cationic polymers are actually histone mimics, sharing a high cationic charge capable of interaction with anionic agents. The researchers at Weizmann Institute also successfully explored the use of such poly amino acids as protein models and studied many of their physical, chemical, and biological properties. In all cases, the effects of the cationic polymers were abrogated by poly anions such as poly L-aspartic acid, poly L-glutamic acid, and the highly sulfated compound heparin.^{3–9} Unfortunately, these pioneering "ancient" studies are hardly ever cited in the modern literature and may be lost to the clinicians forever.

The retardation of clot lysis by basic poly cations was reconfirmed in 2015.^{10,11} In sepsis, both coagulation (activation of the coagulation cascade) as well as fibrinolysis are

enhanced. Therapeutic strategies (eg, activated protein C) have previously been directed at reducing clot formation in the microcirculation to reduce organ dysfunction in septic patients. The authors showed that histone mimics (poly L-lysine and poly L-arginine) and neutrophil extracellular traps exerted antifibrinolytic effects in a plasma environment and that the combination of histones and DNA also significantly prolonged clot lysis by forming thicker fibers accompanied by improved stability and rigidity.^{10–12}

The properties of natural and synthetic antimicrobial cationic peptides designed to replace antibiotics

In 1956 and later on in 1958 and in 1960, poly L-lysine was shown to possess potent bactericidal effects against a variety of microorganisms and also against certain viruses, all abrogated by poly-anions such as heparin poly-glutamic and poly-aspartic acids.^{9,13,14} However, being toxic to mammalian cells, their clinical use should be considered with caution.¹⁵

Antimicrobial peptides (AMP) are mainly small peptides (12–50 amino acids) containing a positive charge and an amphipathic structure. The AMPs, which are rich in proline, tryptophan, arginine, lysine, or histidine, are actually mimics of nuclear cationic histones (see section on "Can histone released from neutrophils") and are able to interact with negatively charged microbial and mammalian membranes to disrupt the bilayer curvature, beyond a threshold concentration of membrane-bound peptide. In bacteria, AMPs rapidly interact with surface lipopolysaccharide (LPS) of Gram-negative organisms and with the membrane-associated lipoteichoic acid (LTA) in Gram-positive organisms, and they also demonstrate toxicity to a variety of mammalian cells. AMPs may also induce bacteriolysis.

Since 1956, an overabundance of publications focused on the chemistry, physics, biology, and bactericidal effects of a large variety of linear and nonlinear cationic AMPs.^{16–22} These cationic agents may be considered evolutionarily ancient weapons against microbial infections. They may also play a pivotal role in innate immunity and as agents for specific uses because of their natural antimicrobial properties and a low propensity for the development of bacterial resistance. Hopefully, one day, AMPs may provide an alternative to conventional antibiotics (discussed later).

Readers who require further background on this topic would be well-served spending time reading the works of and paying tribute to the pioneers: M Zasloff, R Hancock, K Brogden, Y Shai, T Ganz, A Peschel, P Elsbach, Robert I

Lehrer, JK Spitznagel, and many others, who were the ones to delve into the complicated chemistry, physics, and biological effects and possible future medical uses of AMPs.

Polycations as opsonizing agents promoting endocytosis/phagocytosis

In general, polycationic agents can interact via electrostatic forces with negatively charged sites, mainly on the surfaces of microbial and mammalian cells. Such interactions may perturb the membrane, induce cell agglutination, and also cause permeability changes that may lead to cell lysis.²³ The attachment of cationic agents to surfaces of negatively charged particles is called opsonization and is similar to the effect of antibodies. Both facilitate the internalization (phagocytosis) of cationic particles by the professional phagocytic cells, neutrophils (PMNs), and macrophages, but surprisingly, also by nonprofessional phagocytes and by certain tumor cells. Although we will not fully cover the topic, it is worth mentioning that a variety of positively charged agents have also been shown to act as transfecting agents and as agents promoting the delivery of drugs as conjugates and as “decorators” of drug-loaded carriers.²⁴

It is important to note that nonspecific plasma globulins and IgGs are positively charged macromolecules. However, nonspecific cationic globulins, which can bind to cell surfaces by electrostatic forces, might interfere with the binding of specific antibodies. Indeed, a thermostable cytotoxic factor, globulin, in normal human plasma inhibited the action of heterologous antibodies on HeLa cells.²⁵

In 1986, it was shown that *Entamoeba histolytica* and *Acanthamoeba palestinensis*, two distinct classical phagocytic cells (possibly evolutionary forefathers of neutrophils and macrophages), which stubbornly refused to internalize/engulf *Candida albicans*, nevertheless did so very avidly if pre-coated by arginine-rich polycations.²⁶ These studies resulted in later experiments that showed phagocytosis–endocytosis of *Candida albicans* and of Group A *Streptococci* by mouse fibroblasts and by epithelial cells in culture.²⁷ In this study, the most potent opsonins for Group A *Streptococci* were specific antibodies supplemented with complement, nuclear histone, poly lysine, poly arginine, ribonuclease, leukocyte lysates, leukocyte cationic proteins, and, to a lesser extent, cationic lysozyme and myeloperoxidase.

Highly cationic histone, RNase, leukocyte extracts, and platelet extracts also functioned as opsonins for phagocytosis of streptococci in the peritoneal cavity.²⁷ However, the phagocytic capabilities of mouse fibroblast poly karyons (cells with multiple nuclei) were much higher than those of ordinary spindle-shaped fibroblasts, probably due to their very large

cytoplasmic area. Calf thymus histone also functioned as a good opsonic agent for the uptake of *Candida* by human fibroblasts, HeLa cells, epithelial cells, monkey kidney cells, and rat heart cells in culture.²⁷ Phagocytosis of *Streptococci* and *Candida* by macrophages and the uptake of *Candida* by fibroblasts were both strongly inhibited by the polyanions hyaluronic acid, DNA, and dextran sulfate. The paucity of nonprofessional phagocytes of hydrolases capable of breaking down microbial cell wall components may contribute to the persistence of nonbiodegradable components of bacteria in tissues and lead to the perpetuation of chronic inflammatory sequelae such as granulomatosis.²⁸

Two excellent, but concerning, examples of phagocytosis of microbes in vivo showed that *Staphylococcus aureus*, by forming microcolonies, could survive unharmed within skin keratinocytes, waiting for the opportunity to attack patients with low immunity. The mechanism of cell uptake was not disclosed.^{29,30}

It was also demonstrated that nonbiodegradable cell wall components could persist for long periods within macrophages in arthritic granulomas. Another example is the chronicity of lesions in tuberculosis.^{31–33} Furthermore, macrophages and neutrophils loaded with opsonized streptococcal cell walls can be translocated to remote sites to induce chronic inflammation.^{34–36}

Polycations and lysozyme as activators of autolytic wall enzymes (muramidases) in bacteria, causing bacteriolysis and tissue damage

The discovery of the bacteriolysis phenomenon dates back to 1893^{32,37} when Buchner³⁷ reported that fresh serum was able to kill certain bacteria, an effect which was lost upon heating to 55°C. He attributed the bactericidal action of serum to a heat-labile constituent that he called “alexine” (from Greek “to ward off”). One year later, Pfeiffer described the dissolution of *Vibrio cholera* by fresh serum of guinea pigs immunized with heated vaccine, which could be correlated with protection against infection in both passively and actively immunized animals.³²

However, the significance of the biochemical degradation of microbes as related to tissue injury in inflammation, infection, and postinfectious sequelae has emerged mainly from a large series of investigations^{32,38,39} that focused on:

1. The structure and function of the bacterial cell walls
2. The role of muramidases (autolytic wall enzymes) in normal bacterial multiplication

3. The role played by lysozyme, leukocyte-derived poly-cations, cationic enzymes, and antibiotics (mostly β -lactams) in bacteriolysis
4. The role of muramidase-deficient strains in pathology
5. Antibiotic resistance
6. Microbial killing and degradation:

The role of the cell wall components: LPS, LTA, and peptidoglycan (PPG) in the activation of leukocytes and in the generation of oxidants, proteinases, and cytotoxic cytokines.

7. The role of microbial cell wall components in the pathogenesis of granulomatous inflammation and in the potentiation of innate immunity to infections and of tumor-cell proliferation

Morphologically, two main patterns of bacterial cell degradation under various physiological and pathological conditions have been defined:

1. The term *plasmolysis* was proposed when a significant degradation of cytoplasmic constituents occurred, leaving apparently intact cell walls
2. The term *bacteriolysis* was proposed when a significant breakdown and degradation of the rigid cell walls, presumably due to the uncontrolled activation of autolytic wall enzymes (muramidases), occurred.

Bacteriolysis can be defined as an event that may occur when normal microbial multiplication is altered due to an uncontrolled activation of a series of autolytic cell-wall breaking enzymes (muramidases). It may happen following treatment of bacteria by β -lactam antibiotics or also by a large variety of bacteriolysis-inducing cationic peptides such as histones, elastase and cathepsin G, lysozyme, and PLA₂. When bacteriolysis occurs in vivo, cell wall- and membrane-associated LPS (endotoxin) from Gram-negative organisms and LTA and PPG from Gram-positive organisms are released. These highly phlogistic agents can act on macrophages to induce the generation and release of reactive oxygen and nitrogen species, cytotoxic cytokines, hydrolases, proteinases, and also activate the coagulation and complement cascades.⁴⁰

Peptidoglycan hydrolysis can result in the rupture of the murein sacculus due to its high osmotic pressure, leading to the release of cytoplasmic constituents and cell wall fragments.³² A possible explanation for the long persistence of highly phlogistic nonbiodegradable microbial cell wall remnants within professional phagocytic cells was offered in 1989.⁴¹

It was proposed that following phagocytosis either by PMNs or by macrophages, the engulfed microorganisms are exposed intraphagosomally to the respiratory burst generating oxidants, LL-37, lysosomal cationic proteinases, and also numerous hydrolases, which inactivate the autolytic wall enzymes thus allowing the survival of highly phlogistic microbial cell wall component.

It was also shown that neutrophil-mediated myeloperoxidase, H₂O₂, and HOCl production inactivated a class of cytoplasmic membrane enzymes (penicillin-binding proteins [PBP's]) in *Escherichia coli*, *Staphylococcus aureus*, and *Pseudomonas aeruginosa*. These PBP's covalently bind β -lactam antibiotics to their active sites. This contributed to the persistence of nondegraded microbial components, leading to unbalanced bacterial growth and cell death.^{42,43}

Degradation of *Staphylococcus aureus* by β -lactams was markedly inhibited by the polyanions suramine and Evans blue, suggesting that accumulation of polyanions and sulfated polysaccharides in inflammatory sites might also interfere with bacteriolysis.^{44,45}

Clindamycin treatment of *Staphylococcus aureus* caused a remarkable thickening of the bacterial cell wall due to increased numbers of O-acetyl groups in the murein, which made the bacterial wall much more resistant to lytic enzymes within bone marrow-derived macrophages, and this was revealed by electron microscopy and radiolabeling experiments. This reduced wall degradation might increase the survival of highly phlogistic walls in inflammatory sites. Furthermore, such clindamycin-treated bacteria were ingested by adherent bone marrow-derived macrophages at a higher rate than untreated bacteria.⁴⁶ The involvement of bacteriolysis in sepsis was also reported.⁴⁷

The lysozyme riddle: is this enzyme a genuine and an effective bacteriolytic enzyme?

In 1922, Alexander Fleming discovered the enzyme lysozyme (*N*-acetylmuramide glycanhydrolase).⁴⁸ Lysozyme is a 139-amino acid cationic protein found in neutrophils, macrophages, saliva, mucous, egg white, milk, and additional body fluids. Patients with myeloid leukemia can be diagnosed by measuring lysozyme in urine by a simple method using suspensions of *Micrococcus lysodeikticus* as a highly sensitive substrate (Ginsburg, unpublished data). Lysozyme was anticipated to kill, lyse, and biodegrade pathogenic microorganisms.

Lysozymecan very rapidly (within 1–2 minutes) lyse certain nonpathogenic Gram-positive cocci (eg, *Micrococcus*

lysodeikticus and also spore-bearing aerobic bacilli) possessing a “simple” PPG. However lysis of *Staphylococcus aureus*, which possesses a more complex PPG, can take up to 6–12 hours. It is of clinical significance that lysozyme rarely induces lysis either of hemolytic *Streptococci*, *Streptococcus viridans*, *Listeriae*, *Mycobacteria*, or *Candida* species, and it lyses enteric bacteria *Staphylococcus aureus* slowly. It could, however be partially lysed by a synergism between lysozyme, lysolecithin and phospholipase C.³² Also, hemolytic streptococci cultivated in the presence of subinhibitory concentrations of penicillin lost their membrane-associated phospholipids to a large extent following treatment with small concentrations of lysothelin and lysozyme.³²

The main reason for the relative resistance to lysozyme action of *Staphylococcus aureus* and perhaps also of the majority of pathogenic microorganisms, may be ascribed to the presence in their peptidoglycans of O-acetyl groups, which hinder the interaction of lysozyme with the N-acetylglucosamine-N-acetyl muramic acid linkages in the PPG.^{32,49,50} However, mild alkaline solutions rendered such cell walls digestible by egg-white lysozyme.⁵¹

The lack of deacetylating enzymes in phagocytes may explain why apparently intact *Staphylococcal*, *Streptococcal*, and *Mycobacterial* cell walls may persist for long periods either within phagolysosomes of macrophages in culture or also in vivo. However, lysozyme does not seem to function as a muramidase but rather as a cationic peptide that activates the microbial autolytic wall enzymes in certain bacteria.³² Usually, this process takes hours and is therefore missed when the bactericidal effects of AMP are tested for very short periods and expressed as colony forming units.^{49–52}

Lysis of bacteria by antibiotics in vivo may also be involved in sepsis and septic shock (see section on “Can histone released from neutrophils”⁴⁷).

Synergistic effects among cationic peptides

It was also demonstrated that mammalian cationic peptides from different structural classes (eg, α -helical cationic peptides such as lactoferrin and most amphipathic membrane-active AMPs) frequently show synergy with each other and also with lysozyme. It is assumed that this reflects the cooperative interactions of the peptides with the outer membranes of Gram-negative bacteria and/or cooperative interaction with lipid bilayers in general. It was concluded that, given the substantial diversity of peptides in any given location in the host, synergistic interactions are an important determinants of the overall effectiveness of the peptides.⁵² Cationic peptides

and cationic proteins can also act in synergy with reactive oxygen species to injure mammalian cells.^{53–56}

Can histone released from neutrophil nets function as a major virulence factor involved in the pathophysiology of septic shock trauma and also in many additional clinical disorders?

It is alarming that today clinicians are still limited when trying to treat the life-threatening sequelae of severe microbial infections, which very often lead to sepsis and septic shock, both of which have a high mortality.^{57,58}

The annual incidence of sepsis in the USA has been estimated to affect as many as 750,000 hospitalized patients with mortality reaching about 40%.^{57,58} Worldwide, sepsis is one of the commonest, deadliest disease entities, and globally, 20 to 30 million patients are estimated to be afflicted every year with what is one of the least well-understood disorders.

Screening the voluminous literature on sepsis treatments reveals the repeated unsuccessful efforts to save patients’ lives by administering antibiotics, sometimes combined with only singly-selected antagonists. The numbers of unsuccessful antiseptic agents that have been tested in clinical trials in the last 30 years is phenomenal, and today, even the most promising agent, activated protein C, has been recently removed from use. Today, there is no specific effective treatment for sepsis and septic shock.^{57,58}

The pioneering studies on poly alpha cationic amino acids^{3–9} and their role as bactericidal agents, as opsonins, and as bacteriolysis-inducing agents, (see the earlier sections) raised interest regarding the possible role of histones and modified histones, actually lysine and arginine-rich peptides, in the pathophysiology of a variety of clinical disorders.

This “new field” of research emerged in 2009 from two “breakthrough” articles by Xu et al⁵⁹ and Chaput et al⁶⁰ in Nature Medicine. These authors had proposed that nuclear histones released from PMN nets may be the main cause of death in sepsis and that this is due to the toxicity of the highly cationic protein to endothelial cells (ECs).

Histones comprise five groups of nuclear proteins rich in the highly basic amino acids L-lysine and L-arginine, which are bound to chromatin in the cell nucleus. Extracellular histones are highly toxic to bacteria and to mammalian cells and can increase plasma thrombin generation by impairing endothelial thrombomodulin-dependent protein C activation, which is responsible for disseminated intravascular

coagulopathy (DIC). Dysregulation of ECs by the released histones leads to a severe immune cytokine storm and coagulation cascade. The toxic effects of histones could be abrogated or slowed down by antibodies to histone-activated protein C (a protease which cleaves histone) and also by heparin. Extracellular histones are also elevated in response to traumatic injury, and this elevation correlates with fibrinolysis^{6,10-12} and activation of anticoagulants. In trauma patients, an increase in histone levels between the time of admission and 6 hours is predictive of mortality. This suggests a possible role for activated protein C in mitigating the sterile inflammatory response after trauma through the proteolysis of circulating histones. However, the question of whether histones alone are the real culprits or just markers of cell damage is still unsettled.

Septic shock was recently redefined as a multifactorial synergistic phenomenon/disorder. While no distinct virulence factor (alarmin) has been identified, if successfully neutralized, the devastating immune responses can be slowed down or even stopped, which will hopefully reduce mortality.^{61,62}

The possible role of histones in additional clinical disorders

The interesting publications regarding the possible pathogenic properties of histones in sepsis has resulted in a plethora of studies, also suggesting that these circulating polycations may be involved in the pathogenesis of DIC, acute lung injury, trauma, pancreatitis, liver, renal and myocardial disorders, trauma, heat stroke, and many other clinical disorders.⁶³⁻⁷⁹ Analyzing these articles, one wonders if the presence of histones in the circulation indicates that these are innate virulence/toxic agents or just an additional marker of tissue damage.

Can histones function alone in vivo as absolute virulence factors?

It was previously demonstrated that the toxicity of histone to ECs and to epithelial cells in culture was markedly further enhanced (in a synergistic manner) in combination with oxidants, proteinases, and additional proinflammatory agents generated by activated neutrophils.^{54,55,62,80,81} Such synergistic phenomenon might actually be a general mechanism of cell injury mediated by activated phagocytes recruited to infectious and inflammatory sites.

The following scenario might be depicted: following adherence to ECs, PMNs undergo NETosis, and the released DNA combined with histones is accompanied by activation of NADPH oxidase and generation of reactive oxygen species.

This happens concomitantly with activation of a large array of proinflammatory agents.

Dysregulation of ECs leads to platelets activation and the generation of cytokine storms and coagulation cascades.

However, it is highly plausible that in vivo, histones and additional toxic polycations (eg, LL-37, elastase) most probably never act on their own but always in synergism with many additional proinflammatory agents.^{54,55,81}

Since histones' action could be abrogated either by antibodies to histone, activated protein C, anionic heparin, and also additional polysulfates, it is reasonable to assume that these inhibitors actually affect not only histones action alone but also the synergism among the various agents. Therefore, if these inhibitors are administered early enough, they could still manage to neutralize the toxic effects to prevent the ongoing deleterious immune and coagulation responses.

Perhaps we could also use antioxidants,⁸² since NETs and the release of histones (from activated PMNs adhering to endothelial cells) are accompanied by the activation of the respiratory burst in PMNs.⁸²

Years ago it was suggested that a multi-faceted approach to sepsis was required and not treatment based on a single antagonist, this paper was largely ignored.⁸³

Why is the mortality from sepsis still so high?

Current clinical management of sepsis

Currently, most efforts in the clinical management of septic patients are directed at early recognition and diagnosis, prompt commencement of treatment, source control of sepsis, and early antibiotic therapy. Following this initial therapy, the rest of the therapeutic armamentarium is based on supportive treatment such as optimal fluid therapy, vasopressor and inotropic therapy, and organ support (eg, mechanical ventilation and renal replacement therapy). In light of our growing understanding of the underlying mechanisms of sepsis and the host response to sepsis, these treatments are directed at "downstream" processes that occur long after the initial injury.

Usually, in clinical practice, sepsis patients showing the main symptoms of tachypnea, tachycardia, confusion, high lactate and procalcitonin levels, leukocytosis or leukopenia arrive in the intensive care unit (ICU) many hours or even days, after developing symptoms. Therefore, even the "miracle" novel nonanticoagulant heparin^{81,84} combined with antibiotics might already be ineffective since by that time all the pathological biochemical processes are already well established. As for the nature of additional sepsis

markers, which may be helpful in the early diagnosis of sepsis, the reader is directed to excellent review articles on the subject.^{61,82,85–89}

Finally, taken together, it is not understood why the latest consensus definition of clinical of sepsis 2016⁹⁰ has not considered and discussed either the possible involvement in sepsis of histones and additional cationic peptides, the use of nonanticoagulant heparin,⁸⁴ or the role of bacteriolysis in pathogenicity.

It seems that antihistone measures as a plausible and accepted therapy (see section on “Can histone released from neutrophils”) have not yet matured sufficiently to reach practicing clinicians.

It is hoped that further studies on the pathophysiology of sepsis might shed more light on the possible validity of combinations of nonanticoagulant heparin, anti-inflammatory cocktails, anti-cytokines, antioxidants, and anti-bacteriolysis agents in the treatment of sepsis. To do so effectively, we have to define and employ very early markers of sepsis.^{81–85}

Looking to the future

Consider a still hypothetical idea that every office of a public/private practitioner (a house doctor) may have available a simple, inexpensive kit to detect early sepsis makers to identify abnormal levels of biochemical, blood, and leukocyte parameters in urine, blood, or other biological fluid. These will replace the more cumbersome and expensive ELISA kits that, today, are usually available in research laboratories but not in the ICU. Thus, sepsis diagnosis will be much faster and more responsive to adequate effective treatment.

Summary

Taken together, our understanding of the possible role of highly charged polymers of basic amino acid in the pathophysiology of infection, in postinfectious and inflammatory sequelae, and following trauma and inflammation has evolved over more than 60 years and is still going on. Septic shock and posttrauma syndromes are considered synergistic multifactorial disorders where not a single virulence factor had been identified, which, if successfully inhibited, might delay or stop the immunological and coagulation cascades leading to a patient's demise. Circulating histones are also tied up with the pathogenesis of pulmonary, renal, cardiac, pancreatic, and liver disorders, as well as in other disorders. Today, we still do not fully know whether histones and additional cationic peptides released into the circulation are major virulence factors or just biomarkers of tissue damage.

Whatever the reasons are for the pathogenicity of poly-cations, the newly described nonanticoagulant heparin, if combined in time with antibodies to histones, activated protein C (a protease, which cleaves histones), nonbacteriolytic antibiotics, antioxidants, steroids, and cocktails of additional antagonists, may be justified for use as a therapeutic regimen. This may finally bring to an end the numerous unsuccessful trials of sepsis conducted, by the administration of single antagonists, over so many years in an attempt to cope with the patient's morbidity and unfortunate mortality.⁷⁹ However, we should also consider the fact that since patients suspected of developing septic shock may usually arrive at the ICU hours or days after the appearance of symptoms, even antihistone strategies may not be fully protective by that time. Therefore, efforts should be made to identify novel, very early markers of tissue damage to allow early treatment.

The complexity of the sepsis syndrome, which involves multiple interactions among biochemical, immunological, and coagulation cascades, and the difficulty in identifying the disorders early enough are still the main stumbling blocks to achieve a consensus of how to prevent and treat the post infectious sequelae of sepsis and posttrauma syndromes.

Disclosure

Erez Koren is currently employed at Teva Pharmaceuticals Ltd, Israel. The authors report no other conflict of interests in this work.

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