

REVIEW

Bench-to-bedside review: Sepsis - from the redox point of view

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Abstract

The pathogenesis of sepsis and its progression to multiple organ dysfunction syndrome and septic shock have been the subject of investigations for nearly half a century. Controversies still exist with regard to understanding the molecular pathophysiology of sepsis in relation to the complex roles played by reactive oxygen species, nitric oxide, complements and cytokines. In the present review we categorise the key turning points in sepsis development and outline the most probable sequence of events leading to cellular dysfunction and organ failure under septic conditions. We have applied an integrative approach in order to fuse current state-of-the-art knowledge about redox processes involving hydrogen peroxide, nitric oxide, superoxide, peroxynitrite and hydroxyl radical, which lead to mitochondrial respiratory dysfunction. Finally, from this point of view, the potential of redox therapy targeting sepsis is discussed.

Introduction

Sepsis is a leading cause of death in critically ill patients; the mortality in patients admitted to intensive care units and given both antibiotics and supportive care varies from 20% up to more than 60% [1,2]. The problem is becoming even more prominent because the frequency of sepsis is increasing due to the prevalence of antibiotic-resistant pathogens [3]. Historically the widely accepted hypothesis was that sepsis was in fact an uncontrolled inflammatory response. Accordingly, the organism is killing itself while the pathogens are considered as bystanders [4,5]. However, this theory was based on animal studies that do not reflect the clinical picture in humans [6]. Studies on animal models usually involve the application of large doses of endotoxin or pathogens, leading to much higher levels of circulating cytokines in

comparison to humans with sepsis and ending in death by 'cytokine storm' [6]. Recent data indicate that the immune system during sepsis actually shows two phases: an initial hyper-inflammatory stage followed by a prolonged hypo-inflammatory phase [7]. In addition, numerous studies and trials have shown that the application of anti-inflammatory agents is futile, or can even worsen chances of survival [7,8]. There are some exceptions, however. For example, Annane and co-workers [9] have shown that the application of selected corticosteroids at moderate doses may decrease the mortality of sepsis. In contrast to prevalently disappointing results for anti-inflammatory therapies, a pro-inflammatory approach seems to be more promising. For example, interferon- γ , a potent macrophage activator, improved survival of patients with sepsis [10]. IL-12, an immune stimulant, reduced mortality from subsequent sepsis when administered after burn injury [11]. Pertinent to these data, Hotchkiss and Karl [7] have recently proposed a coherent immunological approach in sepsis treatment, consisting of moderate suppression of the immune system during the hyper-inflammatory mode in early sepsis and intensive promotion of activity of the immune system during the hypo-inflammatory phase.

Immunohistochemical analysis has shown that in the majority of patients with sepsis, only lymphocytes and gastrointestinal epithelial cells die [12]. These two types of cells normally undergo rapid turnover through apoptosis, and sepsis most likely accelerates this process. Cell death in the heart, lungs and kidneys has been observed to be minor, resulting in insignificant damage to these organs [13]. Studies in patients with acute renal failure have illustrated discordance between the level of renal dysfunction and the degree of tubular necrosis [14]. It is very interesting that most sepsis survivors who showed renal failure recover baseline renal function [15], suggesting that changes that may cause death are in fact reversible. It can be concluded that organ failure in sepsis is not provoked by tissue damage, but instead by the dysfunction of a substantial number of cells. A central paradigm in organ dysfunction is the concept of tissue hypoxia; the failure to adequately supply organs with oxygen [16]. However, non-survivors of sepsis exhibit an

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impaired capacity to increase tissue oxygen consumption in response to the augmentation of oxygen delivery [17]. This is reinforced by the fact that well-perfused endotoxaemic model animals still present acidosis [18]. In line with these facts, Fink [19] has postulated that impaired cellular O₂ utilisation, rather than inadequate oxygen delivery, may play an important role in sepsis development. As mitochondria represent the main consumers of O₂, significant attention has been focused on the role of mitochondrial dysfunction in sepsis [20]. Brealey and co-workers [21] demonstrated that there is an association between mortality in sepsis and mitochondrial dysfunction. In addition, a decrease in the number of mitochondria has been observed in patients with prolonged septic conditions [22], implying that sepsis-related dysfunction of mitochondria may result in mitoptosis. The direct consequence of mitochondrial dysfunction is a significant fall in the level of ATP, as observed in the tissues of model animals and septic patients [21].

An activated immune system and dysfunctional mitochondria represent the two most potent sources of reactive molecular species in organisms [23]. Pertinent to this, it has been documented that sepsis is characterised by excessive production of reactive oxygen species (ROS) and reactive nitrogen species (RNS), both in the circulation (by activated immune system cells and endothelial cells) and in the affected organs (by dysfunctional mitochondria and due to a modified anti-oxidative status) [16,24]. Many authors agree that ROS and RNS (Table 1) play a crucial role in the pathophysiology of sepsis [24-26]. Some of these species interfere with signalling cascades, while others provoke deleterious effects on various biological molecules and structures. It is clear that the increased production of signalling species (for example, nitric oxide (•NO) and hydrogen peroxide (H₂O₂)) and strong oxidants (the hydroxyl radical (•OH) and hypochlorous acid (HOCl)) act in synergy with collapse in energy metabolism to provoke cell dysfunction, which may result in organ failure and death.

When considering possible treatment strategies for sepsis patients, it is crucial to identify the main events in the development of sepsis and the schedule of changes of pathogenic factors. In the present review we focus on redox mechanisms of sepsis without under-estimating any other possible contributors to this pathology. We outline the most probable sequence of redox events leading to cellular dysfunction in sepsis. In addition, the current potential of and prospects for the application of redox therapy in sepsis are discussed.

The chain of redox events

Oxidative burst

A plasma oxidative burst takes place early in sepsis orchestrated by neutrophils, macrophages and endothelial cells

Table 1. The main reactive species implicated in the pathogenesis of sepsis

Free radicals	Non-radicals
Reactive oxygen species	
Superoxide, •O ₂ ⁻	Hydrogen peroxide, H ₂ O ₂
Hydroxyl radical, •OH	Hypochlorous acid, HOCl
Hydroperoxyl radical, •OOH	
Peroxyl radical, •ROO	
Reactive nitrogen species	
Nitric oxide, •NO	Peroxynitrite, ONOO ⁻
Nitrogen dioxide radical, •NO ₂	Peroxynitrous acid, ONOOH

[27]. These cells represent the innate immune system, our first line of defence against invading microorganisms. Accordingly, they are equipped with different ROS and RNS [23]. When activated by pathogens they generate superoxide (•O₂⁻), H₂O₂, •NO and HOCl, while •OH, peroxynitrite (ONOO⁻), the nitrogen dioxide radical (•NO₂), the hydroperoxyl radical (•OOH), and the peroxyl radical (•ROO) are produced in secondary reactions (Figure 1). The lack of this intrinsic oxidant system can be deleterious to the host as demonstrated in subjects lacking •O₂⁻ generation. Mice deficient in NADPH oxidase subunits and humans with chronic granulomatous disease who also lack NADPH oxidase activity are incapable of adequately performing bactericidal functions [28,29]. However, the oxidant system may also significantly modify redox settings in the organism.

Septic plasma shows a number of positive markers of oxidative stress (Table 2). Anti-oxidants scavenge reactive species, which leads to a decrease in the concentration of anti-oxidants. The level of free thiols decreases due to oxidation or nitrosylation. As a result, the total anti-oxidant capacity of plasma is reduced. In addition, uncharged radicals attack membranes, initiating a lipid peroxidation chain reaction (Figure 1, reactions 6 to 8). Interestingly, two research groups have reported that the plasma total anti-oxidant capacity in sepsis is initially decreased but increases over time as the condition progresses [30,31]. This indicates that the redox status of plasma during sepsis follows a two-phase pattern characteristic of the immune system.

It is important to note that redox conditions in the plasma of neonates with sepsis seem to be different in comparison to other patient groups. Cherian and colleagues [32] reported that there were no significant changes in the levels of ascorbate, glutathione and oxidative stress biomarkers in the blood of neonates and babies with sepsis when compared to age-matched controls. This may be attributed to the immaturity of the immune system. Neonates have a low neutrophil pool and reduced release of pro-inflammatory cytokines (for

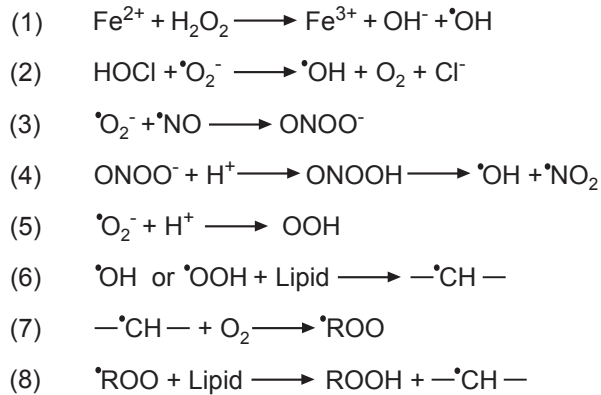


Figure 1. Key redox reactions in living systems [23]. (1) Fenton reaction; (6) to (8) lipid peroxidation (ROOH) chain reactions.

example, TNF- α), and their neutrophils have an impaired capacity to migrate to sites of infection, which may account for the less prominent oxidative burst in response to infection [33]. Fortunately, neonatal sepsis is very uncommon and shows a relatively low mortality rate of 5 to 10% (in developed countries) [34]. The presented

facts regarding neonatal sepsis strongly imply that pronounced immune system-mediated redox changes may play an important role in sepsis initiation in adults.

Modified redox status of septic plasma affects the intracellular milieu of surrounding tissues due to: (i) the capability of some reactive species, such as H_2O_2 , $\cdot\text{NO}$, and HOCl, to cross cellular membrane [23,35]; (ii) oxidative changes in the membrane (Figure 1, reactions 6 to 8) [36]; and (iii) the activation of specific receptors. After entering the cell, H_2O_2 and $\cdot\text{NO}$ participate in secondary reactions (Figure 1, reactions 1 to 4) to produce more reactive species, which provoke intracellular oxidative stress. Due to its high reactivity with biomolecules, HOCl is predicted to have a very limited diffusion radius in biological systems [37,38]. Therefore, only a small fraction of HOCl manages to enter the cell, where it may provoke protein, DNA, lipid and mitochondria damage [38]. On the other hand, H_2O_2 and $\cdot\text{NO}$ represent signalling molecules crucial for normal cell function [39]. They activate specific sets of genes via a number of transcription factors at the nuclear level, of which NF- κB appears to have a central role in sepsis [40]. Other factors, such as nuclear factor E2-related factor 2 (Nrf2), may also be

Table 2. Redox characteristics of plasma in adults with sepsis

	Patient groups		Method	Reference
Anti-oxidant				
Ascorbate (mg/L)	Control: 1.71 \pm 0.46	Sepsis: 0.55 \pm 0.28	Spectrophotometric assay	[116]
	Control: 4.5 to 12	Critically injured: 1.1 \pm 0.3		HPLC
Retinol ($\mu\text{g/L}$)	Control: 735 \pm 183	Septic shock: 265 \pm 193	HPLC	[118]
Tocopherol (mg/L)	Control: 11.5 \pm 1.3	Septic shock: 3.6 \pm 2.0	HPLC	[118]
β -Carotene ($\mu\text{g/L}$)	Control: 619 \pm 259	Septic shock: <101	HPLC	[118]
Lycopene ($\mu\text{g/L}$)	Control: 522 \pm 184	Septic shock: <154	HPLC	[118]
Oxidative stress biomarkers				
Total plasma thiols (R-SH) ($\mu\text{mol/L}$)	Control: 364 \pm 22	Sepsis: 138 \pm 14	Ellman's method	[119]
Lipid peroxidation ($\mu\text{mol/L}$)	Control: <2	Patients with >3 failing organs: 4.9 \pm 2.3	TBARS assay	[118]
	Control: 0.92 \pm 0.39	Sepsis survivors: 0.99 \pm 0.14	TBARS assay	[120]
		Sepsis non-survivors: 3.33 \pm 2.29		
Lipid peroxidation (nmol/mg of protein)	Sepsis (day 1): 2.4	Sepsis (day 7): 2.9	FOX assay	[121]
Total anti-oxidant capacity (nmol/mg of protein) ^a	Sepsis (day 1): 1.5	Sepsis (day 7): 1.1	ABTS assay	[121]
Total anti-oxidant capacity (%)	Control: 57.3 to 60.3	Survivors: 51.4 to 55.4	ABTS assay	[30]
		Non-survivors: 46.1 to 47.6		
Total anti-oxidant capacity ($\mu\text{mol/L}$)	Control: 354 \pm 9	Sepsis: 265 \pm 32	TRAP assay	[31]
		Septic shock: 452 \pm 34		

^aThe use of total anti-oxidant capacity assays is associated with some pitfalls; therefore, the results obtained with such methods should be taken with some caution. For example, urate interferes with both the ABTS and the TRAP assays, potentially leading to falsely high values [23]. Urate increases in many pathophysiological conditions, for example, in renal failure, which may develop in sepsis. Another problem is that plasma contains a complex mixture of anti-oxidants that react at different rates, so some of them may not be detected by rapid assays [23]. ABTS, 2,2'-azino-bis(3-ethylbenzthiazoline-6-sulphonic acid); FOX, ferrous oxidation-xylenol orange; HPLC, high-performance liquid chromatography; TBARS, thiobarbituric acid reactive substances; TRAP, total radical-trapping antioxidant potential.

involved. Nrf2 is a transcription factor that regulates anti-oxidant response element (ARE)-mediated transcription of a plethora of anti-oxidant and protective genes to counteract the harmful effects of ROS or environmental carcinogens. Non-carcinogenic ARE inducers from a variety of chemical classes have been identified to enhance the transcriptional activity of Nrf2 through S-alkylation of reactive cysteines within the cellular redox sensor protein Keap1 (Kelch-like ECH associated protein 1) [41]. Following dissociation from the cytosolic protein Keap1, a scaffolding protein that binds Nrf2 and Cul3 ubiquitin ligase for proteasome degradation, Nrf2 rapidly accumulates in the nucleus and transactivates the ARE in the promoter region of many anti-oxidant genes [42]. Nrf2 seems to represent an important regulator of oxidative stress in sepsis [43]. For example, it has been reported that lipopolysaccharide-induced shock and polymicrobial sepsis induces early and greater mortality in Nrf2-deficient mice, which is related to the lethal innate immune response [44]. In contrast, transgenic mice with a decreased rate of Nrf2 degradation showed lower mortality in polymicrobial sepsis [45]. Consequences of the down-regulation of the Nrf2 host defence mechanism still remain to be fully explored.

Intracellular $\cdot\text{NO}$ generation

Pro-oxidative and pro-inflammatory events in the plasma have a strong impact on the intracellular milieu. H_2O_2 causes a cascade of intracellular events resulting in the liberation of NF- κB from its inhibitor I κB [46]. NF- κB can be activated via several other pathways, as reviewed by MacDonald and co-workers [40]. For example, TNF- α , one of the primary mediators of the innate immune system, binds to its specific receptor (TNFR), which leads to the activation of multi-component protein kinase (I κB kinase), I κB phosphorylation, and NF- κB release and translocation into the nucleus [40]. Clearly NF- κB has a special place in sepsis development. In an animal study lipopolysaccharide administration resulted in NF- κB activation in several organs [47]. Two different research groups have reported that NF- κB activity is significantly higher in septic non-survivors in comparison to controls and survivors and that NF- κB activity predicts sepsis-related mortality [48,49]. Finally, encouraging results of clinical trials with activated protein C could be at least partially attributed to its ability to block the expression of NF- κB -regulated genes [50].

NF- κB binds to DNA to enable the initiation of the transcription process of a number of genes, two of them encoding redox-active enzymes: inducible $\cdot\text{NO}$ synthase (iNOS) [51] and cyclooxygenase (COX)-2, which generates $\cdot\text{O}_2^-$ as a by-product [52]. It is noteworthy that iNOS expression can be stimulated by hypoxia inducible factor-1, which is also activated by ROS [53]. In contrast to

constitutively expressed endothelial NOS, which produces local $\cdot\text{NO}$ concentrations in the nanomolar range, iNOS has been estimated to produce $\cdot\text{NO}$ in the micromolar range [54,55]. The role of iNOS expression in non-septic infections is to induce $\cdot\text{NO}$ -mediated vasodilatation, thus enabling adequate circulation in the affected areas and normal cardiac function [56]. However, over-expression of iNOS in sepsis affects normal cellular functions [57]. Pertinent to such dual (patho)physiological roles of iNOS in sepsis, it has been demonstrated in two separate studies that iNOS-deficient animals exposed to endotoxin show lower cardiac output [58] but have improved survival [59] in comparison to control subjects.

Over-expression of iNOS occurs early in sepsis [54,60], taking place in nearly all the vital organs [61]. Patients with sepsis show increased $\cdot\text{NO}$ production and elevated nitrite and nitrate levels, which have been correlated with the severity of illness [62]. The negative role of iNOS in sepsis is also indicated by the fact that the administration of the iNOS substrate arginine results in a poorer outcome in sepsis [63]. The mechanism that makes $\cdot\text{NO}$ production deleterious to cells seems to be related to protein nitrosylation and nitration, which can modulate the function of various enzymes and ion channels as well as different signalling pathways [23]. In addition, $\cdot\text{NO}$ efflux from cells and a substantial increase in the $\cdot\text{NO}$ level in plasma may occur. $\cdot\text{NO}$ converts haemoglobin to methaemoglobin (MetHb), which leads to increased rigidity of red blood cells (RBCs) and RBC lysis [64]. Another potential cause of RBC lysis in sepsis is the ability of HOCl to induce membrane-pore formation in RBCs [65]. RBC lysis results in the release of MetHb into the plasma [64], reflecting the severity of sepsis [66]. An increased level of free MetHb in the plasma may have several propagating effects on sepsis development: (i) it leads to increased availability of iron to pathogens, which may fuel their proliferation [67]; (ii) it has pro-inflammatory effects by stimulating interleukins (IL-6 and IL-8) and E-selectin [68]; and (iii) MetHb is an activator of NF- κB [64], which may lead to further iNOS expression, thus closing a $\cdot\text{NO}$ -generating loop (Figure 2).

Intracellular superoxide generation

Septic insult significantly increases intracellular $\cdot\text{O}_2^-$ [69]. This is substantiated by the high activity of superoxide dismutase (SOD) in sepsis [22,27]. There are several intracellular $\cdot\text{O}_2^-$ generators implicated in sepsis: NADPH oxidase, COX-2, xanthine oxidase and the electron transfer chain (ETC) in mitochondria. The entry of ROS from the plasma into the cell can initiate the expression of NADPH oxidase and COX-2 [40,52,70]. The early rise in $\cdot\text{O}_2^-$ production in endothelial cells in sepsis is abolished by NADPH oxidase inhibitors [69], while the inhibition of

Peroxynitrite production and its effects

The interaction between $\cdot\text{NO}$ and $\cdot\text{O}_2^-$ plays a fundamental role in cell and organ failure [78]. Superoxide reacts with $\cdot\text{NO}$ at a slightly faster rate than with SOD so when the level of $\cdot\text{NO}$ is in the high nanomolar or micromolar range $\cdot\text{NO}$ may outcompete SOD and react with $\cdot\text{O}_2^-$ to generate ONOO \cdot [79] (Figure 1, reaction 3), which is capable of inhibiting enzyme and membrane functions [23]. In particular, ONOO \cdot is responsible for decreased activity of the H_2O_2 -removing enzyme glutathione peroxidase in sepsis [80]. The relevance of this interaction in sepsis is demonstrated by the fact that selenium, which is crucial for glutathione peroxidase metabolism, shows clinical benefits in sepsis treatment [25,80].

The most important effects of ONOO \cdot in sepsis are exerted indirectly on mitochondria. It causes DNA single-strand breakage in the nucleus, which signals poly(ADP-ribose) polymerase (PARP) to be activated. PARP catalyses the cleavage of NAD^+ , which leads to a decrease in the NAD^+ level and to an increase in the NADH/NAD^+ ratio [54,81]. A high NADH/NAD^+ ratio is known to raise $\cdot\text{O}_2^-$ production on the mitochondrial complex I [74]. It appears that the ONOO \cdot -provoked increase in $\cdot\text{O}_2^-$ production on complex I represents the third major event in sepsis following the immune system-generated oxidative burst and iNOS over-expression. *In vitro* studies have shown that PARP activation causes energy depletion and cellular injury, while pharmacological inhibition of PARP improves the response of mitochondria to ONOO \cdot -challenge [81]. In addition, PARP inhibitors have been shown to diminish sepsis-provoked mitochondrial impairment [56]. The importance of the ONOO \cdot /PARP/ NAD^+ /complex I sequence of events in sepsis is illustrated by the fact that PARP-deficient mice with sepsis show improved survival, while the administration of PARP inhibitors is beneficial in a porcine model of sepsis [82,83]. Finally, Khan and co-workers [84] have shown that the application of liposomal NAD^+ on immunostimulated cells (modelling sepsis), leading to a decreased NADH/NAD^+ ratio, prevents diminished consumption of O_2 provoked by ONOO \cdot .

As in the case of $\cdot\text{NO}$ -promoted $\cdot\text{O}_2^-$ production in mitochondria, ONOO \cdot -provoked generation of $\cdot\text{O}_2^-$ results in increased mitochondrial H_2O_2 , which leaks from mitochondria to further promote NF- κB activation, thus closing another redox loop (Figure 2). In line with this, application of the ONOO \cdot -decomposition catalyst FeTPPS has been reported to reduce NF- κB activation in endotoxaemic mice [85].

Mitochondrial dysfunction

Mitochondria are specifically targeted by highly reactive products of ONOO \cdot decomposition - $\cdot\text{OH}$ and $\cdot\text{NO}_2$ [23] (Figure 1, reaction 4). The hydroxyl radical non-selectively

damages membranes, proteins and DNA [24] while $\cdot\text{NO}_2$ binds to proteins leading to increased susceptibility to protease activity [23]. Pertinent to this, research regarding mitochondrial structure after lipopolysaccharide insult revealed oxidative stress and pronounced mitochondrial DNA damage [16,20]. In combination with ETC inhibition, oxidative modifications lead to mitochondrial dysfunction, which is intimately linked to a depleted cellular ATP pool and an energy deficit characteristic of sepsis (Table 3).

Dysfunctional mitochondria show increased susceptibility to mitoptosis. Morphological changes characteristic of mitoptosis have been documented in a feline model of acute endotoxaemia [86], while a decrease in mitochondrial content has been observed in patients with prolonged septic conditions [22]. A lower number of mitochondria may further compromise energy balance. Together, a fall in ATP concentration and oxidative stress exerted on mitochondrial metabolites lead to mitochondrial permeability transition pore opening with eventual organelle swelling and cytochrome c release. Once in the cytoplasm, cytochrome c can bind to apoptotic protease activating factor 1 (APAF-1) and thus activate caspase-9, which in turn activates caspase-3 and -7, culminating in apoptosis in some cell types. Anti-oxidant treatment can prevent mitochondrial swelling and cytochrome c release, pinpointing the role of ROS in mitochondrial dysfunction [87]. Mitochondrial dysfunction represents the fourth and final major redox event in the pathogenesis of sepsis.

Patients typically die of sepsis days to weeks after the onset of the initial infection and in some cases even after the original infection has been eradicated [1,2]. Interestingly 10 to 40% of sepsis patients show negative cultures [1,2]. At the later stages of sepsis the immune system is in a hypoactive mode. Therefore, the plasma oxidative burst can no longer be directly responsible for cell/organ dysfunction [7]. These facts imply that infection could, under specific conditions, activate a self-sustaining and self-promoting redox cycle. Instead of shutting down upon pathogen eradication or in the immune hypoactive stage the 'sepsis redox cycle' continues to drain energy and prevents normal function from being re-established due to several positive feedback loops (Figure 2), even when there is no further inflow of reactive species from plasma. It is important to note that in the 'sepsis redox cycle' H_2O_2 crosses both cellular and mitochondrial membranes. This transport is facilitated by aquaporin-8 [88]. Interestingly, aquaporin-8 is down-regulated in sepsis [89], which may be a mechanism of defence targeted at stopping or slowing down the cycle. The pathophysiological redox profile of one cell can spill over to surrounding tissue, due to the ability of H_2O_2 and $\cdot\text{NO}$ to cross membranes, thus leading to organ failure.

Table 3. The level of ATP in sepsis

Subjects	Results	Reference
Critically ill septic patients and healthy subjects undergoing elective surgery	ATP (nmol/mg dry weight) in skeletal muscle biopsies: Controls (8): 12.5 (9.7 to 13.7) Survivors (12): 15.8 (12.1 to 18.6) Non-survivors (9): 7.6 (6.6 to 10.0) ($P < 0.001$)	[21]
Human umbilical vein endothelial cells incubated with healthy or septic sera	ATP (nmol/L) in the cells exposed to the serum of: Controls: 13.5 ± 1.9 Septic shock patients: 6.7 ± 1.2 ($P < 0.01$)	[76]
Murine model of endotoxaemia (LPS-induced)	ATP ($\mu\text{mol/g}$ wet weight) in liver: Controls: 3.51 ± 0.21 Endotoxaemic: 3.00 ± 0.13 ($P < 0.05$)	[122]

LPS, lipopolysaccharide.

Polymorphisms of redox-relevant genes and sepsis

The question why the activation of the immune system in some patients leads to sepsis and in others does not remain unresolved. On the basis of studies in identical twins genetic factors have been implicated in the determination of susceptibility to death from infections [90]. It seems that specific gene polymorphisms of some redox-active enzymes are more frequent in patients with sepsis and other related conditions in comparison to the general population (Table 4). However, the exact role of gene polymorphisms in sepsis susceptibility and sepsis development, progression and outcome remains to be further elucidated.

Redox therapy in sepsis

Sepsis is treated by controlling the source of infection, administering antimicrobial therapy, assuring hemodynamic support with fluid resuscitation and vasopressor drugs, inducing sedation or analgesia as needed and by other supportive therapies [91]. Despite these interventions, about one-third of all septic patients die before leaving hospital [1,2]. Therefore, the development of additional treatment approaches, such as anti-oxidative therapy, is a research topic of urgent priority.

Although many promising results have been obtained from animal models, believable benefit from anti-oxidant application in sepsis has rarely been translated into success in human clinical trials [92]. This may be explained by several reasons. Animal models of sepsis do not completely recapitulate the human disease or involve identical care delivered to human sepsis patients [93-95]. The onset and progression of sepsis to multi-organ failure occurs over days to weeks whereas in most animal models these processes are much faster, occurring within hours to days. Animal models of sepsis almost exclusively use young animals, which mimic the physiology of a young human. The pitfall here is that epidemiologic studies show that the older human population is more susceptible to sepsis. Moreover, in contrast to animal models, patients with sepsis commonly have co-morbidities [95]. Finally, models and clinical sepsis show very

different cytokine dynamics [93,94]. It should be stressed that therapeutic studies on animals sometimes involve the application of high doses of anti-oxidants exceeding recommended human allowances that clinical researchers are reluctant to use in trials. In addition, animal studies frequently employ early application of therapy (or pre-treatment), which is often not the case with clinical sepsis. Some clinical trials do not take into account all the main principles of anti-oxidative therapy. Halliwell and Gutteridge [23] made a list of questions that generally should be answered preceding the use of an anti-oxidant in humans: (i) Is it non-toxic? (ii) Which biomolecules will the anti-oxidant protect, and are such molecules implicated in the pathogenesis of the disease? (iii) How does the anti-oxidant work - by scavenging reactive species, increasing endogenous defence or by repairing the damage? (iv) Can the anti-oxidant-derived products cause damage? (v) Does the anti-oxidant interfere with redox signalling? Kohen and Nyska [96] sublimed the principles of anti-oxidant therapy as follows: 'Taking into consideration the high reactivity of ROS, their short life span, their continuous production in close proximity to biological targets and their ability to be modified into other more reactive species, one realises that, in order to cope with these deleterious metabolites, the anti-oxidant should be administered to the body continuously, in high concentrations and targeted to the biological site susceptible to oxidative damage.' Pertinent to this and the mechanism proposed here, it is clear that a redox approach in sepsis treatment should be targeted at the specific steps in the pathogenesis. For example, very promising effects of therapies targeting dysfunctional mitochondria have been reviewed recently by Dare and colleagues [16] and Galley [20].

Targeted delivery of anti-oxidants into intracellular space is limited by mechanisms regulating the level of reducing agents in order to maintain normal functioning of signalling pathways that involve reactive species [97]. Therefore, even if long-term supplementation raises blood anti-oxidants, it will have a limited effect on intracellular levels or redox status [97]. Hence, the majority of

Table 4. Polymorphisms of genes involved in redox regulation that are implicated in sepsis and related conditions

Gene	Polymorphism	Effect of polymorphism	Population	Genotypes	Reference
<i>p22-PHOX</i> (NADPH oxidase subunit)	C242T	T allele is associated with higher plasma levels of nitrotyrosine (circulating biomarker of $\cdot\text{O}_2^-$ and $\cdot\text{NO}$ generation)	Patients with acute renal failure (200)	T-allele carrier state was associated with 2.1-fold higher odds for dialysis requirement or hospital death ($P < 0.01$)	[123]
<i>MNSOD</i>	Ala9Val	Valine variant is associated with decreased intra-mitochondrial transport of MnSOD	Healthy subjects (100) and patients with sepsis (40)	A/A: 20% (frequency in control group) and 27.5% (in sepsis) V/V: 36% and 10% A/V: 44% and 62.5%	[124]
<i>eNOS</i>	G894T	T allele is related to low levels of eNOS activity	Patients with <i>Escherichia coli</i> bacteraemia (42)	Carriage of the T allele was associated with low mean arterial pressure and high Sequential Organ Failure Assessment score	[125]
<i>MPO</i> (gene promoter)	G-129A	A allele causes loss of transcriptional factor binding site and lower amount of MPO in cytoplasmic granules	Patients that underwent chemotherapy (136) or autologous stem cell transplantation (113)	Chemotherapy: GG: 11% developed sepsis GA: 17% developed sepsis ASCT: GG: 7% developed sepsis GA: 25% developed sepsis	[126]
<i>NRF2</i> (gene promoter)	C-617A	A allele presents less transcriptional activity than C allele	Humans with major trauma (164)	Carriage of the A allele was related to a greater risk of acute lung injury ($P = 0.021$)	[127]
<i>HMOX1</i> (gene promoter)	(GT) _n	Longer (GT) _n repeats in the HMOX1 promoter are associated with higher plasma HO-1 levels	ICU patients who developed (437) or did not develop (1014) acute respiratory distress syndrome	Longer (GT) _n repeats are associated with reduced acute respiratory distress syndrome risk	[128]
<i>TNF</i>	TNF1 and TNF2 allele	TNF2 is associated with higher levels of inducible and constitutional TNF	Critically ill infected patients (112)	TNF2 frequency: General population: 5% Studied group: 12% ($P < 0.01$) Overall mortality: TNF1/TNF1: 48% TNF1/TNF2: 62% The frequency of septic shock: TNF1/TNF1: 34% TNF1/TNF2: 50% Mortality in septic shock patients: TNF1/TNF1: 62% TNF1/TNF2: 92% ($P < 0.05$)	[129]
<i>HIF-1α</i>	C85T	T variant presents more transcriptional activity	Patients with acute kidney injury (241)	T-allele carriers had significantly higher odds for in-hospital death ($P < 0.001$)	[130]

eNOS, endothelial NOS; HIF, hypoxia inducible factor; HO-1, haem oxygenase (haem degrading enzyme); MnSOD, manganese superoxide dismutase; MPO, myeloperoxidase (HOCl generator).

anti-oxidants perform their actions predominantly in plasma, which may have beneficial effects during the hyper-inflammatory stage of sepsis but are ineffective against the self-sustaining 'sepsis redox cycle' taking place inside the cell. This raises the question of how to overcome refractory mechanisms and help the organism to fight intracellular oxidative stress against its own 'will'. In other words, in conditions related to intracellular oxidative stress, such as sepsis, reducing agents targeted to intracellular space and capable of accumulating inside the cell are needed.

Potential candidates for such an approach are fructose 1,6-(bis)phosphate and ethyl pyruvate, which act as both antioxidants and energy molecules [98,99], and organisms in times of crisis will not reject energy supplies. These compounds have already shown positive effects in sepsis models [100-103], but are yet to be studied in

clinical trials. Molecular hydrogen that reacts directly with ROS should also be capable of overcoming the refractory response. Recently, Xie and co-workers [104] reported that H₂ inhalation significantly improved the survival rate of septic mice in a concentration- and time-dependent manner.

An alternative anti-oxidative approach is to suppress ROS production in mitochondria. It has been shown that maintenance of glucose in blood within the range 4.4 to 6.1 mM, instead of the normally higher range of 10.0 to 11.1 mM, leads to a decrease in mortality by more than 10% [105]. We propose that decreased availability of glucose could exert positive effects in sepsis by preventing the production of $\cdot\text{O}_2^-$ on mitochondrial complex I. A lower level of glucose leads to decreased generation of NADH and hence to a decrease in the NADH/NAD⁺ ratio. Another way to decrease the NADH/NAD⁺ ratio is

to apply liposomal NAD⁺, which has been shown to prevent diminished O₂ consumption in mitochondria under septic conditions [84]. As an alternative to feeding complex I, which represents a point of electron leakage and [•]O₂⁻ production, mitochondrial complex II could be supplemented with succinate in order to bypass [•]O₂⁻ production on complex I and in parallel to maintain normal energy metabolism. Several studies have confirmed positive effects of succinate in sepsis [106-108].

Pertinent to the important role of iNOS and [•]NO in sepsis-related cell/organ dysfunction, therapeutic strategies capable of decreasing the level of [•]NO could be beneficial in sepsis. The supplementation of selective iNOS inhibitors, such as aminoguanide, L-NAME and L-N⁶-(1-imminoethyl)lysine, has had positive effects in different sepsis models [109-113]. It is interesting that a clinical trial of the non-selective NOS inhibitor L-NMMA in patients with septic shock was terminated early because of increased mortality. Such an adverse effect of high-dose L-NMMA may have resulted from the inhibition of endothelial NOS, whose activity is essential for microvascular homeostasis [114]. In addition to iNOS inhibition, [•]NO excess can be tackled using [•]NO scavengers. For example, diethyldithiocarbamate (DDC) demonstrates positive effects in septic shock mice, partially due to the ability of the iron-DDC complex to sequester [•]NO [115].

Conclusions

Organ failure in sepsis is provoked by cellular dysfunction, a process related to pro-oxidative conditions and dysfunctional mitochondria. The oxidative burst performed by the innate immune system is 'transferred' from the plasma into the cell, where it activates the production of ROS and RNS, resulting in non-physiological redox conditions, disturbed redox signalling, oxidative damage of diverse molecules and cellular structures and energy depletion. These events are interconnected in a self-sustaining and self-promoting pathological process named here as the 'sepsis redox cycle', which may result in the total loss of cellular function. Redox therapies targeting specific steps of the 'sepsis redox cycle' have shown some very promising results for sepsis treatment, but require further clinical evaluation. Although it is obvious that sepsis represents a multi-factorial pathophysiology, studies investigating the effects of multi-target therapies are still scarce. We believe that instead of looking for a single 'magic bullet' for sepsis treatment, a multi-component 'magic bullets' therapeutic approach combining different redox agents capable of suppressing specific steps in the pathogenesis of sepsis is clearly the way forward.

Abbreviations

ARE, anti-oxidant response element; COX, cyclooxygenase; ETC, electron transfer chain; H₂O₂, hydrogen peroxide; HOCl, hypochlorous acid;

IL, interleukin; iNOS, inducible nitric oxide synthase; MetHb, methaemoglobin; MnSOD, manganese superoxide dismutase; NF, nuclear factor; [•]NO, nitric oxide; [•]NO₂, nitrogen dioxide radical; NOS, nitric oxide synthase; Nrf2, nuclear factor E2-related factor 2; [•]O₂⁻, superoxide; [•]OH, hydroxyl radical; ONOO⁻, peroxynitrite; [•]OOH, hydroperoxyl radical; PARP, poly (ADP-ribose) polymerase; RBC, red blood cell; RNS, reactive nitrogen species; [•]ROO, peroxy radical; ROS, reactive oxygen species; SOD, superoxide dismutase; TNF, tumour necrosis factor.

Competing interests

The authors declare that they have no competing interests.

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