

# Intensive care unit occupancy: Making room for more patients\*

**M**onday, March 30, 1981 was the last day of my week on clinical call. It had been a busy weekend and all the intensive care unit (ICU) beds were full. Because three patients were ready for floor transfer and only two patients were scheduled for admission, one bed would be available later in the day. Monday's normal routine was suddenly interrupted by a TV news announcement—President Ronald Reagan had been shot at 2:30 PM. We soon heard that the President, his Press secretary, James Brady, and a Secret Service agent, Tim McCarthy, were in our emergency department.

Even with no clinical information, it never occurred to me that they would not be coming to the ICU. A discussion with the ICU nurse manager and Associate Director of Critical Care Nursing led to a list of potential ICU admissions and discharges. We also made arrangements for the care of patients who would be admitted, remain in ICU, or be discharged with special care needs. Under these circumstances, the efficiency of ICU discharge was unprecedented and our terrific staff ensured that nursing resources were practically unlimited.

Agent McCarthy was admitted to the ICU in stable condition after successful repair of a liver injury at 7:30 PM, but we heard nothing about the President. I called the recovery room and was told that the President had been there for an hour and had some disturbing blood gas results. I spent the night there providing him with respiratory care, while my colleague Bill Knaus cared for James Brady in the ICU, who had undergone a surgery for severe brain injury. The President's low Pao<sub>2</sub> was due to blood in his airways and loss of lung volume, which quickly

responded to suctioning and lung re-expansion. He was extubated at 2:30 AM, transferred to ICU at 6:15 AM, and was stable throughout his 12-hour ICU stay.

ICU clinicians know it is not unusual for a day to begin at full unit capacity and end with even higher occupancy. Beyond caring for the President, his Press secretary, and a Secret Service agent, what was unusual about the events described earlier was having practically unlimited nursing resources and an extremely efficient discharge process. In my 30 years in intensive care, I was more accustomed to begging nurses to work overtime, to perform a discharge process more difficult than “pulling teeth,” and to stretching assignments to care for more patients. Sometimes we had to “board” patients in alternative locations, and occasionally requested that ambulances be rerouted or surgery cancelled because there were no more beds and nurses.

An increased patient load was a major cause of anxiety. Could we obtain sufficient resources to care for more patients? Will staff “burnout” be aggravated by the increased workload? Will admitting more patients compromise the quality of care? In this issue of *Critical Care Medicine*, Iwashyna et al (1) provide a reassuring answer to the latter question. In 108 ICUs at 46 hospitals, consistent patient mortality was maintained, despite an unusually high census on the day of admission. These ICUs were able to cope with a census that ranged from 65% to 147% of their mean daily census for a single day or as long as 2 weeks. They also found that census fluctuations had little impact on ICU stay and the rate of transfer to other hospitals.

It is comforting to know we are likely to maintain consistent outcomes when we “play musical beds” or “stretch the staff” or stated more politely, safely scale up operations to meet operating conditions. The scaling up of care capabilities in preparation for admitting President Regan, Press secretary Brady, and Agent McCarthy represents an extreme example, and I had never thought of this often painful process as the functioning of a “high-reliability organization.” But I do

feel that ICU physicians and nurses are often negatively impacted when workload is markedly increased and sustained. The literature on ICU burnout supports this opinion (2–4).

Iwashyna et al (1) state that their study “implies, but does not prove, the viability of regionalization strategies.” The authors of the study have previously published convincing evidence about the potential value of regionalization and improved outcomes in association with concentrating patients in high-volume hospitals (5–7). I urge caution, however, in using this study to support regionalization.

There are several reasons why this study might not support regionalization: First, the study hospitals and ICUs are probably not representative of US ICUs. In addition, the study included nonteaching (34%) and small teaching hospitals (28%), institutions that are unlikely to be high-volume centers of excellence or referral hospitals. Second, the ability to successfully adjust to high occupancy for 1–14 days does not mean that caregivers could do this permanently. Third, the study did not address the ability to respond to the demands of regionalization. Additional ICU beds, professional staff, and financial resources might not be available to meet the demands of increased patient referral. Finally, the authors provide no data that identify the structural and managerial mechanisms used by these ICUs to meet these temporary increases in care demand.

All studies have limitations, but the take-home message of this one is pleasing. When faced with the need to scale up operations to meet increased patient demands, ICUs maintain consistent patient mortality and operate as high-reliability organizations. This conclusion is well supported and is good news.

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## \*See also p. 1545.

Key Words: intensive care; patient outcome assessment; mortality; burnout; workload

Dr. Zimmerman receives research support from and provides consulting services to Cerner Corporation, which provided the data used for this study.

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DOI: 10.1097/CCM.0b013e3181a0956a

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## Profiling pediatric sepsis\*

“**T**here are known knowns. These are things we know that we know. There are known unknowns. That is to say, there are things that we know we do not know. But there are also unknown unknowns. There are things we do not know we do not know.” This quotation by Donald Rumsfeld (the former Secretary of Defense) is applicable not only to the subject Mr. Rumsfeld was discussing, but also to our understanding of the pathophysiology of the sepsis syndrome from systemic inflammatory response syndrome to septic shock. While we would like to think that we know much about the biology and pathophysiology of complex diseases, such as the sepsis syndrome, and that we know the areas where we need additional information to fill in the details, it seems as although there is still much that falls into the category of “unknown unknowns.” Indeed, revealing those “unknown unknowns” is likely to generate new areas of research with potential for the development of novel therapies.

The sepsis syndrome is an exceedingly complex clinical syndrome. It exists as a continuum from minimal systemic inflammatory response syndrome manifested by fever and tachycardia to septic shock. Several complex molecular and cellular processes are involved that contribute to the clinical syndrome, including activation of the inflammatory and coagulation pathways,

disruption of vascular integrity, and apoptosis. Many other processes and pathways are also activated and/or deactivated, some are known, others we have yet to imagine.

Over the past couple of decades, advances in molecular biology and biochemistry, as well as bioinformatics, have provided important tools to better understand complex human diseases. These tools have also allowed for the discovery of processes not previously thought to play important roles in disease. One such tool, genome-wide expression microarrays, has been used to examine the expression of all genes within the genome. In this issue of *Critical Care Medicine*, Wong et al (1) use this tool to examine genes that are up-regulated and down-regulated in children with varying degrees of the sepsis syndrome, from systemic inflammatory response syndrome to septic shock. The authors' aim was to determine whether there are expression signatures that are specific to septic shock. Their data demonstrate that septic shock seems to be uniquely characterized by a decreased expression both in genes involved in zinc-related biology and in genes involved in adaptive immunity-specific signaling pathways. There were also expression signatures common across the systemic inflammatory response syndrome to septic shock spectrum of the sepsis syndrome. Such signatures were primarily characterized by increased expression of genes involved in inflammation and innate immunity.

Before this and previous studies of Wong et al using an expression profiling, the possibility that proteins involved in zinc biology were associated with septic shock was not considered, it was one of the “unknown unknowns.” However, now that these studies have pointed the field in this direction, it almost seems obvious

that this group of proteins could be involved with septic shock. Zinc is considered to be an essential trace element for the immune system, and deficiencies in zinc can result in immune system dysfunction, most notable in acrodermatitis enteropathica, a zinc malabsorption syndrome (2). Zinc deficiency is also known to result in decreased antibody formation, natural killer cell killing activity, and lower levels of phagocytosis and intracellular killing by granulocytes, monocytes, and macrophages (3). In addition, zinc blunts the inflammatory cascade and protects against lipopolysaccharide-induced shock and liver injury in animal models (4, 5).

There are many genes involved in zinc-related biology that could be crucial players in septic shock, but the data on the degree of their involvement are largely lacking. Previous reports by Wong et al have demonstrated that in pediatric patients with septic shock two isoforms of metallothionein (intracellular metal-binding proteins that bind zinc) are up-regulated in nonsurvivors compared with survivors. Consistent with an increase in the expression of metallothionein, serum zinc levels are significantly lower in non-surviving children of septic shock compared with survivors (6). In addition, in the mouse cecal ligation and puncture model of sepsis, metallothionein null mice have a significant survival advantage compared with wild-type mice (6). These studies suggest that zinc levels may impact survival in pediatric patients with septic shock.

In some instances, zinc is also an important trace element in bacterial pathogens (7). Zinc metalloproteases have been reported to be virulence factors for *Streptococcus pneumoniae* (8), *Pseudomonas aeruginosa*, *Legionella pneumophila* (9), and *Streptococcus suis* 2 (10), a strain noted for outbreaks of toxic shock syn-

\*See also p. 1558.

Key Words: sepsis; pediatrics; expression profiles; zinc

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a1a220

drome in China in 1998 and 2005. Consequently, it will be of interest to determine whether the association of zinc levels in patients correlates with survival in sepsis with various bacterial etiologies.

In summary, the current study of Wong et al characterizes expression profiles in children with varying degrees of the sepsis syndrome and indicates that septic shock is characterized by decreased expression in genes involved in zinc-related biology and in adaptive immunity. The impact of this work relates not only to identifying novel pathways and genes involved in septic shock as a way to identify novel targets for therapeutics, but it may also help identify potential markers for disease severity in children with sepsis. Ultimately, this information will further our understanding of sepsis by converting the “unknown unknowns” into “known unknowns” and eventually “known knowns.”

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## Magic bullets and surrogate biomarkers circa 2009\*

With the discovery almost 25 years ago that administration of tumor necrosis factor- $\alpha$  could produce shock and tissue injury similar to severe sepsis, the fundamental role for cytokines in experimental sepsis was established (1, 2) The dramatic improvement in survival, first seen in mice and then confirmed in primates with bacteremic shock, produced simply by inhibiting a single cytokine (3, 4), resulted in unbridled optimism that sepsis mortality in hospitalized patients could be dramatically reduced. This outlook pervaded the collective conscious for well over two decades, spawned a sundry of biological in-

hibitors, tested in a multitude of human clinical trials, all with the goal of recapitulating the survival benefit imparted by these “magic bullets” in preclinical animal models (5). Unfortunately, these efforts consumed an enormous amount of energy and financial resources, and demonstrated only minimal benefit to the treatment of human sepsis. Fortunately, the past decade has been spent in self-evaluation (some might also say self-denial), asking why such therapies have not yielded the clinic successes observed with preclinical models. Only now are we beginning to understand the complexity of human sepsis and the limits of our preclinical models (6).

In this issue of *Critical Care Medicine*, Osuchowski and co-workers (7) argue that although most antisepsis therapies, in general, and anticytokine therapies, in particular, have failed in severe sepsis clinical trials, the challenge has been to prospectively identify individuals who might benefit from such therapies (5). There is little disagreement that what we call “severe sepsis” is presently so poorly defined that our study populations are too broadly heterogenous to optimize

drug efficacy. There is a strong precedent to suggest that anti-inflammatory therapies, in general, and anticytokine therapies, in particular, are most effective in the sickest individuals at the highest risk(s) of mortality. In a large meta-analysis including both preclinical and clinical studies, Eichacker et al (8) demonstrated a linear relationship between anti-inflammatory drug efficacy and overall mortality in the placebo groups. How then do we better identify prospectively those patients with severe sepsis who may benefit from such targeted therapies? Do surrogate biomarkers exist that can prospectively identify individuals who would benefit from such therapies?

Osuchowski et al has previously shown that even individual outbred mice subjected to a reproducible cecal ligation and puncture manifest a broad range of survival responses, but mortality could be predicted by their early circulating interleukin (IL)-6 concentration (9, 10). In this report, the investigators use the same stratification system to demonstrate that only mice predicted to die based on their IL-6 concentration responded favorably to goal-directed early

### \*See also p. 1567.

Key Words: biomarkers; sepsis; corticosteroids; IL-6; mortality

Supported by a training grant in burn and trauma research (T32 GM-08431) awarded by the National Institute of General Medical Sciences, U.S.P.H.S. (to MJD).

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a09440



supraphysiologic dexamethisone. In a 21st century landscape tainted by the history of earlier failed therapeutic interdiction (11), the authors should be applauded for their demonstration that early stratification based on circulating IL-6 levels can guide corticosteroid therapy to improve survival. Equally important, the authors demonstrate that the entire septic mouse cohort and the cohort of mice predicted to survive based on their plasma IL-6 concentrations gained no survival benefit from corticosteroid therapy. Thus, the authors have emphasized the challenges and shortcomings of past trials aimed at empirical treatment of poorly defined, heterogeneous septic cohorts (5, 11).

Although the authors have provided convincing evidence that targeted therapies based on IL-6 concentrations can identify cohorts that might benefit from steroid therapies in a murine model of polymicrobial peritonitis, a number of perplexing questions remain. These studies were performed in a model of generalized peritonitis; however, different findings have been witnessed in a murine pneumonia model. Li et al (12) reported that corticosteroid therapies were broadly beneficial in a murine *Escherichia coli* model, regardless of the severity of the initial infection, and the steroids significantly lowered the plasma IL-6 concentrations across the board.

These latter findings are again different than those observed by Osuchowski and co-workers (6). In this report, the authors hypothesized that septic mice succumb to an early overwhelming systemic inflammatory response, which the authors believe may be ameliorated by strategic corticosteroid administration. Surprising was the fact that although early targeted corticosteroid therapy improved outcome, it had little impact on the circulating levels of IL-6, tumor necrosis factor- $\alpha$ , IL-1 $\beta$ , IL-2, macrophage inflammatory protein-1 $\alpha$ , macrophage inflammatory protein-2, keratinocyte-derived cytokine, and monocyte chemoattractant protein-1, all bona fide inflammatory mediators, many with prognostic value in themselves. In addition, no significant reduction was observed on circulating numbers of neutrophils, platelets, or lymphocytes as is a common effect of corticosteroids and was observed earlier (13).

Although the authors present a rational, well-considered approach to use IL-6

as a prognosticator that highly predicts early mortality in their model, and a potential 29% survival benefit from corticosteroids, the fact remains that individual animal models are rather poor surrogates for human sepsis (5, 11, 14, 15). Even though the cecal ligation and puncture model was used to replicate human peritoneal sepsis (and to many represents the "gold standard"), there are numerous intangibles such as preexisting comorbidities, age, continuous fluid resuscitation, nutritional support, guided antibiotic therapy, and operative intervention that make human sepsis more complex, and routinely difficult to replicate in mice (16). Juxtaposed with the murine vs. human sepsis conundrum stands the mortality disparity between the cecal ligation and puncture model, which was 50% across the board and 90% in the group that showed benefit, compared with an overall mortality of approximately 25% in humans (14, 17, 18). It would be interesting to know whether a more modest (LD<sub>20</sub>) model of murine sepsis (19) would yield as strong a sensitivity and specificity index for predicting 48-hour mortality, and whether the mice predicted to die would experience as great an improvement in survival from steroids. In a large clinical trial in severe sepsis with anti-tumor necrosis factor therapies using plasma IL-6 as an entry criterion, the study showed an 11% relative reduction in morbidity ( $p = 0.041$ ) with anti-tumor necrosis factor antibodies in patients with elevated IL-6 concentrations (20). In patients with an elevated IL-6 level, placebo mortality was nearly 48%. It should be noted, however, that a smaller earlier study could not confirm these results (21).

Considering the current state-of-the-art medical care, and evidence-based protocol-driven practices employed in most tertiary referral centers, the dilemma experienced by practicing clinicians is not to improve 50% but to prevent 20% mortality. This translates into a higher cost in man hours and financial resources per percentage point of survival gained (5, 18). Indeed, the ability to predict, based on IL-6 responses or some other surrogate marker, which subjects will succumb early to sepsis would be invaluable; however, given the variable etiology of sepsis and the individuals experiencing the syndrome, early identification is unfortunately dependent on multiple circumstances that typically remain far from clinician control, and difficult to

summarize with a single biological prognosticator (14).

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## Acute respiratory distress syndrome and the *Art of War*\*

*If you know the enemy and know yourself, you need not fear the result of a hundred battles.  
If you know yourself but not your enemy, for every victory gained you will also suffer a defeat.  
If you know neither the enemy nor yourself, you will succumb in every battle.*  
Sun Tzu, The Art of War, 6th century BC

**K**nowingly or unknowingly, in the fight against any disease, clinicians and researchers often apply age-old principles adapted from Sun Tzu's classic, the *Art of War*, the best-known of which reads, "know the enemy and know yourself" (1). The Acute Respiratory Distress Syndrome (ARDS) Network's 15-year battle is no exception, and the enemy—ARDS and acute lung injury (ALI)—cannot be underestimated. The defining article on ARDS by Ashbaugh et al (2) in 1967 introduced us to a highly lethal clinical entity; seven of their 12 patients died. There has since been some cause for optimism. Several observational studies documented a decrease in mortality from the 1980s to the mid-1990s, but their generalizability was limited given their single-center nature (3, 4). Enter the ARDS Network in 1994, and its multicenter randomized controlled trials (RCTs) documented mortality rates markedly lower than previous figures—in the 25% to 40% range (5–7). These observations led to hopeful assertions that

mortality from ALI/ARDS has decreased over time (8). The all-important question, nevertheless, remains: do we have objective evidence of such a decrease, and if so, what are the contributory factors? In other words, are we winning the war, and if so, do we now know the enemy, or ourselves, or both?

In this issue of *Critical Care Medicine*, Erickson et al (9) present data that provide further insight into these questions. Mortality among 2,451 patients who were enrolled in three landmark ARDS Network trials was analyzed: the ARMA trial on delivered tidal volumes (5), the Assessment of Low tidal Volume and elevated End-expiratory volume to Obviate Lung Injury trial on positive and end-expiratory pressure (6), and the Fluid and Catheter Treatment Trial on fluid strategies and the pulmonary artery catheter (7). They found a significant trend for decreasing 60-day mortality over time from 1996 to 2005. This trend persisted on multivariable analysis, which adjusted for changes in case mix, disease severity, and receipt of low tidal volume ventilation. The authors concluded that "other advancements in critical care" apart from tidal volume limitation had contributed to the improvement in survival.

The study's findings are undeniably positive news for the critical care community. Are we finally winning the war on ALI/ARDS? Some caveats should be highlighted to allow a balanced interpretation of these results. First, although RCTs can lend themselves to many complex analyses because of the rich datasets

available for each included patient, we must exercise caution when using them to make epidemiologic inferences because they generally represent only a highly selected subpopulation of all individuals affected with a disease or syndrome. Only consenting patients who met both the inclusion and exclusion criteria for these trials are included in the analysis. Indeed, these ARDS Network RCTs had at least 15 exclusion criteria each—many of which are associated with increased mortality—and excluded more than 90% of screened patients. Unfortunately, although these excluded patients had a higher mortality (10), information on their mortality trends is unavailable.

Second, the decline in mortality was predominantly restricted to the first 4 years (1996–1999) of the study period: 60-day crude mortality was 35%, 32%, 27%, 27%, and 26%, respectively, in the five 2-year time periods between 1996 and 2005. Changes in enrolment practices could have contributed to the dramatic 8% decrease in mortality between 1996 and 1999, although the temporal increase in Acute Physiology and Chronic Health Evaluation III scores argues against this. The bulk of the patients included during 1996–1999, when the large mortality reductions took place, were from the ARMA study, in which half of the population received higher tidal volumes of 12 mL/kg predicted body weight. However, in this specific case, the effects of high tidal volumes should have been accounted for in the multivariable analysis; it is known which pa-

### \*See also p. 1574.

Key Words: acute respiratory distress syndrome; acute lung injury; randomized controlled trials; mortality  
Dr. Ferguson is supported by a New Investigator Award from the Canadian Institutes of Health Research (Ottawa, Canada).

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a09310

tients were randomized to this group, and protocol adherence to lower tidal volume was good across all other groups. The authors hypothesized that “other advancements in critical care” led to the decrease in mortality, but the available data do not tell us just what these “other advancements” are. Tidal volume limitation remains the only intervention that has been convincingly shown to reduce mortality in ALI/ARDS (5), and even this is more a limitation of iatrogenic injury than a therapeutic intervention *per se*. Meanwhile, controversy surrounds other therapies for sepsis (a major cause of ALI/ARDS), including glucose control, human recombinant activated protein C, and corticosteroids (11).

What messages should we then take home from the findings of Erickson et al? The authors were rightly cautious to emphasize that their findings apply specifically to patients “treated at ARDS Network Centers.” The improving trends suggest that patients with ALI/ARDS may benefit by being enrolled in an ARDS Network RCT. We can only speculate why this may be so. First, clinicians and researchers may have learnt more about the enemy over time, i.e., small tidal volumes can subdue the enemy. Second, clinicians and researchers may have learnt more about themselves over time, i.e., the ARDS Network centers may have steadily fine tuned their clinical and research infrastructure, thereby resulting in earlier case detection, better use of management bundles, such as early resuscitation, appropriate antibiotics, and sedation protocols (12), and/or they may have improved care simply through the experience of expert clinicians in centers with high volumes of patients with ALI.

The data of Erickson et al may be interpreted in light of two recent systematic reviews (including one from our group), which also dealt with mortality trends in ALI/ARDS (13, 14). While the first smaller review found a similar decline in mortality (13), when we evaluated 53 observational studies and 36 RCTs published between 1984 and 2006 (including the ARMA, Assessment of Low

tidal Volume and elevated End-expiratory volume to Obviate Lung Injury, and Fluid and Catheter Treatment trials), we found a static mortality between 1994 and 2006, plus a significantly lower mortality in RCTs than observational studies.

Erickson et al must be applauded for shedding more light on this topic. They have clearly shown that outcomes among patients randomized in ARDS Network trials have improved over time. Yet, given the caveats we have highlighted, and in answer to our original question: no, we cannot say that we are winning the war on ALI/ARDS. We have come a long way in the last 40 years, however, we still do not know enough of this enemy that remains a mixed bag of heterogeneous disorders; and we do not yet know enough of our own armamentarium (8). The battle against this important clinical entity goes on.

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# Prolonged glucocorticoid treatment in acute respiratory distress syndrome: Evidence supporting effectiveness and safety\*

In this issue of *Critical Care Medicine*, Tang et al (1) provide a systematic review and meta-analysis of nine controlled studies ( $n = 648$ ) evaluating the effectiveness of prolonged glucocorticoid treatment (PGCT) in patients with acute lung injury and acute respiratory distress syndrome (ARDS) (2–10). These studies (Table 1) consistently report a significant improvement in  $\text{PaO}_2/\text{FiO}_2$  (2–10) and a significant reduction in markers of systemic inflammation (2–10), multiple organ dysfunction score (2, 5, 6, 8–10), duration of mechanical ventilation (2, 4–6, 10), and intensive care unit length of stay (2, 3, 5, 6, 10) (all with  $p$  values  $<0.05$ ) without an increased rate of complications. The reduction in duration of mechanical ventilation is two- to three-fold greater than the reduction reported with low tidal volume ventilation or the conservative strategy of fluid management (11, 12). The more rapid resolution of lung injury and multiple organ dysfunction scores observed in these trials could positively affect long-term physical recovery (13) and survival (5). The relevance of these findings to public health and healthcare economics urges investment in clinical investigation of this inexpensive and highly effective anti-inflammatory therapy.

Because of differences in study design and patient characteristics and the limited size of the studies (1–4), the cumulative mortality summary should be interpreted with some caution. Nevertheless, all four studies (2–5) ( $n = 334$ ) investigating treatment initiated within 3 days of meeting acute lung injury and ARDS criteria showed a significant reduction in mortality, with an overall 24% absolute reduction in

mortality (risk ratio, 0.69; 95% confidence interval, 0.56–0.84). Two of the five studies (6, 9) investigating treatment initiated after 5–7 days of meeting ARDS criteria showed a significant reduction in mortality, whereas three studies (7, 8, 10) found no difference in mortality, with an overall 15% absolute reduction in mortality (risk ratio, 0.75; 95% confidence interval, 0.55–1.02) that increased to 19% for the larger subgroup of patients ( $n = 233$ ) randomized before day 14 of ARDS (risk ratio, 0.65; 95% confidence interval, 0.45–0.94) (6, 8–10). In the three studies without a mortality benefit (7, 8, 10), treatment was associated with significant early physiologic improvement; however, rapid dosage reduction (8) or premature removal after extubation (as acknowledged by the authors) (10) might have affected final outcome. In the ARDS Network trial, the treated group had—before removal of treatment—a noteworthy 9.5 days' reduction in duration of mechanical ventilation ( $p = 0.006$ ) (10). The ARDS Network trial reported a lower mortality for patients ( $n = 132$ ) randomized before day 14 (27% vs. 36%;  $p = 0.14$ ) and higher mortality for patients ( $n = 48$ ) randomized after day 14 of ARDS (8% vs. 35%;  $p = 0.01$ ) (10). The latter subgroup, however, had large imbalances in baseline characteristics and an uncharacteristically low mortality in the control group, and the mortality difference lost significance ( $p = 0.57$ ) when adjusting for these imbalances (14, 15).

Treatment decisions involve a tradeoff between benefits on the one hand and risks, burdens, and, potentially, costs on the other (16). As an aggregate ( $n = 648$ ), absolute and relative reduction in mortality is substantial for all patients (18% and 35%) and even greater when treatment is initiated before day 14 of ARDS (21% and 38%). While awaiting a larger confirmatory trial in early acute lung injury and ARDS, this meta-analysis provides evidence of a sizable reduction in duration of mechanical ventilation and intensive care unit stay and a considerable survival benefit with the potential

saving of one life for every four treated patients (1). In the United States alone, this could translate to tens of thousands of lives saved per year and several billion dollars in reduced healthcare expenditures (17). Furthermore, the low cost of off-patent methylprednisolone—in the United States approximately \$240 for 28 days of intravenous therapy (5)—makes this treatment globally and equitably available.

In their systematic review, Tang et al (1) report that PGCT at low-to-moderate doses was not associated with an increased rate of major complications, including infections and neuromyopathy. This counterintuitive finding deserves further elucidation and provides an opportunity to debunk common fallacies about glucocorticoid treatment-associated complications. Most misconceptions originate from the findings of sepsis and ARDS trials conducted in the 1980s that investigated a massive daily dose of glucocorticoids (methylprednisolone, up to 120 mg/kg/day) over a short time interval (24–48 hours). The experimental model supporting this treatment protocol relied on the intravenous administration of a lethal bolus of lipopolysaccharide that could be offset only by administering an equally massive dose of glucocorticoids before or within a brief experimentally generated inflammatory window (18–20). This experimental model did not replicate human sepsis or acute lung injury and was discredited in the early 1990s (18). Since then, longitudinal measurements of inflammatory cytokines in ARDS patients have clearly shown that significant systemic and pulmonary inflammation persists 28 days into the disease process (21–24) and that PGCT (14–21 days) followed by slow tapering is essential to achieve and sustain biological resolution of the disease process (21, 25).

In recent years, substantial evidence has accumulated showing that systemic inflammation, the central pathogenetic process in ARDS (21, 22, 24), is also implicated in the type of morbidity—

## \*See also p. 1594.

Key Words: acute respiratory distress syndrome; complications; duration of mechanical ventilation; glucocorticoid treatment; infection; mortality

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e31819d2b43

**Table 1.** Prolonged glucocorticoid treatment in ALI-ARDS: Overall mortality, improvement in markers of systemic inflammation, gas exchange, duration of mechanical ventilation, and ICU stay

Study	Hospital Mortality <sup>a</sup>	Reduction in Inflammation	Improvement in PaO <sub>2</sub> :FiO <sub>2</sub>	Reduction in MV Duration	Reduction in ICU Stay
Early ALI-ARDS (n = 334)	38% vs. 62%	3 of 3	4 of 4	4 of 4	3 of 3
Confalonieri (2)	0.0% vs. 30%	Yes	Yes	Yes	Yes
Lee (3)	8% vs. 88%	NA	Yes	Yes	Yes
Annane (4)	64% vs. 73%	Yes	Yes	Yes	NA
Meduri (5)	24% vs. 43%	Yes	Yes	Yes	Yes
Late ARDS (n = 314)	28% vs. 43%	5 of 5	5 of 5	2 of 3	2 of 3
Meduri (6) <sup>b</sup>	12% vs. 62%	Yes	Yes	Yes	Yes
Keel (7)	38% vs. 67%	Yes	Yes	NA	NA
Varpula (8)	19% vs. 20% (30 d)	Yes	Yes	No	No
Huh (9)	43% vs. 74%	Yes	Yes	NA	NA
Steinberg (10) <sup>b</sup>	29% vs. 29% (60 d)	Yes	Yes	Yes	Yes
Early and Late ARDS	34% vs. 52%	8 of 8	9 of 9	6 of 7	5 of 6

Early ALI-ARDS, treatment initiated within 3 days of meeting acute lung injury-acute respiratory distress syndrome criteria; Late ARDS, treatment initiated after 5–7 days of meeting ARDS criteria; NA, not available or not applicable; d, days; MV, mechanical ventilation; ICU, intensive care unit.

<sup>a</sup>Mortality is reported as hospital mortality unless specified otherwise in parenthesis; <sup>b</sup>Two trials reported mortality in patients randomized before day 14 of ARDS: Meduri et al (6) (13% vs. 57%), and Steinberg et al (10) (27% vs. 36%). Comparisons are reported as glucocorticoid-treated vs. control.

hyperglycemia (26), neuromuscular weakness (27), increased risk for nosocomial infections (28), and delirium (27)—that is otherwise attributable to glucocorticoid treatment. Two sets of consequences follow. First, in uncontrolled studies, when the use of glucocorticoid treatment is limited to the “rescue” of the sickest patients, it is difficult to separate disease from treatment-related complications. Second, treatment-induced downregulation of systemic inflammation could theoretically prevent, or partly offset, development and progression of these complications. The findings reported by Tang et al (1) add credit to a new line of reasoning that views control of systemic inflammation as indispensable to decreased short- and long-term morbidity in ARDS and sepsis. We review evidence accumulated in the last decade that supports this paradigm shift in our traditional thinking. Because of space limitations, we will address only the topic of infection and neuromuscular weakness.

None of the reviewed trials reported an increased rate of nosocomial infection (1, 29), whereas two reported a significant reduction (5, 10). In the ARDS network trial (10), the infection rate for treated and control groups was 31% vs. 47% (relative risk 0.59, 95% confidence interval 0.40–0.88), respectively. A prospective ARDS study investigating the longitudinal relationship between inflammatory cytokine levels (plasma and bronchoalveolar lavage) and infections (28) showed that nosocomial infections are an epiphenomenon of dysregulated systemic inflammation (28, 30), a finding supported by recent experimental evidence

(31). It is now appreciated that bacteria have receptors for cytokines tumor necrosis factor- $\alpha$  and interleukin (IL)-1 $\beta$  and that exposure of bacteria to inflammatory cytokines enhances their growth and virulence (reviewed in Ref. 30). Although a moderate degree of local inflammation is required to control infection, high levels of inflammatory cytokines favor bacterial proliferation and virulence following a U-shaped response. When fresh isolates of *Staphylococcus aureus*, *Pseudomonas aeruginosa*, and *Acinetobacter* spp. obtained from patients with ARDS were exposed *in vitro* to a lower concentration (10–250 pg) of tumor necrosis factor- $\alpha$ , IL-1 $\beta$ , or IL-6—similar to the plasma values detected in ARDS survivors—extracellular and intracellular bacterial growth was not promoted, and human monocytic cells were efficient in killing the ingested bacteria (32, 33). On the contrary, when bacteria were exposed to higher concentrations of proinflammatory cytokines—similar to the plasma values detected in ARDS non-survivors—intracellular and extracellular bacterial growth was enhanced in a dose-dependent manner (32, 33). In separate parallel experiments, impairment in intracellular bacterial killing by activated monocytes correlated with the increased monocyte expression of proinflammatory cytokines, whereas restoration of monocyte killing function upon exposure to methylprednisolone coincided with the downregulation of the expression of tumor necrosis factor- $\alpha$  and IL-1 $\beta$  (34). In experimental sepsis, the magnitude of induced systemic inflammation affects

macrophage-associated host antibacterial innate immunity and susceptibility to infection (31).

Glucocorticoid treatment has been proven beneficial and safe for a wide variety of infections (35). In experimental pneumonia, animals randomized to glucocorticoid treatment had a significant reduction in bacterial burden (bronchoalveolar lavage and lung tissue) and histologic severity of pneumonia (36) and improved survival (37). Two randomized trials in patients with septic shock (most caused by pneumonia) reported that hydrocortisone infusion—while decreasing circulating IL-6 and IL-8 levels and neutrophil spontaneous release of hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>)—was associated with *ex vivo* preserved or increased (in comparison to placebo) monocyte and neutrophil phagocytosis and preserved respiratory burst (38, 39). The cumulative evidence indicates that in ARDS and severe sepsis, glucocorticoid-induced downregulation of life-threatening systemic inflammation improves innate immunity (38, 39) and provides an environment less favorable to intracellular and extracellular bacterial growth (34, 36).

A number of reports, including muscle histopathologic studies (40), support an association between systemic inflammation and critical illness neuromuscular abnormality (CINMA) (27). In this regard, downregulation of systemic inflammation in ARDS could potentially prevent development and progression of CINMA. A recent systematic review of the literature found no clear association between glucocorticoid treatment and electro-



physiologically proven CINMA in patients on mechanical ventilation (41). Because CINMA is recognized as an independent predictor of prolonged weaning (42) and ARDS trials consistently report a sizable and significant reduction in duration of mechanical ventilation (2,4–6,10), clinically relevant CINMA caused by glucocorticoid or glucocorticoid-induced hyperglycemia is unlikely.

The ARDS network study (10) reported nine serious, unspecified (not defined in the text of the article) events associated with myopathy or neuropathy among treated patients ( $p = 0.001$ ). Specific information about these patients, however, was not provided. Because greater than 40% of patients in the ARDS network trial were exposed to neuromuscular blocking agents (NMBA), it is difficult to factor how much the combination NMBA–glucocorticoid affected this finding. In ventilated patients with severe asthma on PGCT, the use of glucocorticoids in combination with an NMBA was associated with a much higher incidence of muscle weakness than was the use of glucocorticoids alone (20 of 69 vs. 0 of 38,  $p < 0.001$ ) (43). In the single study that identified by multivariate analysis a positive relationship between glucocorticoid treatment and CINMA, most patients were exposed to NMBA for 3 days (44). For this reason, the use of NMBA is strongly discouraged in ARDS patients receiving glucocorticoid treatment (15).

It can be argued that, lacking a large confirmatory trial proving a definitive mortality benefit, caution is warranted in recommending PGCT in ARDS. Weighting in favor of this approach are concerns related to glucocorticoid-induced complications, a line of reasoning that is weakened by the findings of this meta-analysis. As reviewed by Tang et al, cumulative findings from multiple controlled ARDS studies have consistently demonstrated that PGCT is associated with a sizable and significant improvement in meaningful patient-centered outcome variables and a distinct survival benefit. Subgroup analysis should be interpreted with some caution. Although the analysis by Tang et al did not find the dosage strategy to affect outcome, we believe that corticosteroids should be initiated early at a dose not exceeding 1 mg/kg/day of methylprednisolone in patients meeting criteria for severe ARDS ( $\text{PaO}_2/\text{FiO}_2 \geq 200$  on positive end-expiratory pressure of 10 cm  $\text{H}_2\text{O}$ ) and within 72 hours in patients without severe ARDS

but failing to improve by day 3. In patients failing to improve lung injury score after days 5–7, present data are limited to the study investigating a dose of 2 mg/kg/day of methylprednisolone. In all cases, duration of treatment should be 14–21 days before final tapering. Most important, PGCT has a strong benefit/risk ratio in ARDS when it is applied in conjunction with measures demonstrated to reduce potential morbidity associated with glucocorticoids (5, 6). These measures include i) intensive infection surveillance, ii) avoidance of paralytic agents, and iii) avoidance of rebound inflammation with premature discontinuation of treatment that may lead to physiologic deterioration and reintubation (5).

## ACKNOWLEDGMENT

We thank Dr. David Armbruster for critical review of the manuscript.

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## How can severity scoring methods maintain clinical and temporal relevance with advances in critical care practice? Here's one way\*

**O**ne dilemma facing users of risk-adjusted severity scoring as a surrogate for quality in the intensive care unit (ICU) is the difficulty in ensuring that predictive modeling accurately reflects advancements

in clinical care, admission criteria, or other practice changes. Earlier in this decade, Glance et al (1), using the standardized mortality ratio (ratio of mortality predicted by a severity score to actual mortality) as a measure of ICU quality, hypothesized that existing individual scoring systems should consistently and reproducibly demonstrate high-quality or low-quality ICU outliers. While establishing that there was only a fair to moderate agreement among Acute Physiology and Chronic Health Evaluation II (2), Simplified Acute Physiology Score II (3), and Mortality Probability Admission Model (MPM<sub>0</sub>-II) (4) in determining

ICU quality outliers using the Project IMPACT (5) database, they also demonstrated an interesting, unintended finding: that most of the ICUs were, with apologies to Garrison Keillor and the Lake Wobegon effect (6), “above average” (i.e., of high quality).

Because the severity scoring methods used in the aforementioned study (1) were published 10–15 years prior, most clinicians would argue that skewing of the bell-shaped curve toward inflation of ICU quality is indicative of the prediction models losing their relevance as care practices change over time. Unfortunately, the only way to refine an exist-

### \*See also p. 1619.

Key Words: severity scoring; outcomes measurement; standardized mortality ratio; quality of care; critical illness

The author has not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a09375

ing predictive model is by subjecting it to a new, contemporary dataset of ICUs or patients—a process that is typically expensive and time consuming. Recognizing that periodic updating of prediction models is necessary, Acute Physiology and Chronic Health Evaluation (7), Simplified Acute Physiology Score (8), and MPM (9) have all recently been recalibrated to align with evolving clinical practices.

However, the time between published updates of these scoring systems, typically a decade or more, may be undesirably long for clinicians to accept that a particular model remains temporally relevant. In this issue of *Critical Care Medicine*, Higgins et al (10) performed a prospective revalidation of MPM<sub>0</sub>-III using a new patient population acquired from Project IMPACT 15 months after the original population, with the goal of demonstrating ongoing robustness of the model. The researchers analyzed more than 55,000 admissions from 77 hospitals and 103 ICUs; nearly a quarter of the included ICUs were not part of the initial MPM<sub>0</sub>-III development or internal validation dataset. They assessed discrimination using the receiver operating characteristic, and calibration using graphic, Hosmer-Lemeshow goodness of fit, and Brier score statistical methodologies (10). Despite limitations inherent in using a closed database like Project IMPACT (such as its retrospective nature, underrepresentation of ICU types, and exclusion of patient subsets), the authors conclude that the MPM<sub>0</sub>-III calibrates and discriminates well using a new, more contemporary patient population (10). In addition, their results remain stable even though the more recent study group had statistically significant reductions in the number of low-risk patients and increases in those receiving active treatment and mechanical ventilation when compared with the original validating population (10). These significant differences occurring only 15 months apart suggest that even relatively brief time lapses may lead to substantial clinical practice changes.

The critical care medicine community has been at the forefront of developing risk-adjusted clinical scoring systems, such as Acute Physiology and Chronic Health Evaluation, Simplified Acute

Physiology Score, and MPM, that could arguably be considered surrogates for helping to define quality of care. Yet, clinical mortality-based measures of outcome (e.g., the standardized mortality ratio) have not become ubiquitous in ICUs, despite being available in one form or another for a quarter century. Reasons for this lack of embracement include the distaste for the high costs associated with what has evolved into a proprietary, for-profit enterprise; the additionally expensive but required labor-intensive data collection process; concerns that a scoring system would not accurately reflect a particular patient population; or, perhaps the lack of transparency in knowing whether a particular severity score may underpredict or overpredict outcomes during those lengthy intervals between published updates. The study by Higgins et al (10) addresses this latter concern by demonstrating that their scoring system remained robust with a different, more contemporary patient population. It may be, then, that one way to alleviate concerns about the relevance of a particular scoring system would be insistence by the critical care community that studies like the one by Higgins et al (10), transparent and subjected to peer review, are published more frequently and continually in the medical literature. Such studies may be necessary to convert skeptical clinicians that outcome predictions and surrogates for ICU quality (such as the standardized mortality ratio) retain reliability sufficient to accept into their practices.

Obviously, there are economic issues at stake regarding future completion of labor-intensive clinical revalidation studies. A major limitation of the current study by Higgins et al (10) is that all of the authors have some financial ties to, and most are direct employees of, the company that owns the rights to MPM<sub>0</sub>-III. That the study was completed at all is illustrative of the integration of science and industry necessary for maintaining validity of their severity scoring model. To ensure frequent, future revalidation studies, it may be time to consider assembling interested investigators, perhaps funded by a novel governmental, third-party payer and for-profit enterprise consortium, with the (admittedly lofty) goal of producing frequent, independent, high-quality, transparent, peer-reviewed

literature demonstrating reliability of (or recalibrating as necessary) the available risk-adjusted scoring systems. The study by Higgins et al (10) might be seen as an initial step by the critical care community in accepting that risk-adjusted severity scores and outcome measurements (like the standardized mortality ratio) have resiliently withstood the test of time and focusing investigative efforts on how to scientifically and transparently ensure their ongoing clinical and temporal relevance.

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# Nosocomial bacteremia a continuing challenge—A challenge we can meet\*

Hospitals, and more explicitly intensive care units (ICUs), are healthcare delivery sites where activities aimed at improving patient outcomes are coupled with risks that may lead to increased patient morbidity and mortality. This harm is a byproduct of the intrinsic characteristics of the critically ill patients who we care for; however, our care delivery systems also have a prominent role.

The topic of nosocomial infections has been extensively reviewed (1, 2), with urinary tract, lower respiratory tract, surgical site, and primary bloodstream infections being the most frequent sources. The epidemiology and outcomes of nosocomial infections have been described, and the results indicate an increasing toll on our patients.

In this issue of *Critical Care Medicine*, Blot et al (3) focus on nosocomial bacteremia, one of the most frequent and lethal nosocomial infections in ICU patients (1). They provide us data on the trends and outcomes over a 15-year timeframe of nosocomial bacteremia in critically ill patients from their clinical database. Their study, a single institution's retrospective cohort study, examines 984 ICU patients (cardiosurgical ICU 55.1%, medical ICU 37.8%, and burn unit 7.1%) with 1228 episodes of microbiologically confirmed nosocomial bacteremia. The source of nosocomial bacteremia was primary (31.8%), catheter related (18.5%), lung (14.5%), intra-abdominal (11.9%), wound (7.9%), urinary tract (7.9%), and either another site or multiple sites (7.5%). A total of 48.7% episodes were caused by Gram-positive organisms, 41.3% by Gram-negative organisms (70% *Enterobacteriaceae* and 30% nonfer-

menting Gram-negative bacteria), 2.2% by anaerobic bacteria, 7.8% by *Candida* species, and 15.3% by polymicrobial bloodstream infections. The article provides some new insights and some thoughts on what needs to be done to improve performance.

The authors categorized patients into middle aged (patients 45–64 years), old (65–74 years), and very old ( $\geq 75$  years). They assessed a number of parameters, with the most important being the age distribution of patients in the ICU, the age distribution of those with nosocomial bacteremia, and the prevalence of nosocomial bacteremia and in-hospital mortality in ICU patients with nosocomial bacteremia as it relates to age.

They observed that from study onset to the end, despite a 10% decrease in overall ICU admissions, the number of patients in the very old group increased by 33%. The prevalence of nosocomial bacteremia increased significantly over the 1992–2006 timeframe from 21.3 of 1000 admissions in 1992 to 33.6 of 1000 admissions in 2006 in the old age group ( $p < 0.001$ ) and from 9.7 of 1000 admissions to 21.3 of 1000 admissions in the very old group ( $p < 0.002$ ).

In-hospital mortality increased significantly with age: 42.9%, 49.1%, and 56.0% for middle aged, old, and very old patients, respectively. Observed mortality (56.0%) significantly exceeded expected mortality (based on Acute Physiology and Chronic Health Evaluation II scores) in the very old group and fell outside the 95% confidence intervals of the expected mortality (35.6% to 45.5%).

The proportion of elderly patients in our population continues to grow and the proportion of elderly patients admitted to intensive care is increasing and will continue to increase (4, 5). Although the authors' results bear out the changing demographic of our ICUs (6, 7), their article reinforces the mandate that age is an important factor to be considered in our care processes.

Blot et al reported that the number of very old increased by 33% during the study timeframe, whereas the prevalence of nosocomial bacteremia with its associated elevated mortality in this population increased disproportionately to more than 240%. This trend is indeed sobering. One unfortunate issue that the authors highlight is that nosocomial bacteremia in the critically ill may be following the path of sepsis in the general population; with a significant overrepresentation of the elderly, this prevalence may be growing at a more rapid rate and with poorer outcomes (8).

The study has a number of limitations, which include the following: nutritional and functional status, comorbidities central components for outcomes for the elderly were not assessed. The authors did not explore whether limitations of care initiated by physicians or patients contributed to mortality. Transfusion history, which is associated with an increased rate of nosocomial bacteremia, was not part of the database (9). The types of ICUs in the authors' institution may not reflect the makeup of many of our facilities. The classic age categories used in a number of studies on nosocomial bacteremia differ from those used in the current study.

The elderly are an ever-present and ever-increasing populace in our ICUs, and they have age-associated factors added to comorbidities that lead to increased rates of nosocomial bacteremias. The roles played by impaired host defenses, altered immune function, anatomical or physiologic derangements, or comorbidities are unclear and warrant further investigation.

Despite our lack of understanding of the cause of nosocomial bacteremias, a number of risk factors for nosocomial bacteremias, such as central venous catheterization, urinary catheterization, ICU-acquired pneumonia, neurologic impairment, increased length of ICU stay, mechanical ventilation, diagnosis of trauma, certain chronic disease processes, and high intensity of care, have been identified (1).

Many of these factors can be mitigated and that is where our focus should be

\*See also p. 1634.

Key Words: nosocomial; bacteremia; elderly; mortality; catheter; prevention

The author has not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a0939e

directed. An excellent example is the prevention of catheter-related bloodstream infections by incorporating into ICU practice five evidence-based procedures that have the greatest effect on the rate of catheter-related bloodstream infections and the lowest barriers to implementation (10). Additional measures to be considered include the Surgical Infection Prevention Process Measures, prevention of ventilator-associated pneumonia with a ventilator-associated bundle, and the use of policies and practices to prevent catheter-related urinary tract infections (11).

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## A marker for posttraumatic-sepsis: Searching for the Holy Grail around intensive care units\*

In the Western world, trauma continues to be one of the major causes of mortality in people younger than 50 years. Patients with trauma die either as a direct consequence of their injuries or from the development of secondary complications such as sepsis and sepsis-related multiple organ failure (the leading cause of death in intensive care units). Risk assessments and prompt diagnosis have become major tools to prevent sepsis in posttraumatic patients. An early diagnosis is extremely important due to a lack of proven treatments for improving the final outcome in septic patients. The host injury level is evaluated by scoring systems that combine clinical signs and measurements of proinflammatory parameters (1). The most relevant sepsis markers currently used are C-reactive protein (CRP), procalcitonin (PCT), proinflammatory cytokine production (such as IL-6, IL-8, and IL-

18), and polymorphonuclear leukocyte (PMNL) count (2). Unfortunately, these parameters are also increased in a variety of other conditions, creating uncertainty at the time of diagnosis and prognosis of sepsis. Thus, searching for new biochemical or immunologic marker(s) has been a major goal for researchers during the past decades. In this issue of *Critical Care Medicine*, Keel et al (3) present data supporting the role of plasma PSP/reg (pancreatic stone protein or reg) levels as a prognosis maker for posttraumatic complications such as sepsis. PSP/reg is a secreted tryptic fragment found in pancreatic juice as well as in pancreatic stones (4, 5). PSP/reg is also produced in the small intestine (6) and stomach (7) and has been involved in cellular regeneration and healing (8, 9). Plasma levels of PSP/reg dramatically increased in patients with acute and chronic pancreatitis (10). In their study, Keel et al used a small group of severely injured young patients (average age of 38 years) who were divided into three groups, patients without infection (n = 18), patients with local infection (n = 32), and septic patients (n = 33), and these groups were compared with a group of healthy volunteers. Their data showed that on the day

of admission, all the four groups had similar levels of plasma PSP/reg. However, within a couple of days, a significant increase in PSP/reg levels was observed in patients with local infection and in those with sepsis. These two groups had different kinetics of increase, which were unrelated to pancreatic damage (measured as lipase and amylase activity). As expected, sepsis markers, CRP, leukocyte count, IL-6, and PCT showed an early peak (around day 7) in septic patients, and their values were significantly different from those observed in patients with local or no infection. In an attempt to elucidate whether this increase in PSP/reg level has a biological function, the authors decided to evaluate the effect of this protein on PMNL function. They found that incubation of control PMNL with recombinant PSP/reg (at levels similar to those found in plasma) resulted in cell surface reduction of CD62L (by shedding or internalization?) and an increase of CD11b. This profile strongly suggests that PMNL exposed to PSP/reg develop a firm adherence phenotype, which may facilitate their infiltration and, therefore, their activation. These authors ruled out potential endotoxin contamination by boiling PSP/reg samples before incubation.

\*See also p. 1642.

Key Words: PSP/reg; neutrophils; inflammation; multiple organ failure; immune response

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DOI: 10.1097/CCM.0b013e3181a1a0b5

tion with PMNL. Additionally, direct binding of PSP/reg to PMNL derived from healthy volunteers was observed. Incubation of PMNL isolated from septic patients with PSP/reg did not show an effect over CD62L or CD11b cell surface levels. These data may suggest tolerance-like phenomena, although further investigations are required. One of the questions that remained unanswered in this work is what is the binding capacity of PSP/reg by sepsis-derived PMNL? In addition, it will be important to know if this unresponsiveness to PSP/reg or altered binding capacity are parameters that may be used as prognosis indexes. To assess the real relevance of PSP/reg as a sepsis marker will require further investigations. Furthermore, PSP/reg will be competing with new sepsis marker candidates such as peripheral endothelial progenitor cells (11), plasma Treg cells, CD25 levels (12), and B-type natriuretic peptide (13) to be crowned as the definitive sepsis marker.

Thus, finding the “perfect” sepsis marker has been one of the most elusive dreams in modern medicine. The list of potential sepsis markers increases day by day, and we still do not have a parameter or a group of them that can accurately and rapidly diagnose sepsis. Most of the current markers (clinical signs and laboratory measurements) are the product of the proinflammatory stage and therefore are nonspecific. Thus, if we can make a wish for the ideal sepsis marker, what would we ask for? Probably an important characteristic would be a parameter that is altered in all types of sepsis,

independently of the agent causing the infection. This characteristic would eliminate “subpopulation of septic patients” and all the nightmares associated with conflicting data at the moment of evaluating a potential sepsis marker. Thus, it would be desirable to have a substance that reports early changes and can be detected in an easy and rapid way. Prognosis potential is also a characteristic that should be added to this wish list. PSP/reg seems to have a certain potential as a predictor of sepsis, although only time will tell if this protein fulfills the minimum requirements to be called a true sepsis marker.

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## Predictive models: The angel is in the details\*

**M**odels for measuring severity of illness and predicting hospital mortality for patients in intensive care units (ICUs) are now in their third and fourth

generation (1–3), and newer models have appeared recently (4, 5). This has come about as a result of the desire to assess ICU performance by comparing observed and predicted mortality (6, 7) and, at least in part, to the ability to capture data electronically. Large data sets containing numerous measurements on all patients have enabled the development of sophisticated predictive models. Simplicity, however, is not a hallmark of these systems. The leanest critical care hospital mortality model, Mortality Probability Model (MPM<sub>0</sub>-III),

still requires the collection of 17 data elements (3).

Statistical modeling in other acute care settings has lagged behind that in critical care. Recent attempts have been made to introduce predictive models outside of the ICU, most particularly in the area of rapid response teams/medical emergency teams (8). One example is the Modified Early Warning System (9), which was developed on 206 patients in a postoperative ward. This scoring system assigns “weights” to six physiologic measurements. The weights are summed and

\*See also p. 1649.

Key Words: intensive care; patient outcome assessment; Sequential Organ Failure Assessment Score; emergency department; mortality

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DOI: 10.1097/CCM.0b013e3181a093cb



a cut-point  $>4$  is used as an early warning signal. What is common among decision algorithms in rapid response teams/medical emergency teams is that they use triggers ("antecedents") and/or uncomplicated scoring algorithms to indicate possible patient deterioration. Here, simplicity is desirable, until such time as electronic data capture mechanisms permit the assimilation and multivariate analysis of high-dimensional data.

In this issue of *Critical Care Medicine*, Jones et al (10) examine how the Sequential Organ Failure Assessment (SOFA) score (11) measured while the patient is in the emergency department (ED), can predict hospital mortality for patients subsequently admitted to an ICU. The authors also gathered data for the SOFA score collected 72 hours postadmission as well as the difference between this value and the value recorded in the ED. These additional measurements may be useful variables for stratification in *post hoc* analyses, but as they are based on information collected after admission to the ICU, they cannot be used as a predictor for patients in the ED.

In addition to the SOFA score, the authors collected information on other physiologic measures while patients were in the ED: vital signs, oxygen saturation, Glasgow Coma Score, white blood cell count, and lactate concentration. These were all recorded prospectively on standardized forms. The patient population is narrowly focused in terms of case mix: 248 patients with severe sepsis who had resuscitation procedures initiated in the ED. This strategy is a wise one, given the difficulty of developing a model using what would be an otherwise heterogeneous population. Limiting the study to a single institution, however, means that the results reported by Jones et al must be considered exploratory.

The primary statistic chosen for determining the ability to predict hospital mortality was the area under the receiver operating characteristics curve (AU-ROC)

(12). This is a measure of "discrimination," i.e., the ability to distinguish between patients who die vs. those who survive. The AU-ROC ranges from 0.50, which indicates that the prediction is no better than flipping a coin, to 1.00, which is a perfect predictor. All of the ICU predictive models cited above have AU-ROC values  $>0.80$ . Values lower than that are considered mediocre. Unfortunately, the AU-ROC to predict mortality using the SOFA score taken in the ED was only 0.75.

The authors compare this value with the AU-ROC generated by other variables they had collected, and found that none had higher AU-ROC values than the SOFA score. Given the additional physiologic variables that Jones et al collected, they might have considered a more sophisticated approach. They certainly could have constructed a pseudo-Modified Early Warning System instrument, which most likely would have increased their AU-ROC beyond 0.80.

The authors should be commended for assessing the value of the relatively simple SOFA score in the ED as a predictor of subsequent in-hospital mortality. But in their attempt to maintain simplicity, they gave away the opportunity to look at a metric that had high discrimination with little additional data capture burden. I agree with the authors that using ICU predictive models in the ED is at present not feasible. But, although simplicity can be alluring, it should not trump a comprehensive inclusion of enough variables to generate a precise yet timely prediction. When it comes to predicting outcomes, the angel is in the details.

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# Old friends: Pneumonia and interleukin-6\*

**P**neumonia has been termed “The old man’s friend” because it may ease suffering at the end of life (1). Previously, investigators have reported that pneumonia has a worse prognosis in elderly men compared with women (2). In this issue of *Critical Care Medicine*, Reade et al (3) provides insights into why older men with pneumonia have a worse prognosis. They specifically measured circulating biomarkers and demonstrated that at admission to the emergency department, men had higher levels of tumor necrosis factor, interleukin-6 (IL-6), IL-10, and lower coagulation factors, including antithrombin-III and factor IX. Men had lower survival compared with women at 30, 90, and 365 days.

The study design strengthens the results reported by this group. First, the study evaluated a large number of patients, 2183, in multiple clinical settings. This increases the likelihood that the results may be generalized to other patients. Second, only community-acquired pneumonia was evaluated rather than multiple different severe, lethal infectious diseases. With this design, the authors increased patient heterogeneity by examining a large number of patients and also increased homogeneity by restricting the study to community-acquired pneumonia. Third, the sophisticated statistical analysis of large, complex data sets with multiple adjustments for confounding factors increases enthusiasm for the work.

Several aspects of this study were anticipated, although there is still value in having a recent publication provide rigorous data documenting widely held beliefs. These beliefs include the observation that men are more likely than women to die of pneumonia. Additionally,

men were more likely than women to receive antibiotics within 8 hours of diagnosis further contributing to the literature demonstrating better health care delivered to men compared with women, although it must be acknowledged that there was no difference in the percent of patients receiving antibiotics for 4 hours.

IL-6 is another old friend and it has been widely reported as a biomarker of inflammatory diseases. Other investigators determined that plasma levels of IL-6 are elevated in patients with community-acquired pneumonia (4–6). A small study by Monton et al (7) evaluated cytokine concentrations both within the serum and the bronchoalveolar lavage of patients with pneumonia. Patients with pneumonia had higher levels of IL-6 than controls, and there was a trend toward higher serum levels of IL-6 in patients who died, and also generally higher IL-6 in the lavage fluid compared with serum. Another report by the GenIMS investigators (the same group reporting the findings in this article) showed that plasma levels of IL-6 at discharge also predict mortality over a 1-year period (8).

Numerous previous reports documented that in another lethal infectious disease, sepsis, high circulating levels of IL-6 correlate with mortality (reviewed in Ref. 9). It is probably safe to say that most investigators agree that biomarkers have utility to diagnose disease and predict outcome. It is also probably safe to say that most investigators strongly disagree about the best biomarker for predicting outcome. Another contentious area is whether the biomarkers merely predict clinical trajectory or if they actually cause the disease. In the cecal ligation and puncture model of murine sepsis, plasma levels of IL-6 strongly predict outcome (10), but probably do not actually induce mortality (11). Certainly, the low levels of plasma IL-6 observed in these patients with community-acquired pneumonia did not induce lung injury or mortality (3), even though the data clearly demonstrate that higher levels are associated with worse outcome.

IL-6 has also been reported to be elevated in patients with cardiac disease who

are at increased risk of dying or myocardial infarction (12). In the current study, significantly more men had cardiac disease (32%) compared with women (20%). This raises the possibility that the men with their preexisting cardiac disease may have had higher baseline levels of IL-6 that was further elevated by pneumonia.

IL-6 is classically considered a proinflammatory cytokine, whereas IL-10 is an anti-inflammatory cytokine, yet both were elevated in patients who would die. This creates a bit of a quandary for the classic paradigm that deaths caused by bacterial infections are secondary to an overwhelming proinflammatory response that injures cells, tissues, and organs. However, an alternative viewpoint would be that a proinflammatory response manifests itself by the host’s ability to produce something, anything, be it a proinflammatory cytokine or an anti-inflammatory cytokine (13). In other words, the capacity to synthesize new proteins not present in the circulation of normal individuals reflects the inflammatory status. Producing either IL-6 or IL-10 shows the capacity to respond. However, additional data in this article would argue against this viewpoint because coagulation proteins were found in patients more likely to die.

The important concept provided by this article is that older men dying of pneumonia have higher levels of the biomarkers tumor necrosis factor, IL-6, and IL-10. A cautious clinical implication of these findings is that men may be a more appropriate target for initial directed anti-inflammatory therapies compared with women.

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## \*See also p. 1655.

Key Words: pneumonia; inflammation; cytokines; sepsis; interleukin-10; tumor necrosis factor; biomarker; cardiac disease

The author has not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a1a12b

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## High-frequency percussive ventilation: An old mode with a great future\*

**H**ypoxemia and atelectasis are well-known and serious complications of one-lung ventilation (1). Intubation using a double-lumen tube facilitates independent lung ventilation, which alleviates hypoxemia, increases lung volume, and overcomes atelectasis in one-lung ventilation. However, the double-lumen tube does not function optimally in high-resistance airways and hinders access for suctioning secretions (2, 3).

High-frequency ventilation is characterized by breathing frequencies higher than 1 Hz where tidal volumes of 1–3 mL/kg are less than the dead space (4). High-frequency percussive ventilation (HFPV) is a mode of high-frequency ventilation that delivers small bursts of gas at 300–600 cycles per minute and relies on chest wall elastic recoil pressure for passive exhalation. HFPV is designed to be used in conjunction with mechanical ventilation or as a stand-alone treatment.

### \*See also p. 1663.

**Key Words:** one-lung ventilation; hypoxemia; lung resection; high-frequency percussive ventilation; lung secretions; mechanical ventilation

Dr. Blanch has received a Research Grant from Hamilton. He has a patent pending on a monitoring system. He serves on the Covidien Advisory Board. The remaining authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a0923a

The amplitude of pressure oscillations depends on the pulsatile flow amplitude and on the impedance of the respiratory system. Therefore, flow oscillations can be delivered on top of mechanical breaths or on top of spontaneous breathing (5, 6). Nowadays, high-frequency ventilation modes are a late option to sustain adequate gas exchange in adult patients with acute respiratory distress syndrome (4), and HFPV has led to improved oxygenation in cohort studies of patients with acute brain injury (7), acute respiratory distress syndrome (8), or acute smoke inhalation (9). In parallel, HFPV has been used to facilitate high-frequency-assisted airway clearance by vibrating the cilia layer. However, there is not enough data to consider HFPV superior to other techniques of high-frequency-assisted airway clearance like chest wall compression or chest wall oscillation (10, 11).

In this issue of *Critical Care Medicine*, Lucangelo et al (12) report the results of a randomized study on a novel use of HFPV in patients undergoing elective partial lung resection. After patients were placed in lateral decubitus and dependent lung ventilation was instituted, the nondependent lung was randomized to receive either humidified continuous positive airway pressure at 5 cm H<sub>2</sub>O or HFPV at a percussion rate of 500 cycles per minute and a mean pulsatile pressure of 5 cm H<sub>2</sub>O. After the surgical procedure was completed, patients had the same stan-

dard of care until hospital discharge. Interestingly, patients in the HFPV group had significantly better oxygenation and airway clearance of secretions, as well as a higher probability of being discharged earlier to the ward.

The study by Lucangelo et al (12) is important because it demonstrates that HFPV is efficacious not only in increasing Pao<sub>2</sub> during one-lung ventilation, but also in improving clearance of secretions, thus, enabling better outcome. These results from the well-controlled perioperative period might also have importance in the more general critical care arena. HFPV helps mobilize secretions from the periphery of the lung to larger airways; however, eliminating secretions depends on the patient's ability to cough them up or, in intubated patients, on the health-care team's ability to aspirate them. This is clear in the article by Lucangelo et al (12), where although the final amount of secretions was the same in both groups, it was eliminated 1 day earlier in the HFPV group. In critically ill patients, secretion removal is crucial. We can speculate that in patients with effective cough, early-assisted airway clearance might increase the number of ventilator-free days, decreasing episodes of nosocomial lung infections and extubation failures. In fact, Clini et al (13) recently reported that the addition of percussive ventilation to the usual chest physiotherapy routine in tracheostomized patients



improved gas exchange and expiratory muscle performance and reduced the incidence of pneumonia. However, these potential benefits might be totally lost in patients who are unable to expectorate, like patients with neuromuscular diseases or intensive care unit-acquired weakness, where secretions, although present in the airways, are extremely difficult to be mechanically suctioned.

Several points must be considered before the broad application of HFPV in mechanically ventilated critically ill patients. First, adding HFPV to conventional ventilation reduces humidity and a heated humidifier on the inspiratory line would be necessary to provide adequate humidification (14). Second, mechanical loads, such as those caused by alterations in resistance and compliance, affect flow, volume, airway pressure, and their waveforms (6, 14). Third, in conjunction with a driving ventilator, intrapulmonary percussive ventilation may add pressure and volume to tidal ventilation and generate intrinsic positive end-expiratory pressure (14). Fourth, before HFPV can be considered suitable for standard intensive care practice, evidence needs to be accumulated to demonstrate better patient-ventilator interaction with HFPV and the absence of significant adverse effects. Finally, when HFPV is added to current ventilators it could adversely affect the ventilator's ability to monitor pressures and volumes and may cause the ventilator alarm to go off incessantly.

In conclusion, HFPV is an old but very attractive technique. In short-term use, HFPV helps earlier secretion mobilization with some clinical benefits and no adverse effects. However, before broadly applying HFPV in intensive care patients, it is

necessary to determine which subgroups of patients might benefit most from HFPV, and further technological developments are necessary to ensure that the technique does not alter ventilator function.

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# Is lactate the “Holy Grail” of biomarkers for sepsis prognosis?\*

When oxygen delivery falls below a critical delivery or anaerobic threshold, the amount of oxygen available for consumption is not able to meet the needs of an organism. These delivery and consumption inadequacies result in cellular hypoxia. Under hypoxic conditions, mitochondrial oxidative phosphorylation fails and energy metabolism becomes dependent on anaerobic glycolysis (1). Anaerobic glycolysis sharply increases the production of cellular lactate that diffuses into the blood during prolonged cell ischemia. Elevated circulating lactate concentration thus indicates widespread inadequate tissue oxygenation due to inadequate oxygen delivery and/or consumption (2).

Numerous previous studies have documented the utility of lactate as a prognostic indicator in shock states (2, 3). The use of lactate measurements is gaining recognition in acute care medicine as a useful indicator of tissue hypoperfusion. In fact, several recent reports document the value of considering lactate levels for inclusion in aggressive resuscitation protocols for septic shock and in monitoring the response of septic patients to resuscitation (4–6). Furthermore, lactate screening as a method of risk stratification and prognosis has been shown to be beneficial in hemodynamically stable patients with suspected infection (7). Underscoring the importance of these data is the fact that point-of-care testing performed at the patient's bedside is becoming more common (8, 9).

In this issue of *Critical Care Medicine*, Mikkelsen et al (10) present a retrospective cohort study of emergency department patients with severe sepsis to investigate the association between initial

serum lactate, organ dysfunction, and shock. These investigators asked a particularly important question: are elevated lactate levels usually seen in subjects with clinically apparent organ dysfunction and shock? Said another way, does the value of a lactate level in septic patients remain, even in the absence of obvious clinical markers of illness severity (organ dysfunction and shock)?

The authors of this report present a methodologically sound study with meticulous detail in assuring the internal validity of the study by using standard definitions and recommended methods for retrospective chart reviews. The study was fairly large, with 830 subjects included. As has been previously reported (11), nonsurvivors had significantly higher initial levels than did survivors in subjects both with and without shock. The authors also found that when lactate levels are grouped (low <2, intermediate 2–3.9, and high >3.9 mmol/L), both intermediate and high levels are associated with significantly higher mortality compared with low levels, as has been previously reported (7). However, one interesting finding in this report (depicted in Fig. 2) was that the prognostic value reached a plateau in the nonshock group around 8 mmol/L, but that same plateau was not reached in the shock group until 18 mmol/L. This suggests that high lactate levels need to be interpreted in context of the clinical scenario and that simply thinking of lactate levels in compartments or groups (low, intermediate, and high) may marginalize the value of this test.

The most important finding in this report was that the association between lactate level and mortality was independent of clinically apparent organ dysfunction and shock. This finding substantially enhances our understanding of the prognostic value of an initial lactate level performed at the time of recognition of sepsis. These data indicate that the clinician should carefully consider the significance of an elevated lactate level in a patient who has no other clinically apparent signs of high acuity.

Many studies have shown that escalation of care in patients with severe sepsis

and hypoperfusion (manifested as persistent hypotension after fluids or a lactate  $\geq 4$  mmol/L) is associated with benefit (5, 4, 12–14). The study by Mikkelsen et al and others (11) have found even intermediate (2–3.9 mmol/L) lactate levels to be associated with higher mortality. What we do not presently know, and the question that this study does not answer, is what should we do with patients who have severe sepsis, are not in shock, and have an intermediate level (the group that makes up the largest lactate stratum in the study by Mikkelsen et al)?

In the end, Mikkelsen et al have presented a study that enhances our knowledge about the value of lactate in patients with sepsis. Namely, lactate measurements in patients with sepsis without clinically apparent organ dysfunction have important prognostic value. Given the cumulative data known about lactate to date, it makes one wonder if lactate is the “Holy Grail” of biomarkers for sepsis.

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\*See also p. 1670.

Key Words: lactate; sepsis; shock; organ dysfunction

Dr. Jones has received grant support from Critical Biologics and Hutchinson Tech. Dr. Puskarich has not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a09487

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## Keratinocyte growth factor in acute lung injury—A work in progress\*

In experimental and clinical studies of lung injury, improvement in alveolar epithelial function is associated with a reduction in pulmonary edema (1–3), indicating that a strategy to accelerate epithelial repair in acute lung injury (ALI) may be beneficial. Keratinocyte growth factor (KGF) modulates a variety of mechanisms recognized to be important in alveolar repair and resolution in ALI. Effects of KGF on alveolar epithelial cells *in vitro* include increased proliferation, increased surfactant protein production, and altered matrix metalloproteinase release. KGF also down-regulates proinflammatory cytokines and enhances endothelial cell resistance to injury (4). In animal models of ALI, KGF treatment reduces alveolar capillary permeability and pulmonary edema, and improves survival (5). Two clinical studies have reported KGF levels in patients with ALI. KGF was higher in nonsurvivors with more severe lung injury. This may indicate that the reparative activity associated with increased KGF in the setting of severe alveolar damage is insufficient, providing a rationale for augmentation of KGF levels as a therapy to restore alveolar function during injury (6, 7).

In this issue of *Critical Care Medicine*, Chandel et al (8) report the potentially interesting finding that bronchoalveolar lavage (BAL) fluid from a small cohort of patients with ALI decreases KGF messenger RNA (mRNA) expression and that this effect is mediated in part by transforming growth factor (TGF)- $\beta$ 1. However, shortcomings in the experimental design and data available limit the interpretation of the data.

In relation to the BAL procedure in this study, several important issues are worth highlighting. The authors report instilling 60 mL aliquots up to three times in keeping with published recommendations (9); however, it is unclear how the reported instilled volumes of  $90 \pm 32$  mL for the patients with ALI,  $72 \pm 26$  mL for the patients with cardiogenic edema, and 60 mL for the control subjects were achieved. Furthermore, although the authors acknowledge this as a limitation in the discussion, there was a 50% variation in the volume of BAL instilled in the patients with ALI compared with the control subjects. The authors argue that a larger volume instilled with the attendant greater dilutional effects on alveolar mediators might be expected to attenuate the ALI BAL fluid-induced fall in KGF mRNA expression. This does not hold good if a mediator that promotes KGF mRNA expression, such as interleukin-1 $\beta$ , tumor necrosis factor- $\alpha$ , or prostaglandin E2 (5), is diluted to a greater extent. The variable dilutional effects of using different BAL volumes is clearly a major weakness in this article. Another consideration is that the control subjects were not strictly healthy subjects with

normal lung parenchyma; it would have been reassuring to have data that the subjects had no underlying pathology. Finally, given that nonbronchoscopic lavage may not sample the alveolar space and does not provide similar results in the assessment of alveolar inflammation in patients with ALI compared with bronchoscopic lavage (10), the use of both bronchoscopic and nonbronchoscopic lavage is inappropriate in the same study. These limitations emphasize the need for standardization on the use of BAL in research in ALI in keeping with published recommendations (9) to allow data to be interpreted with confidence.

In this study, BAL fluid from patients with ALI decreased KGF at the gene level. However, a fundamental question remains as to how much translates to the protein level. In this study, although KGF levels in BAL from patients with ALI are reported, KGF levels from the patients with cardiogenic pulmonary edema and control subjects are not presented and, therefore, it is not possible to determine the effects on protein secretion. Furthermore, although discussed, the absence of a correlation between ALI BAL fluid KGF protein levels and KGF mRNA suppressing activity is not adequately explained. These data contrast with a recent study that comprehensively examined the effect of BAL fluid from a larger well-characterized cohort of patients with ALI, ventilated controls, and control subjects (defined as normal bronchoscopy, chest computed tomography, and microbiologic examination) on both KGF protein secretion and mRNA expression (11). In this study, Quesnel et al found the oppo-

\*See also p. 1678.

Key Words: ALI; keratinocyte growth factor; bronchoalveolar lavage

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a1a927



site effect, with up-regulation of KGF from fibroblasts by BAL from patients with ALI. A possible reason, which is discussed, for this difference relates to the differing time points of BAL sampling in the two studies (within 48 hours after intubation in this study compared with a mean of 4 days). However, clearly more data on temporal mRNA expression and protein levels would add to our understanding of the role of KGF during the course of ALI. This may have implications on the timing of pharmacologic treatment of ALI with exogenous KGF.

That the effect of KGF down-regulation is mediated by TGF- $\beta$ 1 is interesting but unconfirmed. TGF- $\beta$ 1 can be activated by many mechanisms and BAL fluid processing can lead to TGF- $\beta$ 1 activation and, therefore, may not accurately reflect *in vivo* activity (12). With the above caveat in relation to accuracy of TGF- $\beta$ 1 bioactivity measurement in BAL fluid *ex vivo*, TGF- $\beta$ 1 activity in BAL fluid was not measured in this study. The authors report historical data that transcriptionally active TGF- $\beta$ 1 is detected at concentrations up to 5 ng/mL in early ALI BAL fluid. Considering the effects of BAL dilution, this would approximate to 250–500 ng/mL in epithelial lining fluid. Therefore, it would have been more useful to use TGF- $\beta$ 1 levels in the *in vitro* experiments, which reflect the pathophysiologically relevant levels seen in patients with ALI.

KGF is available as a recombinant human protein and is approved for human administration in the treatment of oral mucositis associated with radiotherapy and chemotherapy and may represent a novel pharmacologic therapy in ALI. Although this study is interesting, further translational research is required to understand the role of KGF in the pathogenesis and resolution of ALI. These data,

if confirmed, support the concept that exogenous KGF may be a therapeutic option, particularly early in the course of ALI, to augment epithelial repair and overcome the inhibitory effect of the inflammatory milieu in the alveolar space in ALI on endogenous KGF production.

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# Steroids in cardiac surgery: Right time, right dose, right patient group\*

Cardiac surgery involving cardiopulmonary bypass (CPB) evokes a predictable systemic inflammatory insult. Despite modification of surgical techniques, improvement in biocompatibility of the bypass circuit, and advances in postoperative intensive care, cardiac surgery remains associated with significant postoperative morbidity—capillary leak, decline in cardiac function, disordered hemostasis, and multiorgan dysfunction. Corticosteroids have been used intermittently for their anti-inflammatory effects in sepsis and trauma, and perioperatively in cardiac surgery. Although administration of perioperative steroids (both immunosuppressive and physiologic doses) have consistently been associated with a reduction of circulating biomarkers following cardiac surgery, clear evidence of clinical benefit remains obscured by conflicting single-center study results (1–4).

In this issue of *Critical Care Medicine*, Weis et al (5) have added to the evidence suggesting that where selectively administered, perioperative continuous dosing with hydrocortisone may improve clinical outcome in adult patients undergoing CPB. In this prospective study, 36 high-risk adults undergoing cardiac surgery were randomized to receive placebo or 100 mg of hydrocortisone intravenously before induction of anesthesia, followed by 24 hours of continuous infusion of 10 mg/hr, with reducing dose during the following 3 days. Reflecting the advantage of single-center studies, anesthesia, CPB, and postoperative intensive care were standardized, and all patients received the anti-inflammatory agent aprotinin. In keeping with the primary end point, this study showed a significant reduction in circulating interleukin-6 levels at 4 and 24

hours in those patients receiving hydrocortisone, with higher levels of circulating interleukin-10 at the slightly earlier time points of 1 and 4 hours postoperatively.

Although the study was powered to detect a difference in postoperative circulating cytokine levels (interleukin-6/interleukin-10 ratio), the group of patients receiving stress dose hydrocortisone ( $n = 19$ ) had a trend toward increased postoperative indices of oxygenation, with a significant reduction in the incidence of atrial fibrillation, duration of catecholamine support, and intensive care unit stay. Of note, however, the improved oxygenation did not translate into shorter duration of ventilation. Therefore, the reader may conclude that the reduced intensive care unit stay was secondary to improved postoperative hemodynamic stability and, consequently, a decreased requirement for catecholamine support. Interestingly, the overall length of hospitalization was not significantly different in the two groups.

Mortality (although easy to define) is no longer an appropriate end point in clinical studies of cardiac surgery. Although morbidity is felt largely to be secondary to the patients' inflammatory response to surgery and CPB, acceptable and measurable end points are contentious. Acute measures of cardiac function and end organ perfusion are desirable, but despite a market flooded with technology to measure cardiac output (pulmonary artery catheter, NICO, PiCCO, LiDCO, CardioQ), we have failed to even show a correlation between cardiac output and outcome in critical care. Perhaps for the patient duration of ventilation, length of hospital stay and quality of life on discharge are what truly matter. However, many factors contribute to delay a patient's discharge and there is still no agreed measure for quality of life.

As a result, many studies rely on modified biomarker levels as evidence of effect and use correlations between biomarkers and outcome measures to suggest efficacy. Weis et al have added to the existing data to suggest stress dose steroids given to patients at risk of significant postoperative systemic inflammatory response syndrome

and organ dysfunction will attenuate the inflammatory response observed in adult patients undergoing CPB. However, a large study showing definitive association with clinical benefit is lacking. As raised by the authors in the discussion, conflicting results from previously published studies may reflect inappropriate selection of target population, type, dose, and dosing regimen of the steroid used. Thirty years of heterogeneous use of perioperative steroids in cardiac surgery has failed to consistently demonstrate benefit and allay concerns about harm. As in septic shock, the time has come for an adequately powered multicentered randomized control trial to scientifically test if perioperative corticosteroids are of clinical benefit to patients undergoing cardiac surgery, to identify those patients most likely to benefit and the most effective dosing regimen to use, and to dispel concerns of potential harm (6).

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\*See also p. 1685.

Key Words: cardiac surgery; corticosteroids; inflammation; interleukin-10; interleukin-6

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a09521

# The cycles of heart, lungs, and science\*

In 1988, Lemaire et al (1) published a landmark investigation into the cardiovascular effects of weaning. By using an array of invasive physiologic and nuclear radiographic measurements, they showed that some patients failed weaning because they developed left ventricular dysfunction and elevated pulmonary artery occlusion pressure (PAOP) on switching from positive pressure to spontaneous breathing. Fast forward to 2009, Dr. Teboul, who was then a junior author of that article, is now the senior author of a follow-up study by Lamia et al (2) published in this issue of *Critical Care Medicine*.

Lamia et al update the invasive diagnostics of 1988 by showing that certain echocardiographic findings during a spontaneous breathing trial correspond to a PAOP >18 mm Hg. This may have substantial clinical utility. On the basis of the work done by Lemaire et al, I suspect left ventricular dysfunction as a cause of weaning failure in patients whose respiratory mechanics would otherwise predict success. However, resting echocardiography may be normal, and I do not routinely insert a pulmonary artery catheter in such patients. Echocardiography during the stress of a spontaneous breathing trial might confirm weaning-induced cardiac dysfunction, and thereby guide therapy. In a letter to the editor, Teboul and Richard (3) had decried the lack of echocardiographic data allowing noninvasive evaluation of the left ventricular response to weaning. I applaud their effort to fill that void. This is not, however, the data from this study that I find most curious.

I was a fellow in training when Lemaire et al (1) published their seminal article. My

mentor, Sol Permutt, reviewed it and wrote an editorial (4). I impetuously questioned the reasoning in his editorial, prompting two decades of periodic friendly and stimulating argument. It is now I who has been given the opportunity to review and editorialize this follow-up article, and I am struck by some of the same findings that intrigued Dr. Permutt.

Patients who seemed ready to wean but had failed two spontaneous breathing trials were studied during a third trial. For this, they were changed from positive pressure ventilation with 5–7 mm Hg positive end-expiratory pressure (PEEP) to a T-piece. When changing from positive pressure with PEEP to spontaneous breathing without PEEP, pleural pressure should fall both during inspiration, due to patient effort, and during expiration, due to the loss of PEEP. One would also expect right atrial pressure (RAP) to fall, because pleural pressure surrounds the right atrium. In the study by Lemaire et al, the patients who manifested left ventricular dysfunction during weaning paradoxically *increased*, rather than *decreased*, RAP when they began breathing spontaneously. Although venous return will rise when pleural pressure falls, this could not cause RAP to rise above its baseline value, because it is the fall in RAP that augments venous return in the first place. Dr. Permutt reasoned that RAP was rising above its starting value because the diaphragm was now contracting around an engorged splanchnic and hepatic vasculature. This directed more blood back toward the heart than would lower pleural pressure alone.

In the current study, patients were divided into those whose PAOP either rose or failed to rise above 18 mm Hg by the end of their trial of spontaneous breathing. Among the latter, there was no change in RAP. However, once again, in the patients whose PAOP rose from a mean of 12 to 26 mm Hg, RAP rose from 5 to 12 mm Hg (both  $p < 0.05$ ; Table 1 in Ref. 2). If one conservatively estimates that pleural pressure fell by 2 mm Hg because of the loss of PEEP, the transmural filling pressure of the right ven-

tricle seems to have risen by 9 mm Hg at end expiration. What factors may have caused this?

First, contraction of the diaphragm seems, at best, a minor cause. Although Dr. Permutt may be rising to rebut as he reads this, the RAP is reported at end expiration when the diaphragm is relaxed. One might speculate that the inspiratory increase in venous return has not yet fully left the right side of the heart by end expiration, despite the passage of at least three heartbeats. However, this end-expiratory remnant of an enormous inspiratory increase in right ventricular preload should have induced a proportionally large increase in cardiac output. Yet, the stroke volume index in these patients fell slightly from 39 to 37 mL/m<sup>2</sup> (not significant; Table 1 in Ref. 2). Therefore, an increase in preload is an unsatisfying unitary explanation for the RAP elevation. Another potential explanation is that the observed increase in pulmonary artery pressure precipitated acute cor pulmonale. This is not supported by such a stable mean stroke volume, but the combined increase in right ventricular preload and afterload from all of the above mechanisms may certainly have combined to increase RAP.

A final explanation for the increase in RAP is that pleural pressure rose, rather than fell, during spontaneous breathing trials in these patients. Dynamic hyperinflation and expiratory muscle recruitment have been shown to increase both esophageal and PAOP when patients with chronic obstructive pulmonary disease exercise or hyperventilate (5, 6). Dyspneic subjects recruit expiratory muscles and can elevate PAOP even in the absence of air flow obstruction (7–9). In the current article, the same group of patients whose PAOP and RAP rose during spontaneous breathing also increased their pulmonary artery pressure (see Ref. 2, Table 1). A potential unifying explanation for this is that pleural pressure was the tide lifting all of these intrathoracic boats.

Regardless of whether RAP rose because of preload, left heart dysfunction, pleural pressure elevation, or a combination of factors, it might serve as another

\*See also p. 1696.

Key Words: heart–lung interactions; weaning; mechanical ventilation; hemodynamic monitoring

The author has not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a0961f



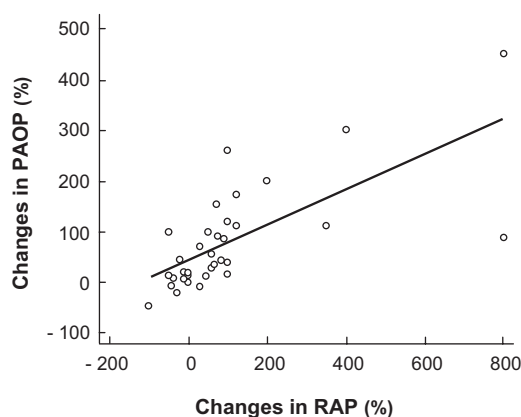


Figure 1. Relationship between percent changes in right atrial pressure (RAP) and pulmonary artery occlusion pressure (PAOP) induced by the trial of spontaneous breathing in the subjects in Ref. 2. Changes in the two variables correlated with each other ( $Y = 0.34x + 44$ ;  $r^2 = .47$ ,  $p < 0.05$ ).

marker of PAOP elevation. The authors kindly shared Figure 1, which shows that the increases in PAOP and RAP correlated with each other (Fig. 1). Although less accurate than the echocardiographic indices, they calculated that a  $>60\%$  increase in RAP predicted a PAOP  $>18$  mm Hg with sensitivity and specificity of 85% and 76%, respectively. In patients with central venous catheters or in hospitals lacking the echocardiographic expertise, observation of the weaning-induced change in RAP may also identify patients who could benefit from diuresis.

Lamia et al provide a beautiful illustration of the many cycles of heart–lung

interactions. To the respiratory cycle and the cardiac cycle, we can add the cycle of scientists. I hope that now one of my impetuous fellows will read these articles and challenge *my* reasoning.

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## Surviving critical illness is not without its perils: A perspective on depression in acute lung injury survivors\*

**T**he experience of surviving an acute life-threatening illness or sustaining a severe physical injury necessitating treatment in an intensive care unit (ICU) is inherently stressful and frightening. There has been a growing body of literature demon-

strating that surviving critical illness carries with it a substantial psychological burden, including depression. Recent systematic reviews of depression in general ICU and acute lung injury (ALI)/acute respiratory distress syndrome (ARDS) survivors found that 28% of patients had substantial depressive symptoms up to 2 years following hospital discharge (1, 2).

Several logical questions are stimulated by these investigations. What is the etiology of depressive symptoms in critical illness survivors? Are endogenous patient-related factors to blame? Are ICU service-delivery characteristics the primary culprit? Emerging studies are producing clues to this enigma. A pre-ICU

history of depression predicted depressive symptoms at 2 and 6 months post-ICU in one study of patients who survived acute respiratory failure (3). In the ALI/ARDS survivor population, longer ICU lengths of stay, durations of mechanical ventilation, and durations of sedation predicted later depressive symptoms (4, 5), a finding not duplicated in general ICU survivors (1). Also, ICU hypoglycemia predicted substantial depressive symptoms in ALI/ARDS survivors 3 months after hospital discharge in one recent study (6).

In this issue of *Critical Care Medicine*, Dowdy et al (7) provide further insights into the answers to these questions by reporting on the prevalence of, as well as

\*See also p. 1702.

Key Words: intensive care units; critical care; depression; respiratory distress syndrome; adult; outcome assessment (health care)

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DOI: 10.1097/CCM.0b013e3181a0967f

potential risk factors for, substantial depressive symptoms in the largest cohort of ALI/ARDS survivors assembled 6 months post-ICU. This study prospectively assessed depressive symptoms in 160 patients using the Hospital Anxiety and Depression scale. The point prevalence of symptoms suggestive of major depression in this sample was 26%, similar to prior studies on ALI/ARDS survivors (2). Also, Dowdy et al found that pre-ICU depressive and/or anxiety symptoms, admission to a surgical ICU, a maximum Sequential Organ Failure score of >10, and a single ICU service-delivery characteristic, a mean daily benzodiazepine dose of  $\geq 75$  mg of midazolam equivalent, all predicted a greater post-ICU depressive symptom burden. Studies on general ICU survivors have found that ICU benzodiazepine receipt may increase the risk of subsequent posttraumatic stress disorder symptoms (8). Dowdy et al are the first to find this association with post-ICU depressive symptoms.

It is important to note that this study does have limitations. First, a screening questionnaire, the Hospital Anxiety and Depression scale, was used to ascertain depressive symptoms. Although a screening questionnaire can assess the burden of symptoms, a diagnosis of major depression cannot be made without a structured or semistructured diagnostic interview. In addition, pre-ICU depressive and anxiety symptoms were assessed retrospectively, creating the possibility of recall bias because of current symptom burden. Furthermore, although ALI/ARDS is a common critical illness, it is possible that the findings of this study might not be generalizable to survivors of other critical illnesses. Also, Dowdy et al do not explain why admission to a surgical ICU may put patients at increased risk for subsequent depression, especially more so than patients admitted to trauma ICUs because patients who survive physical injuries are known to be at increased risk for subsequent depression (9). Finally, what explains the risk for post-ICU depressive symptoms conferred by benzodiazepine sedation? The findings by Dowdy et al do not answer this question. Are benzodiazepines the actual risk factor or are they a proxy for other possible risk factors? Patients with pre-ICU histories of depression and/or anxiety may be more likely to receive benzodiazepines during their ICU stay (10). Additionally, patients

taking benzodiazepines, which are commonly prescribed for the treatment of anxiety symptoms, before acute care hospitalization for physical injuries were found to be at increased risk of developing posttraumatic stress disorder 12 months after hospital discharge in a national multisite study of trauma survivors (11). Perhaps a prior history of depression and/or anxiety is what actually confers the greatest amount of risk for post-ICU psychopathology. Furthermore, benzodiazepines are a known delirigenic class of medications, and a prior history of depression seems to increase the risk of delirium while hospitalized (12, 13). Also, in-hospital delirium may predispose to subsequent depressive symptoms (14). Could in-ICU delirium be the actual risk factor for subsequent depression being suggested by benzodiazepine use? It is difficult to say, but merits further investigation.

The study by Dowdy et al on depression 6 months after surviving ALI/ARDS is a noteworthy contribution to furthering our understanding of the outcomes of patients who survive critical illnesses. It is only the second study to show that a pre-ICU history of psychiatric morbidity is associated with a significantly increased risk of depressive symptoms months after discharge from the hospital. Additionally, Dowdy et al highlight that an ICU service-delivery characteristic, benzodiazepine sedation, may increase the risk of subsequent depressive symptoms, suggesting a possible area of intervention and further study by using alternative sedative agents that might be less delirigenic (15), and stimulating research into other ICU service-delivery characteristics that may confer risk for psychopathology in critical illness survivors. Because millions of patients are admitted to ICUs annually in the United States (8), further research into the causes of post-ICU psychiatric morbidity is an important public health concern, and Dowdy et al provide an important step in the direction of furthering our understanding of the etiology of these problems.

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# Caveolin: Another reason why lipid membranes play an important role in ventilator-induced lung injury\*

Despite identification of numerous potential therapies for critical illness and lung injury, one of the few that has resulted in significantly reduced mortality is low tidal volume ventilation. The Acute Respiratory Distress Syndrome-Net Clinical trial (1) solidified the biological concept of volutrauma in humans and the need to limit it by applying tidal volumes appropriate for an injured, “baby lung” and was the fruition of decades of investigation into the role of mechanical forces in alveolar-capillary barrier injury (2). Although mechanical force is the initiating insult, this force must be transduced into biological effects that are a (im)balance of injurious and protective responses. The nature and balanced effect of these transduction mechanisms are complex and remain incompletely understood. In this issue of *Critical Care Medicine*, Hoetzel et al (3) highlight two novel and interacting etiological factors: carbon monoxide (CO) and caveolin (Cav).

The authors had previously identified a protective effect of CO in ventilator-induced lung injury (4), and therefore this report is novel and exciting in so far as Cav has been identified by them as a potentially important modulator of mechanical injury to alveolar epithelial cells.

That caveolae, which are lipid membrane structures, might play a role in cell injury responses is not, in retrospect, surprising because there has been accumulating evidence for the role of lipids in promoting and inhibiting ventilator-induced lung injury (5). Initially thought to be a simple barrier between the intracellular and extracellular environments, it is now clear that the plasma membrane is in fact a complex array of lipids and

lipoproteins, and that it is a dynamic structure that through lipid-lipid and lipid-protein interactions directly interacts with and influences both the intracellular and extracellular environments (6). Ceramide is just one example of such a membrane lipid involved in lung injury. Investigators showed that cellular stresses can induce ceramide's enzymatic activation and, in turn, its biological activity, which plays an important role in facilitating pulmonary edema and inflammatory cytokine production (7, 8). Furthermore, lipids and their movement to and from the plasma membrane play a protective role. It has been shown both *in vitro* and *in vivo* that alveolar cell membrane damage resulting from injurious mechanical forces can be both prevented and repaired by the active process of lipid membrane trafficking from intracellular compartments to the stressed cell surface (4, 9).

In the last decade, it has also become clear that there is a “domain” structure within the lipid membrane that modulates endocytic and exocytic processes and transmembrane protein functions (10). Caveolae, which represent one such membrane domain structure, are characterized morphologically by flask-like membrane invaginations and biochemically by cholesterol enrichment and abundance of the resident protein, Cav. Cav is a multifunctional protein that can serve both directly and indirectly as a scaffold and signaling protein to facilitate effective communication between and within the intracellular and extracellular environments (6). It seems to play a key role in lung biology because knock out of the protein in mice results in a lung injury phenotype characterized by increased lung weight and epithelial cell proliferation (11). Increasingly, Cav has been implicated in human diseases, including cancer (10). In this issue of *Critical Care Medicine*, Hoetzel et al show that it may also be important in deformation-induced epithelial cell function. Like all reports that provide novel data, this article also raises numerous important

mechanistic questions and, in turn, the potential for bench-to-bedside translatability.

First, is Cav a mechanotransducer that is able to sense and convert mechanical forces to measurable and biologically relevant processes such as cytokine production or injury repair? Previous studies have implicated Cav as a pulmonary endothelial mechanotransducer (12), but this study does not directly address whether this is true for the pulmonary epithelium. Presumably, Cav functions similarly in pulmonary epithelium; certainly, it is plausible given Cav's integral position within the plasma membrane, if not in a direct fashion, at least indirectly through the influence of factors such as CO. This, in turn, raises the question whether Cav functions simply as a facilitator of CO's protective effect or has inherent protective effects. Both would seem plausible given the ubiquitous expression of Cav and its numerous interactions with proteins that have previously been implicated in ventilator-induced lung injury (12) and the extent of injury in lungs ventilated in the absence of CO.

Second, the nature of CO-Cav interaction both physically and biochemically remain incompletely understood. The transduction mechanism that underlies the “crosstalk” that occurs between CO and Cav will be important to delineate. The fact that the effect of CO requires Cav in smooth muscle cells suggests their interaction may not be cell type specific (13). Furthermore, understanding whether they function in a synergistic, additive, or redundant fashion is important in tailoring sensible pharmacotherapies. This can be considered much like chemotherapy, which may be either aimed at one particular mechanistic pathway or multiple.

How might all this basic science be relevant for the bedside? The answer to this may lie in one of the mechanisms of lung injury: surfactant lipid dysfunction. Surfactant composition and surface tension properties are markedly abnormal in all forms of acute lung injury and there

\*See also p. 1708.

Key Words: caveolin; lipids; acute lung injury; ventilator-induced lung injury; carbon monoxide

The author has not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a1a028



have been clinical attempts to supplement the surfactant pool with exogenous lipid-lipoprotein mixtures. However, solid clinical trial evidence that this therapy is effective have been lacking (14). It may not be because of poor biological reasoning but rather the lack of necessary compounds or proteins within the supplemented lipid emulsion (15). Other than the surfactant lipoproteins, one other such protein may be Cav-1 along with CO. In conclusion, this report, in conjunction with others in the pulmonary vasculature, establishes Cav as a novel protein to be considered in the treatment of acute lung injury.

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## Vascular dysfunction in septic shock—Any ROCKing news?\*

Septic shock remains a challenging clinical problem with very high morbidity and mortality rates (1). The dominant hemodynamic feature is a persistent vasodilation that is refractory to fluid resuscitation and vasoconstrictors. Current treatment guidelines aim to restore the effective circulatory volume and arterial blood pressure by the administration of intravenous fluids and vasoactive drugs (2). Such a protocol-driven, early goal-directed therapy has significantly improved outcome (3), yet, the exact pathophysiological mechanisms behind vascular dysfunction in sepsis shock remain unknown.

Frontline research in vascular biology has revealed new insights into the complex interaction between the vascular endothelium and vascular smooth muscle (4, 5). In this issue of *Critical Care Medicine*, da Silva-Santos et al (6) add another puzzle to this picture by providing novel data on the functionality of Rho-A/Rho-kinase (ROCK) in mesenteric vessels in rats suffering from sepsis.

In normal vessels, activation of Rho-A leads to stimulation of ROCK, which, in turn, can phosphorylate and subsequently inactivate the myosin light chain phosphatase, leading to myosin light chain phosphorylation, actin-myosin interaction, and cell contraction (7). The authors aimed to investigate whether a reduced activity of the Rho-A/ROCK pathway could be involved in the hypotension and decreased reactivity to vasoconstrictors observed in patients suffering from septic shock. Interestingly, they were able to demonstrate an impaired Rho-A/ROCK-mediated phosphorylation of my-

osin phosphatase targeting subunit (MYPT)1 (the regulatory subunit of the smooth muscle myosin phosphatase) in vascular smooth muscle from endotoxemic-treated rats. Furthermore, data from this study also suggest that changes in cyclic guanosine monophosphate-dependent mechanisms involved in calcium sensitization might play an important role in the vascular response to sepsis. Although the present *in vitro* study is far removed from clinical utility, da Silva-Santos et al deserve credit for opening up another niche in this research field.

Outcome in septic shock is strongly related to timing of treatment and rapidly instituting cardiovascular support and antibiotic therapy (8). Current management guidelines suggest noradrenalin or dopamine as first-line vasopressors, and adrenaline as second-line treatment, if blood pressure is poorly responsive (2). One might argue that it is the timing, rather than the specific agents, that is likely to make the difference (9). How-

\*See also p. 1716.

Key Words: septic shock; Rho-kinase; endothelium; nitric oxide; sepsis

The author has not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a09408

ever, this will remain rather speculative until we gather more information about the functionality of all relevant vascular pathways in septic shock. The article by da Silva-Santos et al contributes significantly in this context.

The focus for this study was downstream from nitric oxide/guanylate cyclase complex. Accordingly, the authors were unable to fully explore the relationship among the endothelium, nitric oxide, and the inhibition of the Rho-A/ROCK pathway seen in the current study. However, the effect of lipopolysaccharide injection on Rho-A/ROCK mediation of vasorelaxation was demonstrated. Furthermore, the nonspecific nitric oxide synthase inhibitor N<sup>ω</sup>-Nitro-L-arginine methyl ester hydrochloride and the inducible nitric oxide synthase inhibitor 1400W effectively blocked this effect, although the 6-hour and 24-hour results are interestingly distinct. However, the most exciting result was the demonstration of involvement of the Rho-A/ROCK system in altered vascular adrenergic responsiveness in the presence of lipopolysaccharide. The authors' findings revealed an impaired Rho-A/ROCK-mediated phosphorylation of MYPT1 in vessels from endotoxemic animals in a cyclic guanosine monophosphate-dependent manner, and subsequent studies clearly showed that blockage of the soluble guanylate cyclase enzyme normalized not only the vasodilatory responses to Y-27632, but also the levels of phosphorylated MYPT1, an indication of Rho-A/ROCK activity.

These data provide a sequence of evidence indicating that the overexpression of myosin phosphatase, and the inability of the Rho-A/ROCK to inhibit it, is a potential important mechanism involved in the vascular response in this model.

Fasudil, another ROCK inhibitor, has been used in several other studies in the past (10), but is not more selective than Y-27632. On the other hand, Y-27632 seems to be a very selective inhibitor of ROCK and works with similar potency for both isoforms, ROCK I and ROCK II (10). Furthermore, Y-27632 is at least twice more potent than fasudil and the choice of Y-27632 for this purpose seems appropriate. So, what is the ROCKing news in this story? Well, da Silva-Santos et al add novel observations to the existing literature, which hopefully will generate new challenging hypotheses and experimental work within an incredible complex research field. In the meantime, we should stick to the current guidelines, look upstream to understand the sequence of events, and start therapy sooner rather than later.

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## How to plug a leak?\*

**T**he incidence of sepsis increased 9% per year between 1979 and 2000 in the United States (1). Infection caused by bacteria accounts for approximately 50%

to 80% of all septic cases (2). Certain bacteria, via their microbial-associated molecular patterns such as endotoxin (lipopolysaccharide [LPS]), are recognized by specific receptors such as danger signals by immune cells, epithelial cells, and endothelial cells. The reaction of these and other cells induces sequential immune reactions including the cytokine expression and response cascade. The subsequent imbalance between proinflammatory and anti-inflammatory mediators causes severely impaired immune functions (3, 4). Pathogen-induced compromise of the integrity of epithelium and endothelium results in the further

breakdown of immune defense and serves as another source of cytokine activation (3, 5). LPS can disrupt intestinal epithelial tight junctions via increased inducible nitric oxide synthase activity (6) and causes breakdown of the endothelial barrier by altering junctional complexes (5). These events may promote bacterial translocation and fluid sequestration, worsening the course of sepsis toward multiorgan system failure. When epithelial integrity is functionally disrupted, mediators including nitric oxide, tumor necrosis factor, or interferon  $\gamma$  (7) might reduce the tight junction protein zonula occludens 1, together with the internal-

\*See also p. 1735.

Key Words: sepsis; lipopolysaccharide; endothelium; small GTPase Rac 1; cAMP

Supported by NSC95-2320-B-182-030-MY3, CMRPD140152, and EMRPD170581.

The author has not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a096c9

ization of several apical junctional complex transmembrane proteins (8). When endothelial barrier breakdown is structural, thrombin might alter actin dynamics and, thus, junctional components, such as adherens and tight junctions in the endothelium (9).

Although many factors have been identified in the progression of pathogen-induced endothelial dysfunction, the signaling pathway initiating the dysfunction of endothelial barriers during sepsis is still in question. The article authored by Schlegel et al (5) has made a significant advance toward answering this question. Combining studies from single-microvessel perfusion of mesenteric postcapillary venules and from cultures of human dermal microvascular endothelial cells, the investigators were able to determine hydraulic conductivity as an index for endothelial permeability in addition to visualizing and quantifying various junctional proteins, and functionally determining dextran permeability and transendothelial resistance. The investigators discovered that increased cyclic adenosine monophosphate (cAMP) abolishes the LPS-induced increase in endothelial permeability and intercellular gap formation. Furthermore, LPS inactivates small guanosine triphosphosphate (GTP)ase Rac 1 rapidly with barrier dysfunction, whereas Rho A activity increase was delayed indicating that Rac 1 rather than Rho A is involved in the LPS-induced disruption of endothelial barrier. Previously, the observation that inhibition of Rho A could inhibit the LPS-induced contractile pattern of endothelial cells in lung edema (10, 11) suggested a role for Rho kinase in LPS-induced endothelial leakage. Schlegel et al (5) showed that no significant activation of Rho A occurs (2 hours post-LPS) despite a many-fold increase in endothelial permeability, both *in vivo* and *in vitro*. This time-course independence is consistent with the lack of effect of Rho kinase inhibitor on LPS-induced endothelial permeability (12). In

contrast, both cAMP decrease and Rac 1 inactivation seem to be an early event (1 hour post-LPS) associated with LPS-induced endothelial barrier breakdown. Thus, the authors propose that both Rac 1 inactivation and cAMP decrease are important for changes in junctional proteins (e.g., fragmentation of vascular endothelial-cadherin and decrease of claudin 5), leading to the formation of large intercellular gaps and increasing endothelial permeability following LPS treatment.

What does it all mean for clinical sepsis? *In vivo* tests of agents interfering with cAMP level and/or Rac 1 activity during experimental and clinical sepsis should provide more evidence to justify the application of the current findings. Recent clinical investigation using cAMP-increasing agents did show beneficial effects on microcirculatory alterations in experimental endotoxemia (13) and patients with septic shock (14). The knowledge on how endothelial cells regulate junctional proteins may provide clues for the therapeutic potential to stop a leak during sepsis.

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## ACKNOWLEDGMENTS

I thank Drs. Ronke Olabisi and David Lau for editing this article.

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# Thrombin-activatable fibrinolysis inhibitor-a inhibitors: Drugs for sepsis or drugs for disseminated intravascular coagulation?\*

In this issue of *Critical Care Medicine*, Muto et al (1) report on experiments with a novel thrombin-activatable fibrinolysis inhibitor-a (TAFIa) inhibitor in two sepsis models in rats. TAFIa, a carboxypeptidase, cleaves C-terminal lysine and arginine residues from partially plasmin-degraded fibrin, thus, inhibiting the binding of plasminogen, tissue plasminogen activator (tPA), and plasmin to the fibrin. This reduces the cofactor activity of fibrin in tPA-induced plasminogen activation and the efficiency of further degradation of the fibrin by plasmin (2). In addition, the cleavage of critical C-terminal lysine residues from partially degraded fibrin increases the proportion of plasmin that is inactivated by alpha-2-plasmin inhibitor (3).

Activation of TAFI requires high concentrations of thrombin and is strongly enhanced by thrombomodulin present on the endothelium. Theoretically, inhibitors of TAFIa should enhance fibrinolysis and this may be beneficial in sepsis, where intravascular fibrin formation may lead to microvascular thrombosis, which then causes organ dysfunction. The authors report that administration of the novel TAFIa inhibitor reduces fibrin deposits in the kidney and the liver of the experimental animals as well as laboratory markers of organ dysfunction in an endotoxemia model, and reduces the systemic inflammatory response in a bacterial infection model.

Fibrin deposits in kidneys, liver, and other organs are a sign for the presence of disseminated intravascular coagulation (DIC) in sepsis (4). DIC is an acquired syndrome characterized by the intravascular activation of coagulation with loss

of localization arising from different causes. It can originate from and cause damage to the microvasculature and, if sufficiently severe, can produce organ dysfunction (5). The diagnosis of DIC is based on the combination of a typical underlying disease, such as sepsis, with laboratory markers, including prothrombin time, fibrinogen concentration, platelet count, and a fibrin-related marker, reflecting intravascular fibrin formation (5–7). D-dimer, fibrin degradation products, or soluble fibrin are used as fibrin-related markers in the DIC scores.

Approximately 30% of patients with severe sepsis display overt DIC (8) and patients with severe sepsis fulfilling the DIC criteria display a considerably higher rate of organ dysfunction and an increased mortality (9–12).

Some patients with specific bacterial infections generate sepsis-induced purpura fulminans, which is characterized by extensive microvascular thrombosis, tissue necrosis, and secondary hemorrhage into the necrotic tissue, including the skin (13, 14). The lesions resemble those found in newborns with severe protein C deficiency (15). Laboratory data show that sepsis-induced purpura fulminans is caused by a failure of the protein C system in the context of sepsis-induced DIC. Otherwise, the clinical signs of DIC in severe sepsis are less obvious, with multiorgan dysfunction and minor bleeding, especially from insertion sites of intravascular catheters, and other tissue lesions, being the principal symptoms.

TAFI levels decrease in the course of DIC, presumably because of consumption of the proenzyme, leading to increased TAFIa activity (16), whereas plasminogen activator inhibitor (PAI)-1 levels are elevated in response to acute-phase reaction and platelet activation. Organ dysfunction and mortality in sepsis-induced DIC are closely linked to the functional integrity of the fibrinolytic system. In patients without multiorgan dysfunction, markers of coagulation activation are correlated with markers for the activation of fibri-

nolysis, indicating efficient clearance of intravascular fibrin (17). In patients with multiorgan dysfunction, in contrast, coagulation activation is not associated with a compensatory activation of fibrinolysis. Elevated levels of PAI-1 are a frequent finding in patients with sepsis. PAI-1 acts as an acute-phase reactant and additional PAI-1 may be released from activated platelets. Genetic variants leading to higher PAI-1 levels are associated with worse outcome in severe bacterial sepsis (18, 19). In experimental animals, administration of tPA reduces sepsis-induced fibrin deposition in the kidneys (20). A similar effect is observed on administration of a PAI-1 inhibitor (21). In view of these experiences, it seems logical to administer profibrinolytic drugs to patients with sepsis-induced DIC to prevent multiorgan dysfunction caused by microvascular thrombosis.

But is a TAFIa inhibitor indeed a promising approach?

This clearly depends on the individual condition. Treatment with high doses of antithrombin concentrate did not result in a clinical benefit concerning survival or organ dysfunction in patients with severe sepsis (22), but a *post hoc* analysis of the same trial revealed that high-dose antithrombin therapy clearly improves the clinical outcome in patients with severe sepsis and DIC (23). Selection of patients on the basis of a DIC score system would, therefore, limit the treatment to those patients who would be expected to benefit.

In patients with sepsis, normal or high fibrinogen levels indicate bad clinical prognosis (24), reflecting suppressed fibrinolytic response mainly because of elevated PAI-1 levels, but possibly also increased generation of TAFIa. In patients with high TAFIa activity, the soluble fibrin complexes and fibrin degradation products in the blood do not effectively promote plasminogen activation by tPA and binding of plasmin. A TAFIa inhibitor may restore the functional properties of the fibrin derivatives. In 1986, Wiman

\*See also p. 1744.

Key Words: TAFIa; disseminated intravascular coagulation; sepsis; thrombin

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a09708

and Ranby (25) developed a functional soluble fibrin assay, based on the cofactor effect of soluble fibrin in tPA-induced plasminogen activation. This assay showed rather variable results in patients with DIC (26, 27), indicating that the soluble fibrin present in clinical plasma samples differed from soluble fibrin prepared *in vitro* for calibration purposes. The differences may be due to TAFIa-induced modifications of the fibrin compounds detected by the assay. For proper selection of patients for treatment with a TAFIa inhibitor, a functional soluble fibrin assay based on the cofactor activity of the fibrin in tPA-induced plasminogen activation, or other markers of fibrinolysis activation, such as plasmin–plasmin inhibitor complexes, may be helpful if used in conjunction with laboratory markers of coagulation activation.

In contrast to anticoagulant drugs, treatment with a TAFIa inhibitor would be expected to enhance the profibrinolytic response caused by systemic coagulation activation without reducing the amount of fibrin available as cofactor in tPA-induced plasminogen activation or reducing the stimulation of tPA release induced by thrombin.

A certain bleeding risk may also be associated with TAFIa inhibitors, especially in hyperfibrinolytic conditions. If these conditions are identified by appropriate laboratory assays, treatment risks may be minimized.

In conclusion, TAFIa inhibitors might not be drugs to be used in sepsis in general. We would expect the clinical indication for a TAFIa inhibitor in patients with sepsis to be the patient with disseminated coagulation activation not appropriately compensated by activation of the fibrinolytic system. Proper selection of patients is essential to ensure maximal clinical benefit.

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# Can postoperative delirium be prevented pharmacologically?\*

**D**elirium occurs in up to 60% of patients after surgery (1). Patients at greatest risk for experiencing postoperative delirium (POD) include those undergoing a major cardiac procedure, who are elderly, have a preexisting cognitive abnormality or who require admission to an intensive care unit (1, 2). POD is associated with increased mortality, a longer hospital stay, and cognitive deterioration and, thus, is important to prevent (3). The hypermetabolic inflammatory state in the brain that occurs secondary to anesthetic-induced vasodilation is felt to be the initial insult inciting POD (1). The resulting oxidative stress leads to abnormalities in the cholinergic, dopaminergic, histaminergic, and noradrenergic neuronal systems. The ability of cholinergic cells to synthesize and release acetylcholine may be reduced leading to disorientation and memory impairment (4). The resulting increased release of dopamine may cause agitation and hallucinations.

Efforts by clinicians to reverse potential causative factors for POD are often not successful and patients who develop POD are frequently treated with psychoactive medications (5). Given the limited evidence that neuroleptic agents improve outcome in patients with POD and the safety concerns associated with their use, efforts to identify a pharmacologic strategy that can prevent POD are increasing (6). However, POD prophylaxis studies conducted to date that have focused on the pharmacologic modulation of the neurotransmitter abnormalities known to induce POD have failed to demonstrate a benefit (7–9). One large randomized trial of low-dose haloperidol in a population of elderly hip surgery patients did

not reduce the incidence of POD (7). Two smaller randomized studies in patients undergoing elective joint-replacement surgery failed to demonstrate a benefit with the prophylactic use of the acetylcholinesterase inhibitor donepezil (8, 9). There are a number of possible reasons that may account for the lack of benefit observed in these studies. The prophylactic regimens were studied in patients undergoing an elective orthopedic surgical procedure (a population at lower risk for POD than patients undergoing cardiac surgery), an inadequate treatment dose may have been used, the duration of therapy may have been too short and therapy was not initiated before surgery.

In this issue of *Critical Care Medicine*, Gamberini et al (10), despite addressing many of the methodologic concerns of previous POD prophylaxis studies, present the results of a clinical trial that fails to demonstrate the benefit of rivastigmine in preventing delirium after elective cardiac surgery. In this randomized, double-blinded, placebo-controlled study, rivastigmine (administered 1.5 mg every 8 hours starting on the evening before surgery and continued for a total of 6 days) failed to reduce the incidence of POD at 6 days (32% vs. 30%,  $p = 0.8$ ) or improve tests of cognitive function, such as the Mini-Mental State Examination or the Clock Drawing Test. For those patients experiencing POD, neither the time to onset of POD after surgery nor the duration of POD differed between the rivastigmine and placebo groups. The effect of rivastigmine on POD severity using a delirium rating scale was not measured. Use of rescue haloperidol and lorazepam was similar between groups.

Can the lack of benefit observed with this study be attributable to specific methodologic issues or does it support the findings from previous studies demonstrating that acetylcholinesterase inhibitors do not prevent POD? Although the investigators did not report a difference in the rate of POD that was felt to be clinically important, delirium occurrence was only 50% of that estimated in the investigator's sample size calculation, suggesting that lack of power may have

accounted for the lack of benefit observed. Reasons for the low incidence of POD observed are unclear, given that subject demographics, including severity of illness, are similar to those of other cardiac surgery studies (11). The low dose of rivastigmine used in the study and the lack of a rapid dose titration schedule in those patients without rivastigmine-associated side effects may also have accounted for the lack of benefit observed. It is, therefore, conceivable that methodologic factors may have accounted for the lack of benefit that was observed in the study. The study by Gamberini et al, therefore, does not conclusively rule out the benefit of acetylcholinesterase inhibitors in preventing POD but does suggest that any treatment benefit, if one exists, is likely small.

Future studies are, therefore, required to determine whether there is a pharmacologic intervention that will prevent POD in surgical populations at high risk for experiencing delirium. It is entirely possible that modulation of acetylcholine is not the only neurotransmitter-related mechanism that may play a role in preventing POD, given the important role that other neurotransmitters, such as dopamine, serotonin, and glutamine, play in the pathogenesis of POD (1). Therefore, a POD prophylactic regimen that modulates multiple neurotransmitters (e.g., acetylcholine and dopamine) may prove to be of greatest benefit. Future investigations in this area should also evaluate the pharmacodynamic response to any pharmacologic intervention to help answer important questions that exist for any potential prophylactic regimen, including the optimal dosing regimen that should be used and when therapy should be initiated relative to surgery.

The anesthetic regimen chosen for surgery and the sedation and analgesia administered both intraoperatively and postoperatively may each affect the incidence of POD (1). For example, sedation with dexmedetomidine is associated with less delirium than lorazepam (12). It may be possible that subsyndromal delirium is a better outcome than delirium for future prophylactic studies, given that it has

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## \*See also p. 1762.

Key Words: rivastigmine; delirium; postoperative delirium; prevention; surgery; cardiac surgery; intensive care unit

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a09731



been found in the intensive care unit to be associated with outcomes that are nearly as bad as delirium (13). Hopefully, a beneficial POD preventive strategy will be identified once the results from a number of ongoing studies become available. Ongoing trials in this area include two studies that are exploring the use of acetylcholinesterase inhibitors with or without concomitant haloperidol prophylaxis, another study that is evaluating olanzapine, and one that is comparing the incidence of POD in cardiac surgery patients receiving dexmedetomidine or propofol sedation regimens (14). With these ongoing trials, and other future investigations, it is hoped that a pharmacologic intervention will be identified that can reduce the incidence of POD and the morbidity and mortality associated with it.

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## Should we be afraid of the Green Monster?\*

**P***seudomonas aeruginosa* is an incredibly versatile opportunistic pathogen that can infect most organs and tissues in humans. The most notorious and/or life-threatening *P. aeruginosa* infections include lung infections in patients with cystic fibrosis, diabetic foot ulcers, urinary tract infections in catheterized patients, and bacteremia in the severely burned (1). Because of its large genome and genetic adaptability (2), *P. aeruginosa* possesses many conserved genes that allow it to survive almost anywhere, under almost any conditions, express an

extensive arsenal of virulence factors, and produce enzymes that convey resistance to many antimicrobials. This dynamic bacterium even produces its own antibiotics, including pyocyanin, which kill competing microbes and gives *P. aeruginosa* cultures, and sometimes infections, their characteristic green/blue appearance (1). In this issue of *Critical Care Medicine*, Veessenmeyer et al (3) discuss potential future *P. aeruginosa* therapeutics that focus on antagonizing specific virulence determinants. However, one could ask “Are alternatives to antibiotics really needed to treat *P. aeruginosa* infections and do we really need to be afraid of this Green Monster?”

The latest news on *P. aeruginosa* is a mixed bag of good and bad. The good news is that over the last 25 years the antibiotic resistance of *P. aeruginosa* does not seem to have increased (4, 5). In fact, the most recent data from the European Antimicrobial Resistance Surveillance System indicate that percentages of

the antibiotic-resistant *P. aeruginosa* strains decreased between 2002 and 2006 (4). These data demonstrated little change in resistance profiles for meropenem and imipenem, and a notable decrease in resistance to piperacillin/tazobactam, ceftazidime, gentamicin, and ciprofloxacin (4). Another piece of good news, if you live in the United States, is that resistance rates for *P. aeruginosa* nosocomial infections are much lower in the United States than in Europe (6). For example, according to data from the International Nosocomial Infection Control Consortium (2002–2007) and the US National Nosocomial Infections Surveillance System (1992–2004), the rates of *P. aeruginosa* isolates resistant to piperacillin, ciprofloxacin, imipenem, and ceftazidime were 50.8 vs. 17.5%, 52.4 vs. 34.8%, 36.6 vs. 19.1%, and 51.7 vs. 13.9% in Europe vs. the United States, respectively (6).

The bad news is that the overall rate of *P. aeruginosa* infections is increasing and

\*See also p. 1777.

Key Words: *Pseudomonas aeruginosa*; nosocomial infection; antibiotic resistance; quorum sensing; multidrug resistance; opportunistic pathogen

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DOI: 10.1097/CCM.0b013e3181a092bc

quite dramatically in some patient populations (7). For example, the percentages of *P. aeruginosa* isolates associated with pneumonia, surgical site infections, and urinary tract infections have essentially doubled since 1975 (7). These data are not surprising. The patient population is growing older and because of new efficacious treatments for chronic and malignant diseases, there is a rapidly expanding pool of immunodeficient patients who face prolonged courses of antibiotics and stays in intensive care units (8). Furthermore, with the United States engaged in two ongoing wars, the wounded patient population, which has been notoriously prone to *P. aeruginosa* infections, has greatly expanded (7). Therefore, the numbers of multidrug-resistant and pan-drug-resistant *P. aeruginosa* isolates are likely to rise. The majority of multidrug-resistant *P. aeruginosa* strains are resistant to the most commonly used antibiotics, including aminoglycosides, fluoroquinolones, and  $\beta$ -lactams (8). Although several new drugs with activities against multidrug-resistant Gram-positive bacteria have recently been marketed, few exhibit activity against Gram-negative bacteria (8). With few antibiotic options available or even in development, physicians will face a future with increasing multidrug-resistant cases and a scarcity of treatment options. Therefore, it seems we should indeed be afraid of the Green Monster, *P. aeruginosa*.

Although Veesenmeyer et al (3) have identified several potential strategies for targeted anti-*P. aeruginosa* treatments in this issue, the source from which these advances will come is currently unclear. Many pharmaceutical companies are scaling back or even abandoning antibiotic development, primarily because antibiotics pose a low return on investment (8). This is partially due to heightened regulatory requirements for approving antibiotics for human use, which add to the enormous cost of development. It has

been estimated that the development of a new antimicrobial can cost up to one billion dollars (9). Recouping these costs can be difficult because most antibiotics are used for short treatment courses and can quickly be made obsolete by resistance-acquiring microbes. Although research in academic laboratories has yielded promising data on specific targets, such as quorum sensing, type III secretion, and biofilm formation for *P. aeruginosa* (3), academic settings are generally poorly suited to perform the high-throughput screening of extensive chemical libraries required for the identification of effective new drug treatments.

Despite the lack of antimicrobial innovation in the last few decades, a new dawn of Green Monster slaying may be at hand. The recent "Pay for Performance" plan enacted by the Centers for Medicare and Medicaid Services will cease paying hospitals for some of the care made necessary by "preventable complications" (10). Among these complications are many types of nosocomial infections caused by *P. aeruginosa*. Consider, for example, the cost of a recent *P. aeruginosa* outbreak that affected 17 patients in an intensive care unit in Spain (11). Among these patients, the mortality rate was 47%, and the added extra length of intensive care unit stay attributable to *P. aeruginosa* infection was 70 days at an additive cost of more than \$400,000 (11). In the United States, the average Medicare payment for admissions in which a *P. aeruginosa* catheter-associated urinary tract infection is present is more than \$40,000/patient (10). Thus, if current nosocomial rates remain high, hospitals will have to start absorbing huge costs. This scenario may have a silver lining that results in improved standards of care as well as putting pressure on pharmaceutical companies to develop improved prophylactics and new antimicrobials.

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# Eighteenth century mindsets, twenty-first century challenges: The physician as team player\*

If you were to take physicians from the late 18th century—let us disturb William Withering, for example, physician to the General Hospital in Birmingham in the 1780s, or René Laennec from the Hôpital Necker in Paris—and invite them into a modern intensive care unit, what would they observe? After recovering from the shock of seeing complex 21st century technology attached to people in varying states of suspended animation, they might well take comfort from a familiar sight: a group of clinicians on the ward round moving slowly from bed to bed or between cubicles, listening to the history, performing a physical examination (1), inspecting the urine, labeling diseases, and prescribing treatments (2), activities which our two predecessors would have well understood.

If our visitors were to spend long time with us, however, they would note with surprise certain distinct differences about this ward round from those of their own experience. There would be constant interruptions and distractions; the persistent background noise of voices, alarms, and machinery; a huge volume of facts—especially numerical facts—would be exchanged or argued about; decisions would be negotiated, not mandated; members of the group would come and go, sometimes at the run; and over the course of 24 hours the members of the group would change several times, requiring repetitive rehearsal of the patients' histories. They might even find that some of the clinicians did not know each other's names. Within an ordered framework, there could be a sense of uncertainty and incipient chaos.

Thus, out of an apprenticeship-based, long-term, hierarchical, authoritarian, and paternalistic model of medical relationships emerges a different construct: the medical "team," a shifting federation of atomized individuals with varying competencies brought together for short periods with the specialist acting as manager-coordinator-educator. This change has come about through a mix of social, political, and professional pressures: the restriction of junior doctors' working hours (48 per week in Europe, 80 per week in the United States), abbreviated training times, shift working, removal of hospital accommodation and reduction in opportunities for social engagement in the workplace, blurring of professional borders and roles, and diminution of professional prestige and authority. At the same time, hospitals have tried to maximize efficiency by increasing throughput, reducing length of stay, and increasing day-case and outpatient work, with the effect of increasing acuity and complexity of case mix (3). This increases opportunities for miscommunication, error, and unreliable care in acutely ill patients, whose rapidly changing clinical state coincides with equally rapidly changing teams of clinicians.

The review of teamwork in the intensive care by Reader et al (4), in this issue of *Critical Care Medicine*, is therefore timely and important. They have examined the literature from an industrial psychology perspective. They conclude that teamwork—the way individuals interact together for a common purpose—has an important effect on patient outcomes, and they produce a descriptive framework based on inputs, processes, and outputs, in which processes are further categorized in terms of communication, leadership, coordination, and decision making. This model will make it easier to frame research questions and compare interventions in what has hitherto been an ill-defined area of quality improvement research.

The review by Reader et al also demonstrates that research from high-

reliability industries can illuminate care processes, which affect quality and safety in the intensive care unit. However, although the research undertaken in industrial environments provides examples of potentially effective interventions, it is important to recognize that health care and industry have as many differences as similarities. For example, the pilot and crew have more of an imperative than the surgeon and operating theater staff to avoid error (5): the worst result for the medical staff is that when the patient dies, there is no risk to their own physical safety as in an aviation setting. Conversely, the model of civilian aviation may oversimplify the situation of acute and emergency care, which may at times resemble a war zone. In the complex sociotechnical environment of the intensive care unit, we need research methodologies that can handle complexity (6) and adequately describe how social, cognitive, and behavioral factors coalesce to create a slick and responsive team, capable of reflecting "heedful interrelating" for verbal and nonverbal communication between team members. These facets of communication in highly complex environments are not easy to observe and measure. Researchers must, therefore, ensure that their research is fully "situated" in the unique context of the intensive care unit and acute care areas.

It is also important to define the borders of a subject as diffuse as teamwork. For example, the review does not include research on medical emergency teams (7), outreach care (8), or the impact of intensivist staffing on outcomes (9, 10), all of which involve the interaction between different specialties and the consequences of these interactions for acutely ill patients. A weakness of studies like these is that they do not define the content of the intervention, only the vehicle. Given the trend toward developing non-physician clinicians (11, 12), many of whose competencies are shared with physicians (13), and the consequent blurring

\*See also p. 1787.

Key Words: team; safety; communication; intensive care unit; critical care

The authors have not disclosed any potential conflicts of interest.

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DOI: 10.1097/CCM.0b013e3181a092ed



of roles, if we are to study teamwork we will have to define more clearly the active components of a team and its impact on processes of care (14, 15), as well as specify what we mean by “doctor,” “nurse,” “physician assistant” in terms of their competencies, roles, and contributions to the intended outcome.

Health systems have distinct cultural heritages, promulgated not only through the immersion of daily working within hospitals and clinics but also through explicit and tacit learning. The ways in which clinicians develop and execute their knowledge is usually partitioned by the boundaries of professional identities and status. This contributes to poor communication, staff being unable to “speak up,” for example (16), with widely differing perceptions of team spirit (17, 18), impacting the understanding of common goals of care as shown previously by Reader et al (19). Undergraduate and postgraduate medical curricula should include communication and behavior skills that teach clinicians how to raise difficult issues constructively and become self-reflective and more aware of the effects and consequences of suboptimal or dysfunctional social interaction. Shame and guilt about clinical errors need to be channeled into constructive learning and quality improvement for the whole team, and away from silence, depression (20), and burnout (21) for individuals. An “organization with a memory” (22) also needs to be an organization with a heart. If we can achieve this, we will not only have helped our patients but ourselves as well (23).

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