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# Σήψη-Πρωτόκολα Αντιμετώπισης

- ΗΠΑ: 2% ασθενών σε νοσοκομεία=σοβαρή σήψη
- Από τους προηγούμενους οι μισοί σε ΜΕΘ
- 10% όλων των εισαγωγών στις ΜΕΘ
- ΗΠΑ: 750000 ανά έτος
- Παγκοσμία: 19.000.000 και ο αριθμός μεγαλώνει

R. P. Dellinger Mitchell M. Levy Andrew Rhodes Djillali Annane Herwig Gerlach Steven M. Onal Surviving Sepsis Campaign: International Guidelines for Management of Severe Sepsis and Septic Shock, 2012

Sepsis is defined as the presence (probable or documented) of infection together with systemic manifestations of infection. Severe sepsis is defined as sepsis plus sepsis-induced organ dysfunction or tissue hypoperfusion (Tables 1, 2) [6]. Infection, documented or suspected, and some of the following: General variables Fever (>38.3 °C) Hypothermia (core temperature <36 °C) Heart rate  $>90 \text{ min}^{-1}$  or more than two SD above the normal value for age Tachypnea Altered mental status Significant edema or positive fluid balance (> 20 mL/kg over 24 h) Hyperglycemia (plasma glucose >140 mg/dL or 7.7 mmol/L) in the absence of diabetes Inflammatory variables Leukocytosis (WBC count >12,000  $\mu$ L<sup>-1</sup>) Leukopenia (WBC count <4,000  $\mu$ L<sup>-1</sup>) Normal WBC count with greater than 10 % immature forms Plasma C-reactive protein more than two SD above the normal value Plasma procalcitonin more than two SD above the normal value Hemodynamic variables Arterial hypotension (SBP <90 mmHg, MAP <70 mmHg, or an SBP decrease >40 mmHg in adults or less than two SD below normal for age) Organ dysfunction variables Arterial hypoxemia ( $PaO_2/FiO_2 < 300$ ) Acute oliguria (urine output  $< 0.5 \text{ mL} \text{ kg}^{-1} \text{ h}^{-1}$  for at least 2 h despite adequate fluid resuscitation) Creatinine increase >0.5 mg/dL or 44.2 µmol/L Coagulation abnormalities (INR >1.5 or aPTT >60 s) Ileus (absent bowel sounds) Thrombocytopenia (platelet count <100,000  $\mu$ L<sup>-1</sup>) Hyperbilirubinemia (plasma total bilirubin >4 mg/dL or 70  $\mu$ mol/L) Tissue perfusion variables Hyperlactatemia (>1 mmol/L) Decreased capillary refill or mottling

SD standard deviation, WBC white blood cell, SBP systolic blood pressure, MAP mean arterial pressure, INR international normalized ratio, aPTT activated partial thromboplastin time Diagnostic criteria for sepsis in the pediatric population are signs and symptoms of inflammation plus infection with hyper- or

hypothermia (rectal temperature >38.5 or <35 °C), tachycardia (may be absent in hypothermic patients), and at least one of the following indications of altered organ function: altered mental status, hypoxemia, increased serum lactate level, or bounding pulses Adapted from [6]

### Table 2Severe sepsis

Severe sepsis definition = sepsis-induced tissue hypoperfusion or organ dysfunction (any of the following thought to be due to the infection) Sepsis-induced hypotension Lactate above upper limits laboratory normal Urine output <0.5 mL kg<sup>-1</sup> h<sup>-1</sup> for more than 2 h despite adequate fluid resuscitation Acute lung injury with Pao<sub>2</sub>/Fio<sub>2</sub> <250 in the absence of pneumonia as infection source Acute lung injury with  $Pao_2/Fio_2 < 200$  in the presence of pneumonia as infection source Creatinine >2.0 mg/dL (176.8  $\mu$ mol/L) Bilirubin >2 mg/dL (34.2  $\mu$ mol/L) Platelet count  $<100,000 \ \mu L$ Coagulopathy (international normalized ratio >1.5)



# Host – pathogen mismatch

The immune profile of this host-pathogen mismatch can be predominately proinflammatory (systemic inflammatory response syndrome, SIRS), mixed (mixed antagonistic response syndrome, MARS), or anti-inflammatory (compensatory anti-inflammatory response syndrome, CARS). The final result is various degrees of hyperinflammation, immunosuppression, abnormal coagulation, and microcirculatory dysfunction, all which may contribute to organ injury and cell death.<sup>2,6</sup>

Clinical diagnosis of *severe sepsis* or *septic shock* although valuable and of significant importance for the management of septic patients may lead to extremely heterogeneous cohorts in terms of patients' immunological status. This heterogeneity offers one explanation for the failure of prior trials of biologic therapies for sepsis, since treatments that focused on attenuating the initial inflammatory response of sepsis in a sense ignored and in fact might have exacerbated the progressive development of immunosuppression in some patients.<sup>8-11</sup>

Severe sepsis occurs as a result of both community-acquired and health care-associated infections. Pneumonia is the most common cause accounting for about half of all cases, followed by intraabdominal and urinary tract infections.7,8,11,12 Blood cultures are typically positive in only one third of cases, and in up to a third of cases, cultures from all sites are negative.<sup>7,11,13,14</sup> Staphylococcus aureus and Streptococcus pneumoniae are the most common gram-positive isolates, whereas Escherichia coli, klebsiella species, and Pseudomonas aeruginosa predominate among gram-negative isolates.<sup>11,14</sup> An epidemiologic study of sepsis showed that during the period from 1979 to 2000, gram-positive infections overtook gramnegative infections.<sup>15</sup> However, in a more recent study involving 14,000 ICU patients in 75 countries, gram-negative bacteria were isolated in 62% of patients with severe sepsis who had positive cultures, gram-positive bacteria in 47%, and fungi in 19%.<sup>12</sup>

#### CRITICAL CARE MEDICINE

Simon R. Finfer, M.D., and Jean-Louis Vincent, M.D., Ph.D., Editors

Severe Sepsis and Septic Shock Derek C. Angus, M.D., M.P.H., and Tom van der Poll, M.D., Ph.D.

other.<sup>30</sup> T<u>he specific response in any patient depends on the causative pathogen (load and virulence) and the host (genetic characteristics and coexisting illnesses), with differential responses at local, regional, and systemic levels (Fig. 1). The composition and direction of the host response probably change over time in parallel with the clinical course. In general, proinflammatory reactions (directed at eliminating invading pathogens) are thought to be responsible for collateral tissue damage in severe sepsis, whereas antiinflammatory responses (important for limiting local and systemic tissue injury) are implicated in the enhanced susceptibility to secondary infections.</u>

### **EVOLUTION TO IMMUNOPARALYSIS (FIG. 62-1)**<sup>31</sup>



Excessive inflammation causing collateral damage (tissue injury)



### Figure 1. The Host Response in Severe Sepsis.

Proinflammatory response

The host response to sepsis is characterized by both proinflammatory responses (top of panel, in red) and antiinflammatory immunosuppressive responses (bottom of panel, in blue). The direction, extent, and duration of these reactions are determined by both host factors (e.g., genetic characteristics, age, coexisting illnesses, and medications) and pathogen factors (e.g., microbial load and virulence). Inflammatory responses are initiated by interaction between pathogen-associated molecular patterns expressed by pathogens and patternrecognition receptors expressed by host cells at the cell surface (toll-like receptors [TLRs] and C-type lectin receptors [CLRs]), in the endosome (TLRs), or in the cytoplasm (retinoic acid inducible gene 1–like receptors [RLRs] and nucleotide-binding oligomerization domain–like receptors [NLRs]). The consequence of exaggerated inflammation is collateral tissue damage and necrotic cell death, which results in the release of damage-associated molecular patterns, so-called danger molecules that perpetuate inflammation at least in part by acting on the same pattern-recognition receptors that are triggered by pathogens.



#### Figure 2. Organ Failure in Severe Sepsis and Dysfunction of the Vascular Endothelium and Mitochondria.

Sepsis is associated with microvascular thrombosis caused by concurrent activation of coagulation (mediated by tissue factor) and impairment of anticoagulant mechanisms as a consequence of reduced activity of endogenous anticoagulant pathways (mediated by activated protein C, antithrombin, and tissue factor pathway inhibitor), plus impaired fibrinolysis owing to enhanced release of plasminogen activator inhibitor type 1 (PAI-1). The capacity to generate activated protein C is impaired at least in part by reduced expression of two endothelial receptors: thrombomodulin (TM) and the endothelial protein C receptor. Thrombus formation is further facilitated by neutrophil extracellular traps (NETs) released from dying neutrophils. Thrombus formation results in tissue hypoperfusion, which is aggravated by vasodilatation, hypotension, and reduced red-cell deformability. Tissue oxygenation is further impaired by the loss of barrier function of the endothelium owing to a loss of function of vascular endothelial (VE) cadherin, alterations in endothelial cell-to-cell tight junctions, high levels of angiopoietin 2, and a disturbed balance between sphingosine-1 phosphate receptor 1 (S1P1) and S1P3 within the vascular wall, which is at least in part due to preferential induction of S1P3 through protease activated receptor 1 (PAR1) as a result of a reduced ratio of activated protein C to thrombin. Oxygen use is impaired at the subcellular level because of damage to mitochondria from oxidative stress.



#### Figure 2. Procoagulant Response in Sepsis.

Sepsis initiates coagulation by activating endothelium to increase the expression of tissue factor. Activation of the coagulation cascade, and especially factors Va and VIIIa, leads to the formation of thrombin-α, which converts fibrinogen to fibrin. Fibrin binds to platelets, which in turn adhere to endothelial cells, forming microvascular thrombi. Microvascular thrombi amplify injury through the release of mediators and by microvascular obstruction, which causes distal ischemia and tissue hypoxia. Normally, natural anticoagulants (protein C and protein S), antithrombin III, and tissue factor-pathway inhibitor (TFPI) dampen coagulation, enhance fibrinolysis, and remove microthrombi. Thrombin- $\alpha$  binds to thrombomodulin on endothelial cells, which dramatically increases activation of protein C to activated protein C. Protein C forms a complex with its cofactor protein S. Activated protein C proteolytically inactivates factors Va and VIIIa and decreases the synthesis of plasminogen-activator inhibitor 1 (PAI-1). In contrast, sepsis increases the synthesis of PAI-1. Sepsis also decreases the levels of protein C, protein S, antithrombin III, and TFPI. Lipopolysaccharide and turnor necrosis factor  $\alpha$  (TNF- $\alpha$ ) decrease the synthesis of thrombomodulin and endothelial protein C receptor (EPCR), thus decreasing the activation of protein C. Sepsis further disrupts the protein C pathway because sepsis also decreases the expression of EPCR, which amplifies the deleterious effects of the sepsis-induced decrease in levels of protein C. Lipopolysaccharide and TNF- $\alpha$  also increase PAI-1 levels so that fibrinolysis is inhibited. The clinical consequences of the changes in coagulation caused by sepsis are increased levels of markers of disseminated intravascular coagulation and widespread organ dysfunction. t-PA denotes tissue plasminogen activator.



Figure 1. Proposed Actions of Activated Protein C in Modulating the Systemic Inflammatory, Procoagulant, and Fibrinolytic Host Responses to Infection.

The inflammatory and procoagulant host responses to infection are intricately linked. Infectious agents and inflammatory cytokines such as tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) and interleukin-1 activate coagulation by stimulating the release of tissue factor from monocytes and the endothelium. The presentation of tissue factor leads to the formation of thrombin and a fibrin clot. Inflammatory cytokines and thrombin can both impair the endogenous fibrinolytic potential by stimulating the release of plasminogen-activator inhibitor 1 (PAI-1) from platelets and the endothelium. PAI-1 is a potent inhibitor of tissue plasminogen activator, the endogenous pathway for lysing a fibrin clot. In addition, the procoagulant thrombin is capable of stimulating multiple inflammatory pathways and further suppressing the endogenous fibrinolytic system by activating thrombin-activatable fibrinolysis inhibitor (TAFI). The conversion of protein C, by thrombin bound to thrombomodulin, to the serine protease activated protein C is impaired by the inflammatory response. Endothelial injury results in decreased thrombomodulin levels. The end result of the host response to infection may be the development of diffuse endovascular injury, microvascular thrombosis, organ ischemia, multiorgan dysfunction, and death. Activated protein C can intervene at multiple points during the systemic response to infection. It exerts an antithrombotic effect by inactivating factors Va and VIIIa, limiting the generation of thrombin. As a result of decreased thrombin levels, the inflammatory, procoagulant, and antifibrinolytic response induced by thrombin is reduced. In vitro data indicate that activated protein C exerts an antiinflammatory effect by inhibiting the production of inflammatory cytokines (TNF- $\alpha$ , interleukin-1, and interleukin-6) by monocytes and limiting the rolling of monocytes and neutrophils on injured endothelium by binding selectins. Activated protein C indirectly increases the fibrinolytic response by inhibiting PAI-1,12-17





### Intravital sidestream dark field images of the sublingual microcirculation:

capillaries with red cells flowing (normal, upper image) and obstructed with minimal flow (sepsis, lower image).

Images courtesy of C Ince, Amsterdam, the Netherlands



**Clinical Review & Education** 

### Special Communication | CARING FOR THE CRITICALLY ILL PATIENT

# The Third International Consensus Definitions for Sepsis and Septic Shock (Sepsis-3)

Mervyn Singer, MD, FRCP; Clifford S. Deutschman, MD, MS; Christopher Warren Seymour, MD, MSc; Manu Shankar-Hari, MSc, MD, FFICM; Djillali Annane, MD, PhD; Michael Bauer, MD; Rinaldo Bellomo, MD; Gordon R. Bernard, MD; Jean-Daniel Chiche, MD, PhD; Craig M. Coopersmith, MD; Richard S. Hotchkiss, MD; Mitchell M. Levy, MD; John C. Marshall, MD; Greg S. Martin, MD, MSc; Steven M. Opal, MD; Gordon D. Rubenfeld, MD, MS; Tom van der Poll, MD, PhD; Jean-Louis Vincent, MD, PhD; Derek C. Angus, MD, MPH

### Box 1. SIRS (Systemic Inflammatory Response Syndrome)

Two or more of:

Temperature >38°C or <36°C

Heart rate >90/min

Respiratory rate >20/min or Paco<sub>2</sub> <32 mm Hg (4.3 kPa)

White blood cell count >12 000/mm<sup>3</sup> or <4000/mm<sup>3</sup> or >10% immature bands

From Bone et al.<sup>9</sup>

### Improved Understanding of Sepsis Pathobiology

Sepsis is a multifaceted host response to an infecting pathogen that may be significantly amplified by endogenous factors.<sup>14,15</sup> The original conceptualization of sepsis as infection with at least 2 of the 4 SIRS criteria focused solely on inflammatory excess. However. the validity of SIRS as a descriptor of sepsis pathobiology has been challenged. Sepsis is now recognized to involve early activation of both pro- and anti-inflammatory responses,<sup>16</sup> along with major modifications in nonimmunologic pathways such as cardiovascular, neuronal, autonomic, hormonal, bioenergetic, metabolic, and coagulation,<sup>14,17,18</sup> all of which have prognostic significance. Organ dysfunction, even when severe, is not associated with substantial cell death.<sup>19</sup>



### Sepsis

The current use of 2 or more SIRS criteria (Box 1) to identify sepsis was unanimously considered by the task force to be unhelpful. Changes in white blood cell count. temperature. and heart rate reflect inflammation. the host response to "danger" in the form of infection or other insults. The SIRS criteria do not necessarily indicate a dysregulated, life-threatening response. SIRS criteria are present in many hospitalized patients, including those who never develop infection and never incur adverse outcomes (poor discriminant validity).<sup>25</sup> In addition, 1 in 8 patients admitted to criti-

### Box 2. Key Concepts of Sepsis

• Sepsis is the primary cause of death from infection, especially if <u>not recognized and treated promptly</u>. Its recognition mandates urgent attention.

• Sepsis is a syndrome shaped by pathogen factors and host factors (eg, sex, race and other genetic determinants, age, comorbidities, environment) with characteristics that evolve over time. <u>What</u> <u>differentiates sepsis from infection is an aberrant or dysregulated</u> <u>host response and the presence of organ dysfunction.</u>

• Sepsis-induced organ dysfunction may be occult; therefore, its presence should be considered in any patient presenting with infection. Conversely, unrecognized infection may be the cause of new-onset organ dysfunction. Any unexplained organ dysfunction should thus raise the possibility of underlying infection.

• The clinical and biological phenotype of sepsis can be modified by preexisting acute illness, long-standing comorbidities, medication, and interventions.

• Specific infections may result in local organ dysfunction without generating a dysregulated systemic host response.

## SOFA Score

### Table 1. Sequential [Sepsis-Related] Organ Failure Assessment Score<sup>a</sup>

	Score				
System	0	1	2	3	4
Respiration					
Pao <sub>2</sub> /Fio <sub>2</sub> , mm Hg (kPa)	≥400 (53.3)	<400 (53.3)	<300 (40)	<200 (26.7) with respiratory support	<100 (13.3) with respiratory support
Coagulation					
Platelets, ×10 <sup>3</sup> /µL	≥150	<150	<100	<50	<20
Liver					
Bilirubin, mg/dL (µmol/L)	<1.2 (20)	1.2-1.9 (20-32)	2.0-5.9 (33-101)	6.0-11.9 (102-204)	>12.0 (204)
Cardiovascular	MAP ≥70 mm Hg	MAP <70 mm Hg	Dopamine <5 or dobutamine (any dose) <sup>b</sup>	Dopamine 5.1-15 or epinephrine $\leq 0.1$ or norepinephrine $\leq 0.1^{b}$	Dopamine >15 or epinephrine >0.1 or norepinephrine >0.1 <sup>b</sup>
Central nervous system					
Glasgow Coma Scale score <sup>c</sup>	15	13-14	10-12	6-9	<6
Renal					
Creatinine, mg/dL (µmol/L)	<1.2 (110)	1.2-1.9 (110-170)	2.0-3.4 (171-299)	3.5-4.9 (300-440)	>5.0 (440)
Urine output, mL/d				<500	<200
Abbreviations: FIO <sub>2</sub> , fraction of inspired oxygen; MAP, mean arterial pressure;		<sup>b</sup> Catecholamine doses are given as µg/kg/min for at least 1 hour.			
Pao <sub>2</sub> , partial pressure of oxygen. <sup>a</sup> Adapted from Vincent et al. <sup>27</sup>		<sup>c</sup> Glasgow Coma Scale scores range from 3-15; higher score indicates better neurological function.			

### Box 3. New Terms and Definitions

- Sepsis is defined as life-threatening organ dysfunction caused by a dysregulated host response to infection.
- Organ dysfunction can be identified as an acute change in total SOFA score  $\geq 2$  points consequent to the infection.
  - The baseline SOFA score can be assumed to be zero in patients not known to have preexisting organ dysfunction.
  - A SOFA score ≥2 reflects an overall mortality risk of approximately 10% in a general hospital population with suspected infection. Even patients presenting with modest dysfunction can deteriorate further, emphasizing the seriousness of this condition and the need for prompt and appropriate intervention, if not already being instituted.
- In lay terms, sepsis is a life-threatening condition that arises when the body's response to an infection injures its own tissues and organs.
- Patients with suspected infection who are likely to have a prolonged ICU stay or to die in the hospital can be promptly identified at the bedside with qSOFA, ie, alteration in mental status, systolic blood pressure  $\leq 100 \text{ mm Hg}$ , or respiratory rate  $\geq 22/\text{min.}$
- Septic shock is a subset of sepsis in which underlying circulatory and cellular/metabolic abnormalities are profound enough to substantially increase mortality.
- Patients with septic shock can be identified with a clinical construct of sepsis with persisting hypotension requiring vasopressors to maintain MAP ≥ 65 mm Hg and having a serum lactate level
   2 mmol/L (18 mg/dL) despite adequate volume resuscitation.
   With these criteria, hospital mortality is in excess of 40%.

Abbreviations: MAP, mean arterial pressure; qSOFA, quick SOFA; SOFA: Sequential [Sepsis-related] Organ Failure Assessment.



### Box 4. qSOFA (Quick SOFA) Criteria

Respiratory rate  $\geq$  22/min

Altered mentation

Systolic blood pressure  $\leq$ 100 mm Hg

### **Screening for Patients Likely to Have Sepsis**

A parsimonious clinical model developed with multivariable logistic regression identified that any 2 of 3 clinical variables— Glasgow Coma Scale score of 13 or less, systolic blood pressure of 100 mm Hg or less, and respiratory rate 22/min or greater—offered predictive validity (AUROC = 0.81; 95% CI, 0.80-0.82) similar to that of the full SOFA score outside the ICU.<sup>12</sup> This model was robust



The Delphi process assessed agreements on descriptions of terms such as "hypotension," "need for vasopressor therapy," "raised lactate," and "adequate fluid resuscitation" for inclusion within the new clinical criteria. The majority (n = 14/17; 82.4%) of task force members voting on this agreed that hypotension should be denoted as a mean arterial pressure less than 65 mm Hg according to the pragmatic decision that this was most often recorded in data sets derived from patients with sepsis. Systolic blood pressure was used as a qSOFA criterion because it was most widely recorded in the electronic health record data sets.

A majority (11/17; 64.7%) of the task force agreed, whereas 2 (11.8%) disagreed, that an elevated lactate level is reflective of cellular dysfunction in sepsis, albeit recognizing that multiple factors, such as insufficient tissue oxygen delivery, impaired aerobic respiration, accelerated aerobic glycolysis, and reduced hepatic clearance, also contribute.<sup>32</sup> Hyperlactatemia is, however, a reasonable marker of illness severity, with higher levels predictive of higher mortality.<sup>33</sup> Criteria for "adequate fluid resuscitation" or "need for vasopressor therapy" could not be explicitly specified because these are highly user dependent, relying on variable monitoring modalities and hemodynamic targets for treatment.<sup>34</sup> Other aspects of management, such as sedation and volume status assessment, are also potential confounders in the hypotensionvasopressor relationship.

By Delphi consensus process, 3 variables were identified (hypotension, elevated lactate level, and a sustained need for vasopressor therapy) to test in cohort studies, exploring alternative

### **Definition of Septic Shock**

Septic shock is defined as a subset of sepsis in which underlying circulatory and cellular metabolism abnormalities are profound enough to substantially increase mortality (Box 3). The 2001 task force definitions described septic shock as "a state of acute circulatory failure."<sup>10</sup> The task force favored a broader view to differentiate septic shock from cardiovascular dysfunction alone and to recognize the importance of cellular abnormalities (Box 3). There was unanimous agreement that septic shock should reflect a more severe illness with a much higher likelihood of death than sepsis alone.

#### **Clinical Review & Education**

Special Communication | CARING FOR THE CRITICALLY ILL PATIENT The Third International Consensus Definitions for Sepsis and Septic Shock (Sepsis-3)

Mervyn Singer, MD, FRCP; Clifford S. Deutschman, MD, MS; Christopher Warren Seymour, MD, MSc; Manu Shankar-Hari, MSc, MD, FFICM; Djillali Annane, MD, PhD; Michael Bauer, MD; Rinaldo Bellomo, MD; Gordon R. Bernard, MD; Jean-Daniel Chiche, MD, PhD; Craig M. Coopersmith, MD; Richard S. Hotchkiss, MD; Mitchell M. Levy, MD; John C. Marshall, MD; Greg S. Martin, MD, MSc; Steven M. Opal, MD; Gordon D. Rubenfeld, MD, MS; Tom van der Poll, MD, PhD; Jean-Louis Vincent, MD, PhD; Derek C. Angus, MD, MPH

Table 2. Guidelines for the Treatment of Severe Sepsis and Septic Shock from the Surviving Sepsis Campai	gn.*			
Element of Care				
Resuscitation	The NEW ENGLAND JOURNAL of MEDICINE			
Begin goal-directed resuscitation during first 6 hr after recognition				
Begin initial fluid resuscitation with crystalloid and consider the addition of albumin	REVIEW ARTICLE			
Consider the addition of albumin when substantial amounts of crystalloid are required to maintain a				
Avoid hetastarch formulations	CRITICAL CARE MEDICINE			
Begin initial fluid challenge in patients with tissue hypoperfusion and suspected hypovolemia, to acl	Simon R. Finfer, M.D., and Jean-Louis Vincent, M.D., Ph.D., <i>Editors</i>			
Continue fluid-challenge technique as long as there is hemodynamic improvement	Severe Sepsis and Septic Shock			
Use norepinephrine as the first-choice vasopressor to maintain a mean arterial pressure of $\geq$ 65 mm	Derek C. Angus, M.D., M.P.H., and Tom van der Poll, M.D., Ph	1.D.		
Use epinephrine when an additional agent is needed to maintain adequate blood pressure		2B		
Add vasopressin (at a dose of 0.03 units/min) with weaning of norepinephrine, if tolerated		UG		
Avoid the use of dopamine except in carefully selected patients (e.g., patients with a low risk of arrhythe function or low heart rate)	mias and either known marked left ventricular systolic dys-	2C		
Infuse dobutamine or add it to vasopressor therapy in the presence of myocardial dysfunction (e.g., ele going hypoperfusion despite adequate intravascular volume and mean arterial pressure	vated cardiac filling pressures or low cardiac output) or on-	1 <b>C</b>		
Avoid the use of intravenous hydrocortisone if adequate fluid resuscitation and vasopressor therapy restore hemodynamic stability; if hydrocortisone is used, ad- minister at a dose of 200 mg/day				
Target a hemoglobin level of 7 to 9 g/dl in patients without hypoperfusion, critical coronary artery disea	ase or myocardial ischemia, or acute hemorrhage	1B		
Infection control				
Obtain blood cultures before antibiotic therapy is administered		1 <b>C</b>		
Perform imaging studies promptly to confirm source of infection		UG		
Administer broad-spectrum antibiotic therapy within 1 hr after diagnosis of either severe sepsis or sept	ic shock	1B/1C		
Reassess antibiotic therapy daily for de-escalation when appropriate		1B		
Perform source control with attention to risks and benefits of the chosen method within 12 hr after dia	gnosis	1C		
Respiratory support				
Use a low tidal volume and limitation of inspiratory-plateau-pressure strategy for ARDS		1 <b>A/1B</b>		
Apply a minimal amount of positive end-expiratory pressure in ARDS		1B		
Administer higher rather than lower positive end-expiratory pressure for patients with sepsis-induced A	RDS	2C		
Use recruitment maneuvers in patients with severe refractory hypoxemia due to ARDS		2C		
Use prone positioning in patients with sepsis-induced ARDS and a ratio of the partial pressure of arteri <100, in facilities that have experience with such practice	al oxygen (mm Hg) to the fraction of inspired oxygen of	2C		
Elevate the head of the bed in patients undergoing mechanical ventilation, unless contraindicated		1B		
Use a conservative fluid strategy for established acute lung injury or ARDS with no evidence of tissue h	ypoperfusion	1C		
Use weaning protocols		1A		

### **REVIEW ARTICLE**

### CRITICAL CARE MEDICINE

Simon R. Finfer, M.D., and Jean-Louis Vincent, M.D., Ph.D., Editors

### Severe Sepsis and Septic Shock

Derek C. Angus, M.D., M.P.H., and Tom van der Poll, M.D., Ph.D.

Central nervous system support	
Use sedation protocols, targeting specific dose-escalation end points	1B
Avoid neuromuscular blockers if possible in patients without ARDS	1C
Administer a short course of a neuromuscular blocker (<48 hr) for patients with early, severe ARDS	2C
General supportive care	
Use a protocol-specified approach to blood glucose management, with the initiation of insulin after two consecutive blood glucose levels of >180 mg/dl (10 mmo liter), targeting a blood glucose level of <180 mg/dl	l/ 1A
Use the equivalent of continuous venovenous hemofiltration or intermittent hemodialysis as needed for renal failure or fluid overload	2B
Administer prophylaxis for deep-vein thrombosis	1B
Administer stress-ulcer prophylaxis to prevent upper gastrointestinal bleeding	1 <b>B</b>
Administer oral or enteral feedings, as tolerated, rather than either complete fasting or provision of only intravenous glucose within the first 48 hr after a diagnosi of severe sepsis or septic shock	s 2C
Address goals of care, including treatment plans and end-of-life planning as appropriate	18

\* Data are adapted from Dellinger et al.<sup>23</sup> ARDS denotes acute respiratory distress syndrome, and ICU intensive care unit.

† For all grades, the number indicates the strength of the recommendation (1, recommended; 2, suggested), and the letter indicates the level of evidence, from high (A) to low (D), with UG indicating ungraded. Recommendations that are specific to pediatric severe sepsis include therapy with face-mask oxygen, high-flow nasal cannula oxygen, or nasopharyngeal continuous positive end-expiratory pressure in the presence of respiratory distress and hypoxemia (2C); use of physical examination therapeutic end points, such as capillary refill (2C); administration of a bolus of 20 ml of crystalloids (or albumin equivalent) per kilogram of body weight during a period of 5 to 10 minutes for hypovolemia (2C); increased use of inotropes and vasodilators in septic shock with low cardiac output associated with elevated systemic vascular resistance (2C); and use of hydrocortisone only in children with suspected or proven absolute adrenal insufficiency (2C).

the guidelines recommend completing the initial fluid resuscitation within 3 hours (UG).

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Original Investigation | CARING FOR THE CRITICALLY ILL PATIENT

## Developing a New Definition and Assessing New Clinical Criteria for Septic Shock For the Third International Consensus Definitions for Sepsis and Septic Shock (Sepsis-3)

Manu Shankar-Hari, MD, MSc; Gary S. Phillips, MAS; Mitchell L. Levy, MD; Christopher W. Seymour, MD, MSc; Vincent X. Liu, MD, MSc; Clifford S. Deutschman, MD; Derek C. Angus, MD, MPh; Gordon D. Rubenfeld, MD, MSc; Mervyn Singer, MD, FRCP; for the Sepsis Definitions Task Force

### Systematic Review and Meta-analysis

The systematic review identified 44 studies (166 479 patients) reporting septic shock mortality<sup>5-7,19-59</sup> from a total of 92 studies reporting sepsis cohorts between 1987 and 2015<sup>5-7,19-107</sup> (**Figure 1**; eTable 2 in the Supplement). Different shock criteria were used for systolic blood pressure (<90 mm Hg; <100 mm Hg; decrease >40 mm Hg; or decrease >50% of baseline value if hypertensive), mean arterial pressure (<70; <65; <60 mm Hg), serum lactate level (>4, >2.5, >2, >1 mmol/L) and base deficit (-5 mmol/L) (**Table 1**;

Table 1. Summary of Septic Shock Definitions and Criteria Reported in the Studies Identified by the Systematic Review<sup>a</sup>

Septic Shock Case Definitions and Corresponding Variables Reported in Literature

### ♥▼ diamantoaretha

### πόσα διαφορετικά κριτήρια σε διαφορες παραμέτρους

	Consensus Definitions		Other Definitions		οιαφορες παραμετρους	
Criteria	Bone et al <sup>1</sup>	Levy et al <sup>2</sup>	SSC <sup>111</sup>	Trial-based <sup>112</sup>		
Infection	Suspected or proven	Suspected or proven	Suspected or proven	Suspected or proven	16/3/16 10:12 µµ	
SIRS criteria, No.	2	One or more of 24 variables <sup>b</sup>	2	3	Reply	
Septic shock description	Sepsis-induced hypotension despite adequate resuscitation OR receiving vasopressors/Inotropes plus presence of perfusion abnormalities	State of acute circulatory failure characterized by persistent arterial hypotension after adequate resuscitation unexplained by other causes	Sepsis-induced hypotension persisting despite adequate fluid resuscitation	Cardiovascular dysfunction defined as hypotension despite adequate resuscitation or need for vasopressors	Precoded data using ICD-9 and ICD-10 codes <sup>c</sup>	
Hypotension, mm Hg						
Systolic BP	<90	<90	<90	<90	<100	
Decrease in systolic BP	Decrease >40	Decrease >40	Decrease >40	NA<70	>50% decrease in hypertension	
MAP	No	<60	<70	Hypotension lasting >1 h after resuscitation	<65	
Adequate resuscitation definition	Not defined	Not defined	Goals set as CVP 8-12 mm Hg; urine output ≥0.5 mL/kg/h; ScvO <sub>2</sub> >70%	Not defined	After resuscitation fluids (0.5 L; 1 L; 1.5 L; 20 mL/kg ideal body weight	
Vasopressor use	Yes (not absolute requirement)	Yes (CVS SOFA score)	Yes (not absolute requirement)	Yes (not absolute requirement)	Vasoactive drugs required for >30 min	
Hypoperfusion abnormalities	Hypoperfusion abnormality defined as lactic acidosis; oliguria; low Glasgow Coma Score	Tissue hypoperfusion defined as serum lactate >1 mmol/L or delayed capillary refill	Tissue hypoperfusion defined as infection-induced hypotension, elevated serum lactate (>4 mmol/L), or oliguria	No description	Serum lactate >2.5 mmol/L; base deficit >5 mEq/L, alkaline reserve <18 mEq/L; CVP <8; PCWP <12	
Data points from included studies, No. (%) <sup>d</sup>	39 (75)		13 (25)			
Sample size, No.	158 354		8125			
Mortality by septic shock definition using random-effects meta analysis, % (95% CI)	47.2 (42.	7-51.7)	44.2 (3	8.5-49.9)		
l <sup>2</sup> , % <sup>e</sup>	99.6		95.9			
$\tau^{2f}$	191.21		94.9			
P value heterogeneity	<.001		<.001			

#### Figure 2. Random-Effects Meta-analysis of Studies Identified in the Systematic Review, ig Septic Shock Mortality

**Patients With** 

<sup>(°)</sup>.8 (60.7-67.0)

.6 (39.8-51.4)

.2 (45.3-59.1)

.1 (36.5-61.8)

.1 (49.6-58.7)

.8 (20.4-29.2)

.1 (49.6-58.7)

.5 (67.1-78.0)

.7 (34.8-38.6)

.0 (44.3-49.7)

.5 (35.7-47.2)

.6 (60.0-61.2)

.3 (41.2-43.3)

.3 (31.1-61.6)

0



	Septic Shock Deaths, No.	Septic Shock, No.	Mortality, % (95% CI)
Consensus Definition			
Degoricija et al, <sup>46</sup> 2006	90	125	72.0 (64.1-79.9)
Angkasekwinai et al, <sup>38</sup> 2007	41	78	52.6 (41.5-63.6)
Nesseler et al, <sup>27</sup> 2013	30	93	32.3 (22.8-41.8)
Sakr et al, <sup>25</sup> 2013	85	145	58.6 (50.6-66.6)
Goncalves-Pereira et al, <sup>23</sup> 2014	418	856	48.8 (45.5-52.2)
Leligdowicz et al, <sup>5</sup> 2014	4146	7974	52.0 (50.9 -53.1)
Ortiz et al, <sup>19</sup> 2014	144	319	45.1 (39.7-50.6)
Hypotension			
Laupland et al, <sup>47</sup> 2004	81	159	50.9 (43.2-58.7)
Gaspraovic et al, <sup>45</sup> 2006	44	129	34.1 (25.9-42.3)
Shapiro et al, <sup>44</sup> 2006	15	53	28.3 (16.2-40.4)
Povoa et al, <sup>35</sup> 2009	202	458	44.1 (39.6-48.7)
Klein Klowenberg et al, <sup>7</sup> 2012	52	98	53.1 (43.2-62.9)
Kaukonen et al, <sup>22</sup> 2014	14609	51079	28.6 (28.2-29.0)
Hypotension + Perfusion Abnormalities a	nd/or Vasopressor The	erapy	
Rangel-Frausto et al, <sup>56</sup> 1995	51	110	46.4 (37.0-55.7)
Salvo et al, <sup>55</sup> 1995	27	33	81.8 (68.7-95.0)

The random-effects meta-analysis showed significant heterogeneity in septic shock mortality (mean mortality, 46.5% [95% CI, 42.7%-50.3%], with a near 4-fold variation from 23.0% to 81.8%;  $I^2$  = 99.5%;  $\tau^2$  = 182.5; and P < .001) (Figure 2). Statistically significant heterogeneity was also observed in random-effects meta-analysis by clinical criteria reported for septic shock case definition in studies (Table 2).

			.1 (45.0-47.2)
Quenot et al, <sup>26</sup> 2013	728	1495	48.7 (46.2-51.2)
Hypotension ± Vasopressor Therapy or Metabo	lic Abnormali	ties	
Peake et al, <sup>36</sup> 2009	75	324	23.1 (18.6-27.7)
Hypotension or Vasopressor Therapy			
Dahmash et al, <sup>59</sup> 1993	14	36	38.9 (23.0-54.8)
McLauchlan et al, <sup>58</sup> 1995	73	101	72.3 (63.5-81.0)
Pittet et al, <sup>57</sup> 1995	7	12	58.3 (30.4-86.2)
Schoenberg et al, <sup>53</sup> 1998	32	80	40.0 (29.3-50.7)
Engel et al, <sup>42</sup> 2007	119	190	62.6 (55.8-69.5)
Esteban et al, <sup>41</sup> 2007	27	59	45.8 (33.1-58.5)
Khwannimit and Bhuayanontachai, 37 2009	164	303	54.1 (48.5-59.7)
Moore et al, <sup>33</sup> 2011	22	61	36.1 (24.0-48.1)
Zahar et al, <sup>30</sup> 2011 (community)	215	530	40.6 (36.3-44.8)
Zahar et al, <sup>30</sup> 2011 (ICU)	123	232	53.0 (47.1-59.0)
Zahar et al, <sup>30</sup> 2011 (nosocomial)	233	580	40.2 (36.1-44.2)
Klein Klowenberg et al, <sup>7</sup> 2012	29	47	61.7 (47.8-75.6)
Park et al, <sup>28</sup> 2012	228	740	30.8 (27.5-34.1)
Hypotension or Serum Lactate Any Value or Va	sopressor The	rapy	
Liu et al, <sup>21</sup> 2014	827	2536	32.6 (30.8-34.4)
SSC database, <sup>16</sup> 2016 <sup>b</sup>	6556	18840	34.8 (34.1-35.5)
International Classification of Diseases Codes			
Annane et al, <sup>51</sup> 2003	13269	26172	50.7 (50.1-51.3)
Flaatten, <sup>50</sup> 2004	457	1562	29.3 (27.1-31.6)
Whittaker et al, <sup>24</sup> 2013	117	321	36.4 (31.2-41.7)
Serum Lactate Level >4 mmol/L			
Levy et al, <sup>6</sup> 2010	242	811	29.8 (26.7-33.0)
Phua et al, <sup>32</sup> 2011	219	466	47.0 (42.0-52.0)
<b>Overall</b> ( <i>I</i> <sup>2</sup> = 99.5%; <i>P</i> = .000)			46.5 (42.7-50.3)





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between groups for mortality increased mortality compared to other groups

Cohorts <sup>a</sup>	Lactate Category, mmol/L <sup>b</sup>	No. (% of total) [n = 18 840]	Acute Hospital Mortality, No. (%) [95% CI]	χ <sup>2</sup> Test for Tre	16/3/16 10:51 μμ	
Group 1 (hypotensive after fluids)						
and vasopressor therapy and serum lactate levels >2 mmol/L)	>2 to ≤3	2453 (13.0)	818 (33.3) [31.5-35.3]	<.001	1 [Reference]	
	>3 to ≤4	1716 (9.1)	621 (36.2) [33.9-38.5]			
	>4	4351 (23.1)	2163 (49.7) [48.2-51.2]			
	All	8520 (45.2)	3602 (42.3) [41.2-43.3]			
Group 2 (hypotensive after fluids and vasopressor therapy and serum lactate levels ≤2 mmol/L)	≤2	3985 (21.2)	1198 (30.1) [28.6-31.5]	NA <sup>d</sup>	0.57 (0.52-0.62)	<.001
Group 3 (hypotensive after fluids						
and no vasopressors and serum lactate	>2 to ≤3	69 (0.4)	15 (21.7) [12.7-33.3]	.04	0.65 (0.47-0.90)	.009
	>3 to ≤4	57 (0.3)	14 (24.6) [14.1-37.8]			
	>4	97 (0.5)	35 (36.1) [26.6-46.5]			
	All	223 (1.2)	64 (28.7) [22.9-35.1]			
Group 4 (serum lactate levels >2 mmol/L						
and no hypotension after fluids	>2 to ≤3	860 (4.6)	179 (20.8) [18.1-23.7]	<.001	0.71 (0.62-0.82)	<.001
	>3 to ≤4	550 (2.9)	105 (19.1) [15.9-22.6]			
	>4	1856 (9.9)	555 (29.9) [27.8-32.0]			
	All	3266 (17.3)	839 (25.7) [24.2-27.2]			
Group 5 (serum lactate levels between						
2-4 mmol/L and no hypotension before fluids	>2 to ≤3	1624 (8.6)	489 (30.1) [27.9-32.4]	NA <sup>d</sup>	0.77 (0.66-0.90)	.001
anu no vasopressors)	>3 to ≤4	1072 (5.7)	313 (29.2) [26.5-32.0]			
	>4	790 <sup>e</sup>				
	All	2696 (14.3)	802 (29.7) [28.0-31.5]			
Group 6 (hypotensive after fluids and no vasopressors and serum lactate ≤2 mmol/L)	≤2	150 (0.8)	28 (18.7) [12.8-25.8]	NA <sup>d</sup>	0.32 (0.20-0.51)	<.001

Abbreviations: NA, not available; OR, odds ratio.

<sup>c</sup> Refers to the adjusted OR generated using generalized estimating equation regression model (eTable7 in the Supplement).

SI conversion factor: To convert serum lactate values to mg/dL, divide by 0.111.

<sup>a</sup> Mean arterial pressure less than 65 mm Hg was used to define hypotension. "After fluids" was defined using the field "crystalloids" coded as a binary term within the Surviving Sepsis Campaign database.

<sup>b</sup> Using  $\chi^2$  tests, trends in mortality across serum lactate categories within groups (>2 to  $\leq$ 3 mmol/L; >3 to  $\leq$ 4 mmol/L and >4 mmol/L) were assessed.

 $^d \chi^2$  test for trend could only be performed if there were 3 or more serum lactate categories.

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<sup>e</sup> Excluded from full case analysis.



### Figure 3. Selection of Surviving Sepsis Campaign Database Cohort





Special Communication | CARING FOR THE CRITICALLY ILL PATIENT

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### Predictive Validity of Potential Septic Shock Groups

Of the 6 groups of potential patients with septic shock (Table 3), the most prevalent was group 1 (hypotension + vasopressor) therapy + serum lactate level >2 mmol/L) (n = 8520); followed by groups 2 (n = 3985) and 4 (n = 3266). Crude hospital mortality rates in these 3 groups were 42.3%, 30.1%, and 25.7%, respectively. Statistically significant increasing trends in crude mortality were observed over increasing serum lactate level categories within groups ( $\chi^2$  test of trend: P < .001 for groups 1 and 4, P = .04 for group 3). The adjusted OR for hospital mortality using group 1 for reference was significantly lower in all other groups (P < .01 for groups 2 to 6), suggesting that group 1 represents a distinct subpopulation with a significantly greater risk of death (eTable 7 in the Supplement). By a majority (cumulative first choice, 72.2%; second choice, 55.6%) (eTable 4 in the Supplement), the task force agreed that group 1 was most consistent with the proposed septic shock definition, thus generating the new septic shock criteria.



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Derivation of Serum Lactate Cutoff Value and Missing Data Analysis In the generalized estimating equation model (shown in eTable 8 in the Supplement), serum lactate level was associated with mortality, and the adjusted OR for hospital mortality increased linearly with increasing serum lactate level. An increase in serum lactate level from 2 to 10 mmol/L increased the adjusted OR for hospital mortality from 1.4 (95% CI, 1.35-1.45) to 3.03 (95% CI, 2.68-3.45) (referent lactate = 1; Figure 4). A serum lactate level greater than 2 mmol/L was chosen as the preferred cutoff value for the new septic shock criteria, the rationale being the trade-off between highest sensitivity (82.5%) when using the n = 18840 subset, and 74.9% when using patients in groups 1 and 2 combined [n = 12 475]), and the decision from the Delphi process to identify the lowest serum lactate level independently associated with a greater risk of death (OR of 1.4 at a lactate value of 2 mmol/L) (Table 4; eTable 9, eFigure 1, and eFigure 2 in the Supplement).



Figure 4. Serum Lactate Level Analysis



### Cutoff value

Adjusted odds ratio for actual serum lactate levels for the entire septic shock cohort (N = 18 840). The covariates used in the regression model include region (United States and Europe), location where sepsis was suspected (emergency department, ward, or critical care unit), antibiotic administration, steroid use, organ failures (pulmonary, renal, hepatic, and acutely altered mental state), infection source (pneumonia, urinary tract infection, abdominal, meningitis, and other), hyperthermia (>38.3°C), hypothermia (<36°C), chills with rigor, tachypnea (>20/min), leukopenia (<4000 cells/µL), hyperglycemia (plasma glucose >120 mg/dL [6.7 mmol/L]), platelet count <100 ×10<sup>3</sup>/µL, and coagulopathy (eMethods 3 in the Supplement). The adjusted odds ratio (OR) for the 6 groups presented in eTable 7 in the Supplement and the adjusted OR for the individual variables (lactate, vasopressor therapy, and fluids) are reported in eTable 8 in the Supplement. To convert serum lactate values to mg/dL, divide by 0.111. Table 4. Characteristics of Serum Lactate Level Cutoff Values for Complete Case Analysis and Imputation Analysis Using Surviving Sepsis Campaign Database

	Serum Lactate Level, mmol/L						
	>2		>3		>4		
Characteristic	Died/Total	% (95% CI)	Died/Total	% (95% CI)	Died/Total	% (95% CI)	
Complete Case Analysis (n = 1	.8 795)						
Hospital mortality, %	5757/18795	30.6 (29.9-31.4)	6101/18795	32.5 (31.8-33.2)	6456/18975	34.3 (33.7-35.0)	
Sensitivity, %	5372/6509	82.5 (81.6-83.4)	3779/6509	58.1 (56.8-59.3)	2811/6509	43.2 (42.0-44.4)	
Specificity, %	2748/12 286	22.4 (21.6-23.1)	6418/12 286	52.2 (51.4-53.1)	8564/12 286	69.7 (68.9-70.5)	
PPV, %	5372/14910	36.0 (35.3-36.8)	3779/9647	39.2 (38.2-40.2)	2811/6533	43.0 (41.8-44.2)	
NPV, %	2748/3885	70.7 (69.3-72.2)	6418/9148	70.1 (69.2-71.1)	8564/12 286	69.8 (69.0-70.7)	
Imputed Missing Serum Lacta	te Level (n = 22 182	?)					
Hospital mortality, %	6965/22 182	31.4 (30.8-32.0)	7363/22 182	33.2 (32.6-33.8)	7772/22 182	35.0 (34.4-35.7)	
Sensitivity, %	6457/7748	83.3 (82.5-84.2)	4461/7748	57.6 (56.5-58.7)	2931/7748	37.8 (36.7-38.9)	
Specificity, %	3341/14 434	23.1 (22.5-23.8)	7833/14434	54.3 (53.5-55.1)	10801/14434	74.8 (74.1-75.5)	
PPV, %	6457/17 550	36.8 (36.1-37.5)	4461/11062	40.3 (39.4-41.2)	2931/6564	44.6 (43.4-45.8)	
NPV, %	3341/4634	72.1 (70.8-73.4)	7833/11 120	70.4 (69.6-71.3)	10801/15618	69.2 (68.4-69.9)	

Abbreviations: NPV, negative predictive value; PPV, positive predictive value.

SI conversion factor: To convert serum lactate values to mg/dL, divide by 0.111.



### Discussion

The systematic review illustrated the variability in criteria currently used to identify septic shock, whereas the metaanalysis demonstrated the heterogeneity in mortality. Informed by this systematic review, a Delphi process was used to reach a consensus definition of septic shock and related clinical criteria. Three large data sets were then used to determine the predictive validity of these criteria. Septic shock was defined as a subset of sepsis in which circulatory, cellular, and metabolic abnormalities are associated with a greater risk of mortality than sepsis alone. The clinical criteria representing this definition were the need for vasopressor therapy to maintain a

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Underlying methodology	What should be considered	Recommended process
A (figh) KC1s B (moderate) downgraded RCTs or upgraded observational studies C (low) well-done observational studies with control RCTs D (very low) downgraded controlled studies or expert opinion	High or moderate evidence (is there high or moderate quality evidence?)	The higher the quality of evidence, the more likely a strong recommendation.
based on other evidence	Certainty about the balance of	The larger the difference
Factors that may decrease the strength of evidence	benefits versus harms and	between the desirable and
1. Poor quality of planning and implementation of available RCTs, suggesting high likelihood of bias	burdens (is there certainty?)	undesirable consequences and the certainty around that
2. Inconsistency of results, including problems with subgroup analyses		difference, the more likely a strong recommendation. The
3. Indirectness of evidence (differing population, intervention, control, outcomes, comparison)		smaller the net benefit and the lower the certainty for that
4. Imprecision of results		benefit, the more likely a weak
5. High likelihood of reporting bias		recommendation
Main factors that may increase the strength of evidence	Certainty in or similar values (is	The more certainty or similarity
1. Large magnitude of effect (direct evidence, relative risk $> 2$ with no plausible confounders)	there certainty or similarity?)	in values and preferences, the more likely a strong
2. Very large magnitude of effect with relative risk $> 5$ and no		recommendation
threats to validity (by two levels)	Resource implications (are	The lower the cost of an
3. Dose–response gradient	resources worth expected	intervention compared to the
RCT randomized controlled trial	benefits?)	alternative and other costs related to the decision-i.e.
		fewer resources consumed—
		the more likely a strong
The GRADE system classifies recommendations as		recommendation

 Table 3 Determination of the quality of evidence

Table 4 Factors determining strong versus weak recommendation

R. P. Dellinger Mitchell M. Levy Andrew Rhodes Djillali Annane Herwig Gerlach Steven M. Onal

### Surviving Sepsis Campaign: International Guidelines for Management of Severe Sepsis and Septic Shock, 2012

 Table 5
 Recommendations: initial resuscitation and infection issues

- A. Initial resuscitation
- Protocolized, quantitative resuscitation of patients with sepsis-induced tissue hypoperfusion (defined in this document as hypotension persisting after initial fluid challenge or blood lactate concentration ≥4 mmol/L). Goals during the first 6 h of resuscitation:

   (a) Central venous pressure 8–12 mmHg
- (b) Mean arterial pressure  $(MAP) \ge 65 \text{ mmHg}$
- (c) Urine output  $\geq 0.5 \text{ mL kg}^{-1} \text{ h}$
- (d) Central venous (superior vena cava) or mixed venous oxygen saturation 70 or 65 %, respectively (grade 1C)
- 2. In patients with elevated lactate levels targeting resuscitation to normalize lactate as rapidly as possible (grade 2C)
- B. Screening for sepsis and performance improvement
- 1. Routine screening of potentially infected seriously ill patients for severe sepsis to allow earlier implementation of therapy (grade 1C)
- 2. Hospital-based performance improvement efforts in severe sepsis (UG)
- C. Diagnosis
- 1. Cultures as clinically appropriate before antimicrobial therapy if no significant delay (>45 min) in the start of antimicrobial(s) (grade 1C). At least 2 sets of blood cultures (both aerobic and anaerobic bottles) be obtained before antimicrobial therapy with at least 1 drawn percutaneously and 1 drawn through each vascular access device, unless the device was recently (<48 h) inserted (grade 1C)
- 2. Use of the 1,3 β-D-glucan assay (grade 2B), mannan and anti-mannan antibody assays (2C), if available and invasive candidiasis is in differential diagnosis of cause of infection.
- 3. Imaging studies performed promptly to confirm a potential source of infection (UG)
- D. Antimicrobial therapy

D. Antimicrobial therapy

- 1. Administration of effective intravenous antimicrobials within the first hour of recognition of septic shock (grade 1B) and severe sepsis without septic shock (grade 1C) as the goal of therapy
- 2a. Initial empiric anti-infective therapy of one or more drugs that have activity against all likely pathogens (bacterial and/or fungal or viral) and that penetrate in adequate concentrations into tissues presumed to be the source of sepsis (grade 1B)
- 2b. Antimicrobial regimen should be reassessed daily for potential deescalation (grade 1B)
- 3. Use of low procalcitonin levels or similar biomarkers to assist the clinician in the discontinuation of empiric antibiotics in patients who initially appeared septic, but have no subsequent evidence of infection (grade 2C)
- 4a. Combination empirical therapy for neutropenic patients with severe sepsis (grade 2B) and for patients with difficult to treat, multidrug-resistant bacterial pathogens such as *Acinetobacter* and *Pseudomonas spp*. (grade 2B). For patients with severe infections associated with respiratory failure and septic shock, combination therapy with an extended spectrum beta-lactam and either an aminoglycoside or a fluoroquinolone is for *P. aeruginosa* bacteremia (grade 2B). A combination of beta-lactam and macrolide for patients with septic shock from bacteremic *Streptococcus pneumoniae* infections (grade 2B)
- 4b. Empiric combination therapy should not be administered for more than 3–5 days. De-escalation to the most appropriate single therapy should be performed as soon as the susceptibility profile is known (grade 2B)
- 5. Duration of therapy typically 7–10 days; longer courses may be appropriate in patients who have a slow clinical response, undrainable foci of infection, bacteremia with *S. aureus*; some fungal and viral infections or immunologic deficiencies, including neutropenia (grade 2C)
- 6. Antiviral therapy initiated as early as possible in patients with severe sepsis or septic shock of viral origin (grade 2C)
- 7. Antimicrobial agents should not be used in patients with severe inflammatory states determined to be of noninfectious cause (UG) E. Source control
- 1. A specific anatomical diagnosis of infection requiring consideration for emergent source control be sought and diagnosed or excluded as rapidly as possible, and intervention be undertaken for source control within the first 12 h after the diagnosis is made, if feasible (grade 1C)
- 2. When infected peripancreatic necrosis is identified as a potential source of infection, definitive intervention is best delayed until adequate demarcation of viable and nonviable tissues has occurred (grade 2B)
- 3. When source control in a severely septic patient is required, the effective intervention associated with the least physiologic insult should be used (e.g., percutaneous rather than surgical drainage of an abscess) (UG)
- 4. If intravascular access devices are a possible source of severe sepsis or septic shock, they should be removed promptly after other vascular access has been established (UG)
- F. Infection prevention
- 1a. Selective oral decontamination and selective digestive decontamination should be introduced and investigated as a method to reduce the incidence of ventilator-associated pneumonia; This infection control measure can then be instituted in health care settings and regions where this methodology is found to be effective (grade 2B)
- 1b. Oral chlorhexidine gluconate be used as a form of oropharyngeal decontamination to reduce the risk of ventilator-associated pneumonia in ICU patients with severe sepsis (grade 2B)

**Table 6** Recommendations: hemodynamic support and adjunctive therapy

G. Fluid therapy of severe sepsis

- 1. Crystalloids as the initial fluid of choice in the resuscitation of severe sepsis and septic shock (grade 1B).
- 2. Against the use of hydroxyethyl starches for fluid resuscitation of severe sepsis and septic shock (grade 1B).
- 3. Albumin in the fluid resuscitation of severe sepsis and septic shock when patients require substantial amounts of crystalloids (grade 2C).
- 4. Initial fluid challenge in patients with sepsis-induced tissue hypoperfusion with suspicion of hypovolemia to achieve a minimum of 30 mL/kg of crystalloids (a portion of this may be albumin equivalent). More rapid administration and greater amounts of fluid may be needed in some patients (grade 1C).
- 5. Fluid challenge technique be applied wherein fluid administration is continued as long as there is hemodynamic improvement either based on dynamic (e.g., change in pulse pressure, stroke volume variation) or static (eg. arterial pressure, heart rate) variables (UG).

H. Vasopressors

- 1. Vasopressor therapy initially to target a mean arterial pressure (MAP) of 65 mm Hg (grade 1C).
- 2. Norepinephrine as the first choice vasopressor (grade 1B).
- 3. Epinephrine (added to and potentially substituted for norepinephrine) when an additional agent is needed to maintain adequate blood pressure (grade 2B).
- 4. Vasopressin 0.03 units/minute can be added to norepinephrine (NE) with intent of either raising MAP or decreasing NE dosage (UG).
- 5. Low dose vasopressin is not recommended as the single initial vasopressor for treatment of sepsis-induced hypotension and vasopressin doses higher than 0.03-0.04 units/minute should be reserved for salvage therapy (failure to achieve adequate MAP with other vasopressor agents) (UG).
- 6. Dopamine as an alternative vasopressor agent to norepinephrine only in highly selected patients (eg, patients with low risk of tachyarrhythmias and absolute or relative bradycardia) (grade 2C).
- 7. Phenylephrine is not recommended in the treatment of septic shock except in circumstances where (a) norepinephrine is associated with serious arrhythmias, (b) cardiac output is known to be high and blood pressure persistently low or (c) as salvage therapy when combined inotrope/vasopressor drugs and low dose vasopressin have failed to achieve MAP target (grade 1C).
- 8. Low-dose dopamine should not be used for renal protection (grade 1A).
- 9. All patients requiring vasopressors have an arterial catheter placed as soon as practical if resources are available (UG).
- I. Inotropic therapy
- 1. A trial of dobutamine infusion up to 20 micrograms/kg/min be administered or added to vasopressor (if in use) in the presence of (a) myocardial dysfunction as suggested by elevated cardiac filling pressures and low cardiac output, or (b) ongoing signs of hypoperfusion, despite achieving adequate intravascular volume and adequate MAP (grade 1C).
- 2. Not using a strategy to increase cardiac index to predetermined supranormal levels (grade 1B).
- J. Corticosteroids
- 1. Not using intravenous hydrocortisone to treat adult septic shock patients if adequate fluid resuscitation and vasopressor therapy are able to restore hemodynamic stability (see goals for Initial Resuscitation). In case this is not achievable, we suggest intravenous hydrocortisone alone at a dose of 200 mg per day (grade 2C).
- 2. Not using the ACTH stimulation test to identify adults with septic shock who should receive hydrocortisone (grade 2B).
- 3. In treated patients hydrocortisone tapered when vasopressors are no longer required (grade 2D).
- 4. Corticosteroids not be administered for the treatment of sepsis in the absence of shock (grade 1D).
- 5. When hydrocortisone is given, use continuous flow (grade 2D).



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November 1, 2016

# **Effect of Hydrocortisone on Development of Shock Among Patients With Severe Sepsis** The HYPRESS Randomized Clinical Trial

Didier Keh, MD<sup>1</sup>; Evelyn Trips<sup>2</sup>; Gernot Marx, MD<sup>3</sup>; <u>et al</u>

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### **Key Points**

**Question** Does adjunctive early hydrocortisone therapy prevent the development of septic shock in patients with severe sepsis who are not in shock?

**Findings** In this randomized clinical trial that included 380 adults, occurrence of septic shock was not significantly different between patients who received hydrocortisone or placebo (21.2% vs 22.9%, respectively).

**Meaning** Administration of hydrocortisone did not prevent the development of shock in patients with severe sepsis.



### SYSTEMATIC REVIEW



### Unexplained mortality differences between septic shock trials: a systematic analysis of population characteristics and control-group mortality rates

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### Abstract

**Purpose:** Although the definition of septic shock has been standardized, some variation in mortality rates among clinical trials is expected. Insights into the sources of heterogeneity may influence the design and interpretation of septic shock studies. We set out to identify inclusion criteria and baseline characteristics associated with between-trial differences in control group mortality rates.

**Methods:** We conducted a systematic review of RCTs published between 2006 and 2018 that included patients with septic shock. The percentage of variance in control-group mortality attributable to study heterogeneity rather than chance was measured by l<sup>2</sup>. The association between control-group mortality and population characteristics was estimated using linear mixed models and a recursive partitioning algorithm.

**Results:** Sixty-five septic shock RCTs were included. Overall control-group mortality was 38.6%, with significant heterogeneity ( $l^2 = 93\%$ , P < 0.0001) and a 95% prediction interval of 13.5–71.7%. The mean mortality rate did not differ between trials with different definitions of hypotension, infection or vasopressor or mechanical ventilation inclusion criteria. Population characteristics univariately associated with mortality rates were mean Sequential Organ Failure Assessment score (standardized regression coefficient ( $\beta$ ) = 0.57, P = 0.007), mean serum creatinine ( $\beta$  = 0.48, P = 0.007), the proportion of patients on mechanical ventilation ( $\beta$  = 0.61, P < 0.001), and the proportion with vasopressors ( $\beta$  = 0.57, P = 0.002). Combinations of population characteristics selected with a linear model and recursive partitioning explained 41 and 42%, respectively, of the heterogeneity in mortality rates.

**Conclusions:** Among 65 septic shock trials, there was a clinically relevant amount of heterogeneity in control group mortality rates which was explained only partly by differences in inclusion criteria and reported baseline characteristics.

Keywords: Septic shock, Heterogeneity, Clinical trials, Methodology, Meta-research, Machine learning

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### Introduction

The fundamental criteria from the consensus definitions of septic shock are used to select patients for inclusion in clinical studies [1–4]. While the mortality rate of septic shock was found to be 46% (95% confidence interval (CI) 43–50%) in a meta-analysis of observational cohorts [5], randomized controlled trials report more diverse numbers. For example, two high-profile septic shock trials published a year apart reported control group mortality rates as disparate as 16% [6] and 80% [7]. Despite the seemingly wide range of mortality rates there has not yet been a systematic inquiry into its patterns and possible causes.

Identifying the correct patient population to benefit from a specific therapy has been recognized as an essential condition for improving critical care research [8–10]. Yet large unexplained mortality differences among trials that all aim to include septic shock patients may hamper reproducibility and generalizability. Insights into the magnitude and sources of between-trial heterogeneity are therefore valuable in the design, reporting, and interpretation of septic shock trials. For example, incorrect prediction of baseline mortality rates has been identified as a major reason for negative critical care trials, as a discrepancy between expected and observed event rates often leads to underpowered studies [11].

We sought to quantify between-trial heterogeneity and identify inclusion criteria and population characteristics associated with differences in control group mortality rates.

### Methods

After a systematic search to identify all trials published in the past decade that aimed to include patients with septic shock, we used linear mixed models to estimate the total heterogeneity in control group mortality rates and its association with reported baseline characteristics. Using both a multivariate linear model and a machine learning algorithm, we estimated the proportion of heterogeneity that can be explained by population characteristics.

The review protocol was prospectively registered [12] and adheres to the PRISMA checklist [13], which is included in the electronic supplementary material (ESM). Study screening, application of the inclusion- and exclusion criteria and data-extraction were performed independently by two reviewers (HJdG and JP). Conflicting entries were resolved by consensus.

### Inclusion criteria and search strategy

PubMed, Embase, and the Cochrane Central Register of Controlled Trials were queried using the search term ["septic shock" AND (random\* or rct)]. Embase was additionally queried using the search term "septic shock" with the randomized controlled trial filter activated. The queries were limited to publications from 1 January 2006 and the queries were last performed on 20 January 2018.

We limited the search to trials published between 2006 and 2018 as a compromise between the number of eligible studies and secular trends in clinical practice, research practice, and reporting standards. Publications from 2006 and later had sufficient lead time to incorporate the 2004 update of the Surviving Sepsis Campaign guidelines [4].

Eligible for inclusion were parallel-group randomized controlled trials with adult patients in septic shock according to the published consensus definitions or Surviving Sepsis Campaign guidelines [1, 2, 4]. Trials were excluded if the report was not written in English, if it was only available in abstract, if no baseline characteristics were reported, or if no mortality outcome was reported. Trials that aimed to include a specific subcategory of septic shock patients (e.g. "septic shock patients requiring renal replacement therapy") were also excluded, as these would be a major source of between-trial heterogeneity.

#### Identification of the control group and variables of interest

Because the nature of the randomized intervention could contribute to heterogeneity, we focused on the control groups. For each trial, we identified the control group as defined by the authors as 'control group,' 'usual care group,' or a variation thereof. When no control group could be identified (in a comparison of two usual care therapies) we defined the control group as the means of the two groups in terms of sample size, mortality, and baseline characteristics. A sensitivity analysis was performed towards this construct by analyzing whether trials with and without specifically defined control groups differed in terms of mean mortality or the amount of between-trial heterogeneity.

For each trial, we recorded the type of intervention, single- or multicenter design, and the primary endpoint. Trials were graded according to the Jadad scale [14]. For the control group in each trial, we recorded the sample size, the reported baseline characteristics, and the mortality rates.

### Estimation of heterogeneity in mortality rates and associations with population characteristics

We used 28-day mortality throughout all analyses. For trials that did not report this outcome, we estimated 28-day mortality based on reported hospital, ICU, or 90-day mortality using linear regression with data from trials that reported both 28-day and another mortality measure.

To analyze mortality rates across trials we used a random-effects meta-regression model with the log odds of mortality as dependent variable and a random intercept for each study. Each trial was weighted by the inverse of the sampling variance of the mortality rates. A maximum likelihood estimator was used to estimate the mean mortality (random effects pooled estimate), the between-study standard deviation due to heterogeneity  $(\tau)$ , and the percentage of variation due to heterogeneity rather than change  $(I^2)$ . To quantify between-trial heterogeneity, we report the 95% prediction interval (mean mortality  $\pm$  1.96  $\tau$ ), which represents the distribution of estimated future mortality rates based on observed mortalities weighted by sampling variance (trial size) and corrected for random chance [15]. In the absence of between-study heterogeneity, the 95% prediction interval is equal to the 95% confidence interval, but when significant heterogeneity is present the prediction interval estimates the bandwidth of expected mortality rates from similar studies [15, 16]. In other words, the 95% prediction interval can be thought of as the estimate of true between-study distribution of mortality rates. The prediction interval can therefore be used to guide power calculations for future studies [16].

The between-trial heterogeneity in mortality rates was calculated for subcategories of trials employing different inclusion criteria: confirmed or suspected infection; confirmed infection only; different definitions of hypotension; mandatory hyperlactatemia; mandatory vasopressor therapy; and mandatory mechanical ventilation. Differences in mortality rates between subcategories were calculated by addition of dummy variables to the mixed-effects model.

To estimate the association between study and population characteristics and mortality, these variables were added to the model as covariates. Residuals were checked for normality with Q–Q plots, and the goodness of fit of the log-linear model was compared with quadratic and power models by selecting the model with the lowest Akaike information criterion (AIC). To facilitate comparisons between variables, we report standardized regression coefficients ( $\beta$ ) and the proportion of between-trial variability in mortality explained by the population variable (unadjusted R<sup>2</sup>) for all univariate analyses.

### Predicting mortality rates using a linear model and recursive partitioning

We then constructed a comprehensive model to predict between-study differences in mortality. Population variables that were reported by at least 25% of the included trials with a univariate regression  $\mathbb{R}^2 \ge 0.10$  were included as regressors in a multivariate model and removed in a stepwise manner for P values  $\ge 0.05$ . The threshold  $\mathbb{R}^2$  of 0.10 was a compromise between the number of variables and the limited number of observations. This model

selection process was not prospectively protocolized as the number of eligible variables could not be estimated a priori. Multiple imputation (generating 20 datasets) with predictive mean matching was used for missing observations (i.e., missing population characteristics). The imputation methods are further described in section 7 of the ESM.

As a complementary approach to predict 28-day mortality rates from population characteristics, we constructed a regression tree model based on recursive partitioning (a machine learning algorithm) [17, 18] for its ability to handle partially missing observations (obviating the need for imputation) and its robustness to nonlinear relations. We set up the model to predict 28-day mortality based on all inclusion criteria and population characteristics. In short, the recursive partitioning algorithm selected the most informative variable, which was then 'split' at the value that best differentiates low from high mortality. The algorithm then selected the most informative variable for each of the two resulting subgroups, and split it again. When a splitting variable was missing for a specific trial, a surrogate variable (the variable most closely correlated to the splitting variable) was used. After multiple splits, this recursive partitioning resulted in a regression tree (similar to a decision tree) with subgroups of trials ranked from low to high expected mortality. R<sup>2</sup> represents the variance in mortality explained by the decision tree. Overfitting was examined using the cross-validated error.

For all analyses, P < 0.05 was considered significant. The analyses were performed in *R* version 3.4.2 using the *metafor*, *mice* and *rpart* packages [19–21].

#### Results

#### Characteristics of the included trials

The search resulted in 65 trials that met all inclusion and exclusion criteria (eFigure 1 in the ESM), representing a total of 8634 control group patients [6, 7, 22–84]. A list of excluded trials is available in the ESM. The trial characteristics are presented in Table 1.

Twenty trials (31%) did not report 28-day mortality but only hospital mortality, ICU mortality, or 90-day mortality. Using trials that reported multiple mortality measures, 28-day mortality was estimated as a linear function of hospital mortality, ICU mortality, or 90-day mortality ( $\mathbb{R}^2$  values 0.99, 0.98, and 0.98, respectively). The estimates and validation plots are presented in eTable 1 and eFigure 2 of the ESM.

In 14 trials (21%) the control group could not be identified because two usual care therapies were compared. For these trials, the control group characteristics and mortality rates were defined as the means of the two treatment

### Table 1 Characteristics of included trials

	No. (%) or median (IQR)
Number of included trials	65
Control group sample size: median (IQR)	34 (20–100)
Multicenter trials: n (%)	28 (43)
Trial country: n (%)	
France	12 (18)
China	9 (14)
Italy	8 (12)
USA	6 (9)
India	3 (5)
The Netherlands	3 (5)
UK	3 (5)
Other countries (1 each)	13 (20)
Multinational trials	9 (14)
Trial intervention: n (%)	
Drug	44 (68)
Treatment bundle	14 (21)
Device	7 (11)
Primary endpoint: n (%)	
Mortality	21 (32)
Other	32 (49)
Not specified	12 (18)
Jadad scale: median (IQR)	3 (2–4)
Jadad scale components: n (%)	
Randomization	65 (100)
Randomization appropriate	45 (69)
Blinding	23 (35)
Blinding appropriate	19 (29)
Description of withdrawals and dropouts	42 (65)

IQR Interquartile range

groups. None of these 14 trials reported significant mortality differences between the treatment groups.

### The distribution of mortality rates

The control group mortality rates ranged between 13.8 and 84.6%, with a random-effects estimated mean mortality rate of 38.6%. There was significant heterogeneity among trials ( $I^2 = 93\%$ ,  $\tau = 0.710$ , p < 0.0001), and the 95% prediction interval was 13.5–71.7%.

Figure 1 shows the mortality rates of trials categorized by inclusion criteria. The mean mortality rate did not differ between trials with different definitions of hypotension, infection (confirmed vs. suspected), or vasopressor or mechanical ventilation inclusion criteria. There were no significant differences in mean mortality rate or in heterogeneity between large vs. small trials, monocenter vs. multicenter trials, unblinded vs blinded trials, high-quality trials vs. low-quality trials, or trials with vs. without a specifically defined control group (eTable 2 in the ESM).

The exclusion criteria employed in the trials were too diverse for statistical analysis, but the total number of exclusion criteria (ranging from 0 to 30) was inversely associated with the mortality rate ( $\beta = -0.375$ ,  $R^2 = 0.14$ , P = 0.007).

The heatmap in Fig. 2 provides an overview of the between-trial differences in mortality rates and population characteristics. The log-linear associations between the mortality rate and reported control group baseline characteristics are presented in Table 2 (goodness-of-fit statistics are reported in eTable 3 in the ESM). There was no significant decrease in mortality over the period 2006–2018, with only ( $R^2$ ) 4% of heterogeneity explained by the year of publication (Table 2, eFigure 3). Baseline variables that were univariately associated with mortality were: mean Sequential Organ Failure Assessment (SOFA) score, the proportion of patients on mechanical ventilation, the proportion of patients on vasopressors, and mean serum creatinine. Regression plots of selected associations are shown in eFigure 3 of the ESM.

#### Predicting mortality rates from population characteristics

Details of the variable selection process for the multivariate model are available in section 7 of the ESM. Significant independent variables in the final multivariate model were: baseline mean SOFA score ( $\beta = 0.39$ , standardized standard error (SSE) = 0.17, P = 0.019), the proportion of patients on mechanical ventilation ( $\beta = 0.42$ , SSE = 0.18, P = 0.019), and mean serum creatinine ( $\beta = 0.31$ , SSE = 0.10, P = 0.0015). The multivariate model R<sup>2</sup> was 0.41 with significant residual heterogeneity (I<sup>2</sup> = 82%,  $\tau = 0.544$ , P < 0.0001). Figure 3 shows the predicted and actual mortality rates of the included trials.

The recursive partitioning algorithm resulted in a regression tree with the following variables as informative determinants of the mortality rate: mean age (split at 64.8 years); the proportion of patients with a respiratory infection (split at 54.5%); the proportion of patients on mechanical ventilation (split at 74.3%); and the proportion of male patients (splits at 63.8 and 53.8%). The  $R^2$  value of the regression tree was 0.42. The cross-validated relative error decreases to below the root (split 0) value, which indicates that the tree was not overfitted. The results from the regression tree analysis are further described in eFigures 4 and 5 of the ESM (section 7).

### Discussion

In this analysis of 65 septic shock trials published in the past decade, we found a statistically significant and clinically relevant amount of heterogeneity in control group mortality rates. The mean mortality rate was 38.6% with



intervals. The 95% prediction intervals (dashed lines) represents the estimated between-trial variability in mortality rates after adjusting for random chance and sample size. I<sup>2</sup> represents the proportion of between-trial variability that cannot be explained by chance. There were no significant differences in mean mortality rates between inclusion criteria. *MAP* mean arterial pressure, *SBP* systolic blood pressure

estimated 95% prediction limits of 13.5–71.7%, revealing a wide range in underlying mortality rates after discounting the effects of random change and small trials.

In contrast to findings from large observational studies that the mortality of sepsis has decreased in the past decade, we found only a small nonsignificant decline in the period 2006–2018 [85, 86]. Different inclusion definitions of septic shock did not affect mean mortality rates, but a higher total number of exclusion criteria was associated with lower mortality. We used three statistical methods to analyze the association between population characteristics and mortality.

The univariate associations reflect how the reader of a trial report could interpret the population characteristics in relation to the mortality rate, and shows that the proportion of ventilated patients, mean SOFA score, and the proportion of patients on vasopressor support were most informative (i.e. have highest standardized regression coefficients).

The multivariate linear model (with missing observations imputed) shows which combinations of characteristics were predictive of mortality if all trials hypothetically reported the same variables. A combination of three independently significant characteristics (mean SOFA score, proportion of ventilated patients, and mean creatinine) explained only 41% of the heterogeneity in mortality rates across trials.

The recursive partitioning algorithm, which is not limited by dependence on multiple imputation and the assumption of linearity, shows which characteristics were most informative, given that different trials report different characteristics. The resulting regression tree explained only 42% of the heterogeneity in mortality.

The linear model and the regression tree arrived at different predictor variables because the linear model is biased towards more informative linear associations, while the regression tree allows for nonlinear relations and is biased towards variables with less missing data.

In all, these results indicate that there are clinically significant between-trial differences in control group mortality rates, and that these differences are not associated with differences in inclusion criteria and only weakly associated with reported baseline characteristics. Visual inspection of the heatmap (Fig. 2) shows that there are no unambiguous patterns in the relation between population characteristics and mortality rates. This heterogeneity is reflected in our finding that different statistical methods result in different predictive variables.

### Possible sources of residual heterogeneity

Residual heterogeneity among trials may be caused by population differences in nutrition and socio-economic status, heterogenous exclusion criteria, incomplete reporting, between-trial differences in variable definitions, the timing of randomization, and differences in post-randomization co-interventions and standards of care.

We found that no single measure of chronic comorbidity was reported in more than 40% of the included trials and that characteristics of causative pathogens were





Table 2 Univariate associations between mortality rates and reported mean or median population characteristics

	Trials reporting variable (% of $n = 56$ )	Mean (SD)	Standardized regression coefficient $\beta$ (R <sup>2</sup> )	P value
Publication year	65 (100)	2013.3 (3.58)	- 0.19 (0.04)	0.197
Age, years	64 (98)	62.9 (3.80)	0.18 (0.03)	0.160
Male patients %	63 (97)	60.5 (5.80)	0.02 (0.00)	0.927
Comorbidity characteristics				
Charlson Comorbidity Index	5 (8)	1.90 (1.11)	0.52 (0.27)	0.183
From long-term care facility %	6 (9)	5.8 (5.6)	0.44 (0.20)	0.312
McCabe class I %	6 (9)	34.1 (15.2)	— 0.40 (0.16)	0.374
McCabe class II %	6 (9)	14.7 (12.9)	0.02 (0.00)	0.948
McCabe class III %	4 (6)	16.2 (15.0)	0.71 (0.50)	0.120
Diabetes mellitus %	23 (36)	24.4 (6.88)	0.01 (0.00)	0.856
Heart failure or coronary disease %	26 (40)	20.7 (8.7)	0.33 (0.11)	0.133
Chronic obstructive pulmonary disease %	25 (39)	15.1 (6.3)	0.04 (0.00)	0.911
Chronic renal disease %	21 (33)	7.6 (5.0)	0.06 (0.00)	0.773
Chronic liver disease %	17 (26)	5.5 (2.8)	0.25 (0.06)	0.320
Cancer %	20 (31)	21.2 (8.1)	0.19 (0.03)	0.426
Severity of illness scores				
APACHE II score	33 (51)	22.5 (3.65)	0.21 (0.05)	0.376
APACHE III score	1 (2)	-	-	-
APACHE IV score	1 (2)	-	-	-
SAPS II score	24 (37)	55.7 (4.42)	0.36 (0.13)	0.079
SAPS III score	3 (4)	77.6 (1.91)	0.01 (0.00)	0.644
SOFA score	37 (58)	9.59 (2.47)	0.57 (0.33)	0.007**
Characteristics of acute illness				
Medical (non-surgical) %	22 (34)	69.7 (13.1)	0.26 (0.07)	0.314
Time from diagnosis to randomization, hours	13 (20)	13.77 (8.84)	0.47 (0.22)	0.069
Mechanical ventilation %	33 (51)	78.1 (28.3)	0.61 (0.38)	0.0005***
Heart rate, 1/min	39 (60)	104 (8.8)	0.13 (0.02)	0.435
Mean arterial pressure, mmHg	43 (66)	70.7 (6.65)	0.06 (0.00)	0.561
Central venous pressure, mmHg	22 (34)	11.2 (2.21)	0.17 (0.03)	0.425
Vasopressor support %	38 (58)	84.6 (30.0)	0.57 (0.32)	0.0019**
Serum lactate, mmol/l	52 (80)	4.00 (1.28)	— 0.13 (0.02)	0.389
Serum creatinine, µmol/l	26 (40)	168 (31.1)	0.48 (0.23)	0.007**
Fluids before randomization, ml	19 (30)	3209 (1637)	0.31 (0.10)	0.194
Infection site characteristics				
Respiratory %	53 (82)	42.6 (13.7)	0.27 (0.08)	0.087
Abdominal %	51 (78)	24.0 (15.0)	0.06 (0.00)	0.686
Urogenital %	41 (63)	11.3 (5.7)	— 0.27 (0.07)	0.094
Central nervous system %	19 (30)	1.2 (1.6)	0.03 (0.00)	0.885
Skin and soft tissue %	28 (43)	6.8 (3.6)	— 0.09 (0.01)	0.803
Bloodstream %	32 (49)	12.9 (8.2)	- 0.11 (0.01)	0.487
Pathogen characteristics				
Gram-negative %	25 (39)	32.0 (16.1)	0.41 (0.17)	0.0573
Gram-positive %	22 (34)	24.6 (7.12)	- 0.41 (0.17)	0.083
Other pathogen %	22 (34)	44.0 (23.3)	- 0.13 (0.02)	0.473
Culture negative %	18 (28)	29.4 (8.3)	- 0.38 (0.14)	0.085

Univariate associations between control group mortality rate and commonly reported mean baseline characteristics. Associations were estimated using a weighted random-effects model with mortality on the log-odds scale. Some baseline characteristics were reported by a minority of trials, which resulted in low power to detect a significant association. R<sup>2</sup> can be interpreted as the proportion of heterogeneity that is explained by the population characteristic for the n trials that report that characteristic

APACHE Acute Physiology and Chronic Health Evaluation score, SAPS Simplified Acute Physiology score, SOFA Sequential Organ Failure Assessment score

reported in only 28–39% of trials. This compromised the power of our analysis to detect associations across all trials, but, more importantly, it also prevents readers of trial reports from evaluating and comparing populations among trials and from judging to what extent a trial population corresponds to the population under their care.

Another source of heterogeneity is the imprecise definition of many variables. It is unclear whether a variable like 'pre-existing kidney disease' in one trial has the same meaning as 'chronic renal insufficiency' in another trial. Minor variations in variable definitions and data capture methods have been shown to lead to significantly different septic shock populations and to inter-observer variability in severity-of-illness scoring systems [5, 87, 88]. The importance of this 'fine print' in defining a population does not receive due attention in the methods section of most trials.

The time of inclusion may be an additional source of heterogeneity. Patients recruited later after the diagnosis of septic shock have not responded to treatment in an earlier phase and are therefore likely to have a worse prognosis. Only 13 trials reported the time from diagnosis to randomization, and for those trials it explained 22% of the heterogeneity.

While we have focused on inclusion criteria and baseline characteristics, the prognosis of septic shock may be largely influenced by post-randomization standards of care and co-interventions. Unfortunately, co-interventions and (control group) treatment standards are often described as 'according to the Surviving Sepsis Campaign guidelines' or not discussed at all in trial reports. Variables describing important post-randomization interventions, such as red blood cell transfusions, vasopressor dose, or fluid balance were recently found to be reported in only 33, 17, and 13% of large septic shock trials, respectively [89].

We did not analyze the association between trial countries and the mortality rate because many countries are represented by a single trial in the present sample. Nevertheless, between-country differences in standards of care or access to early healthcare may account for part of the residual heterogeneity. Large international observational studies are a more appropriate instrument for the investigation of differences in mortality rates among countries.

### Implications for investigators and clinicians

Clinicians demand of clinical trials that they are relevant, reproducible, and generalizable to a clearly defined patient population. The results of this study indicate that many of the baseline characteristics upon which clinicians rely to gauge the applicability of trial results to their practice are in fact only weakly or not at all associated with mortality outcomes across trials.

The association between the number of exclusion criteria and mortality suggests that many seemingly inconsequential criteria together may have a significant effect on the composition of a trial population. Investigators should therefore be aware of this phenomenon in the design phase of a trial, as it affects the generalizability and external validity of trial results.

The wide prediction limits of control-group mortality have consequences for sample size calculations. Detecting a relative risk reduction of 25% with 80% power requires 245 patients if mortality is estimated to be 71.7%, while it requires 795 patients if control group mortality is 38.6% or 2980 patients if mortality is 13.5%. In practice, misestimation of the mortality rate by more than 7.5% occurred in 65% of critical care trials [11]. We therefore suggest that sample size calculations should not be based on the mean of reported control-group mortality rates in the literature but should be robust towards a wider range of expected event rates.

Reproducibility and generalizability also require a common phenomenological structure with respect to diagnostic definitions, inclusion criteria, patient characteristics, concomitant treatment, and outcomes. A recent review of large septic shock trials found that only half of the information deemed necessary for evaluation of the control group was reported in the investigated trials [89]. In the present study, we now find that many of the reported characteristics are not associated with controlgroup mortality rates, possibly due to variations in variable definitions.

The third consensus definitions for sepsis and septic shock were partly developed to harmonize the inclusion criteria for clinical studies [3]. We were unable to analyze a subset of trials with populations that might fit the Sepsis-3 septic shock definition, as none of the included trials employed both delta SOFA score and vasopressor inclusion criteria. We do note that SOFA score is independently associated with mortality rates, although baseline SOFA explains only 33% ( $\mathbb{R}^2$ ) of the variation in mortality rates in the 37 trials that report it. Furthermore, we found significant heterogeneity within subsets of trials employing similar inclusion criteria (Fig. 2).

We suggest that an international consensus is necessary to standardize variable definitions, data collection, and reporting of patient characteristics and outcomes for sepsis trials, as has been proposed before [89–92]. The feasibility of harmonizing study protocols has been demonstrated in three large trials investigating early goaldirected therapy [93]. The present results indicate that SOFA score, the proportion of ventilated patients, and creatinine independently reflect baseline risk across trials and should therefore be reported for each trial.



**Fig. 3** Included trials ordered by predicted control group mortality rate (diamonds). The predicted mortality rates were based on a multivariate weighted random-effects regression model with baseline mean Sequential Organ Failure Assessment (SOFA) score, the proportion of patients on mechanical ventilation, and mean serum creatinine as significant independent variables. The squares and brackets are the observed control-group mortality rates with 95% confidence interval. The figure illustrates that the model explained ( $R^2$ ) 41% of the variability in mortality rates, with significant residual heterogeneity (P < 0.0001). The red dots are the reported a-priori expected mortality rates used for sample size calculations

The results from this study also support the practice of data sharing, as we have shown that aggregated population characteristics are less informative than expected. Sharing individual patient data will not only increase the power to detect treatment effects across multiple studies but can also be used to test the generalizability of trial results vis-à-vis large cohorts with septic shock.

#### Strengths and limitations

This study was performed with a prospectively registered protocol and analysis plan. We chose to include only trials published between 2006 and 2018 to minimize the influence of long-term secular trends in septic shock diagnosis, treatment, and mortality [94, 95]. The search strategy was broad and comprehensive, but we excluded 40 trial reports not written in English, which compromised power and generalizability. We excluded trials that recruited only septic shock patients with specific organ dysfunction (such as kidney or liver failure) to rule out this source of between-trial heterogeneity.

For 20 trials, 28-day mortality was estimated using another reported mortality rate. Although the prediction equations were very precise ( $\mathbb{R}^2$  values  $\geq 0.98$ ), we cannot rule out the possibility that this influenced the results. Excluding these 20 trials would have eroded the power of the study.

Importantly, using study-level data means that, to avoid the ecological fallacy, we cannot make inferences about predictive characteristics at the individual patient level, although several predictor variables are known to be individually associated with mortality (e.g. high SOFA score as a risk factor [96, 97]).The fact that there was substantial variation in the reporting of baseline variables was an important finding in itself, but also limited our power to detect associations across trials. A more in-depth investigation into the heterogeneity among trial populations would require individual patient data, but we think that obtaining such data would lead to significant selection bias.

### Conclusion

Septic shock is a syndrome with various etiologies, biochemical characteristics, and phenotypes [9, 98]. Onto this inherently heterogeneous syndrome, a layer of investigator-induced heterogeneity is added when trials employ different inclusion criteria, report different variables, and use different variable definitions. This compounded complexity causes heterogeneity among trial populations that may go unnoticed. We have shown that control-group mortality rates are very dissimilar across trials, and that the majority of this heterogeneity remains unexplained after accounting for reported population characteristics. The lack of standardized reporting limits the usefulness of the variables explaining the mortality differences found in this study. In all, the substantial between-trial heterogeneity limits the reproducibility and generalizability of septic shock research and may inhibit the discovery of beneficial therapies for specific (sub)populations. The findings of this study therefore strongly support the argument for profound standardization and harmonization of septic shock trial reporting as well as data-sharing policies to test the external validity of trial populations.

#### Electronic supplementary material

The online version of this article (https://doi.org/10.1007/s00134-018-5134-8) contains supplementary material, which is available to authorized users.

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#### Author contributions

HJdG and AG designed the study. HJdG and JP performed the study screening and extracted the data. HJdG performed the statistical analyses. SL, HO, and AG provided technical, material, and administrative support. HJdG drafted the manuscript. JP, SL, JJP, HO and AG revised the manuscript for important intellectual content.

#### Compliance with ethical standards

#### **Conflicts of interest**

All authors declare that they have no conflicts of interest.

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# Hydrocortisone plus Fludrocortisone for Adults with Septic Shock

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### Abstract

**BACKGROUND** Septic shock is characterized by dysregulation of the host response to infection, with circulatory, cellular, and metabolic abnormalities. We hypothesized that therapy with hydrocortisone plus fludrocortisone or with drotrecogin alfa (activated), which can modulate the host response, would improve the clinical outcomes of patients with septic shock.

In this multicenter, double-blind, randomized trial with a 2-by-2 factorial design, we evaluated the effect of hydrocortisone-plus-fludrocortisone therapy, drotrecogin alfa (activated), the combination of the three drugs, or their respective placebos. The primary outcome was 90-day all-cause mortality. Secondary outcomes included mortality at intensive care unit (ICU) discharge and hospital discharge and at day 28 and day 180 and the number of days alive and free of vasopressors, mechanical ventilation, or organ failure. After drotrecogin alfa (activated) was withdrawn from the market, the trial continued with a two-group parallel design. The analysis compared patients who received hydrocortisone plus fludrocortisone with those who did not (placebo group).

**RESULTS** Among the 1241 patients included in the trial, the 90-day mortality was 43.0% (264 of 614 patients) in the hydrocortisone-plus-fludrocortisone group and 49.1% (308 of 627 patients) in the placebo group (P=0.03). The relative risk of death in the hydrocortisone-plus-fludrocortisone group was 0.88 (95% confidence interval, 0.78 to 0.99). Mortality was significantly lower in the hydrocortisone-plus-fludrocortisone group than in the placebo group at ICU discharge (35.4% vs. 41.0%, P=0.04), hospital discharge (39.0% vs. 45.3%, P=0.02), and day 180 (46.6% vs. 52.5%, P=0.04) but not at day 28 (33.7% and 38.9%, respectively; P=0.06). The number of vasopressor-free days to day 28 was significantly higher in the hydrocortisone-plus-fludrocortisone group than in the placebo group (17 vs. 15 days, P<0.001), as was the number of organ-failure–free days (14 vs. 12 days, P=0.003). The number of ventilator-free days was similar in the two groups (11 days in the hydrocortisone-plus-fludrocortisone group and 10 in the placebo group, P=0.07). The rate of serious adverse events did not differ significantly between the two groups, but hyperglycemia was more common in hydrocortisone-plus-fludrocortisone group.

**CONCLUSIONS** In this trial involving patients with septic shock, 90-day all-cause mortality was lower among those who received hydrocortisone plus fludrocortisone than among those who received placebo. (Funded by Programme Hospitalier de Recherche Clinique 2007 of the French Ministry of Social Affairs and Health; APROCCHSS ClinicalTrials.gov number, NCT00625209.)

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A complete list of investigators in the APROCCHSS trial is provided in the Supplementary Appendix, available at NEJM.org.

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