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Immune Unresponsiveness to Secondary Heterologous Bacterial Infection after Sepsis Induction Is TRAIL Dependent

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Sepsis is the leading cause of death in intensive care unit patients, taking >215,000 lives per year in the United States (1), and is classically defined as the body’s massive hyperinflammatory response to systemic infection. Not surprisingly, the majority of early studies on sepsis in humans and experimental animals focused on understanding the early hyperinflammatory phase, which is characterized by the increased production of a number of proinflammatory cytokines, including IFN-γ, IL-1β, IL-6, and TNF (2–4), and other proinflammatory mediators, such as HMGB1, MIF, TREM-1, and MRPS8/14 (5–9). Although the neutralization of IL-1β, TNF, or HMGB1 improved survival of septic mice and baboons (10–14), similar clinical attempts designed to neutralize IL-1β and/or TNF in septic patients failed to improve survival (15). Other anti-inflammatory–based therapeutic trials, such as antiendotoxin, steroids, anti–platelet-activating factor, also failed to produce a positive effect on the pathogenesis of sepsis (12, 15, 16).

The inability to decrease sepsis-induced mortality by dampening the hyperinflammation led many to question the idea that the early, massive inflammation in sepsis was the sole cause of mortality. It has since been shown that the initial hyperinflammation during sepsis is followed by a prolonged immunosuppressive phase that significantly increases morbidity and the chance of mortality caused by secondary (often nosocomial) infections (17). The increased susceptibility to secondary infection can be attributed to many things, but one hallmark of sepsis is the widespread apoptotic death of immune cells that provide protection. Adoptive transfer of apoptotic cells during sepsis increases mortality (18), whereas prevention of lymphocyte cell death using pharmaceutical agents (such as zVAD) or modified expression of pro- or antiapoptotic proteins (such as Bim−/− or Bcl-2 overexpressing mice) improves survival following sepsis (19–21). It is important to keep in mind that the sepsis-induced lymphopenia is not the only means by which the immune system is altered, because the function of the remaining immune cells is also significantly decreased. Using the well-defined cecal-ligation and puncture (CLP) mouse model (19, 20, 22), we recently found a clear relationship between sepsis-induced immune cell apoptosis and the ability to mount delayed-type hypersensitivity (DTH) (23). Moreover, this previous study showed that the lack of DTH responses during sepsis was dependent on TRAIL–mediated active immune suppression. Thus, the sepsis-induced hypoimmunity phase is likely the result of multiple mechanisms that together lead to the compromised overall health of the septic individual.

The present study continues our analysis of the role of TRAIL-mediated immune suppression during sepsis by modifying the CLP model to include a secondary (heterologous) bacterial infection with the Gram-positive, facultative intracellular bacterium Listeria
monocytogenes. Murine infection with L. monocytogenes is a widely used model for the analysis of cell-mediated immunity to an intracellular bacterial pathogen (24–26). Although both CD4 and CD8 T cells are activated in an Ag-specific manner after L. monocytogenes infection (27), CD8 T cells are the most effective mediators of anti-L. monocytogenes immunity (28–30). Thus, incorporation of secondary L. monocytogenes infection into the CLP model serves as a tool to specifically evaluate sepsis-induced alterations in CD8 T cell immunity to this heterologous bacterial infection. The decision to use L. monocytogenes as the secondary infectious agent in this model was further buttressed by the fact that CD8 T cell-mediated immunity against L. monocytogenes is TRAIL independent (31), so any changes in the CD8 T cell response to L. monocytogenes were predicted to be due to the sepsis-induced immune suppression. The data presented in this study further implicate the prominent role of TRAIL during sepsis-induced immune suppression and suggest the TRAIL/TRAIL receptor pathway as a therapeutic target for restoring T cell-mediated immunity in septic patients.

Materials and Methods

Mice
C57BL/6 (B6) and BALB/c mice were purchased from The National Cancer Institute. Trail−/− and Dr5−/− B6 mice were obtained from Agenon (Seattle, WA) (32) and Dr. W. El-Diery (University of Pennsylvania, Philadelphia, PA) (33), respectively. Trail−/−BALB/c mice were obtained from Dr. Thomas Sayers (National Cancer Institute, Frederick, MD). All mice were housed in the same facility for at least 4 wk, regardless of their source, before use, and animal procedures were performed according to National Institutes of Health guidelines and approved by the University of Iowa Institutional Animal Care and Use Committee. In all in vivo experiments, groups consisted of four or more animals, and experiments were repeated at least two times with similar results before reporting.

Cecal-ligation and puncture
Sepsis was induced by CLP, as described previously with slight modifications (34). Briefly, mice were anesthetized, and an abdominal incision was made to identify the cecum. The distal one-third of the cecum was ligated with 4–0 silk suture and punctured once using a 25-gauge needle. A small amount of cecal contents was extruded through the perforation. This level of injury was used to create a chronic septic state characterized by the loss of appetite and body weight, ruffled hair, shivering, diarrhea, and/or periorbital eductates and with <5% mortality. The peritoneum was closed with a continuous suture after returning the cecum into the abdomen. The skin was then glued shut with 3M Vetbond (3M Animal Care Products, St. Paul, MN), and 1 ml saline was injected i.p. for resuscitation. For sham-treated mice, all of the same steps were performed, except for ligation and puncture of the cecum. In some cases, the CLP-treated mice received 300 μg anti-TRAIL blocking mAb N2B2 (eBioscience) the day after surgery.

Bacterial preparation, infection, and titer
The attenuated L. monocytogenes strains DP-L1942 (attenuated L. monocytogenes) and OVA257-expressing (attenuated LM-OVA), which are ActA deficient, were grown, injected i.v., and quantified as previously described (35). Briefly, the bacteria were grown to log phase (OD600 0.06–0.1) in tryptic soy broth media. Bacteria were diluted to 107 CFU in 0.2 ml sterile saline (Hospira, Lake Forest, IL) and injected i.v. Serial dilutions of the same culture used for injections were plated and grown overnight at 37°C and colonies counted to independently determine the actual number of injected bacteria. Spleens and livers were harvested from infected mice on day 3 postinfection and homogenized in 0.2% IGEPAL buffer (Sigma-Aldrich, St. Louis, MO). The homogenized suspension was incubated at room temperature for 30 min, and serial dilutions were plated on streptomycin agar plates. Plates were cultured overnight at 37°C, and bacterial counts were determined.

Flow cytometry
For staining of surface markers, cells were incubated with fluorochrome-conjugated mAbs at 4°C for 30 min. The cells were then washed with FACS buffer (PBS containing 2% BCS and 0.2% NaN3) and then fixed with 2% paraformaldehyde in PBS. mAbs used for surface staining were FITC-CD11a (clone M174; BioLegend), PerCP- or allophycocyanin-CD8 (clone 53-6.7; Biolegend), PE- or PerCP/Cy5.5-CD3 (clone 17A2 for PE and clone 145C211 for PerCP/Cy5.5; BioLegend), PerCP- or allophycocyanin-CD4 (clone GK1.5; BioLegend), and PE-CD19 (clone 6D5; BioLegend).

For intracellular staining, 106 splenocytes were cultured at 37°C in the presence of OVA257 and brefeldin A (BD Biosciences) for 6 h. The cells were then washed and stained for surface markers as described above. The cells were then washed once with FACS buffer and fixed with 2% paraformaldehyde in PBS for 20 min at room temperature. After fixing, cells were washed and resuspended in saponin buffer for 30 min at 4°C. PE- or allophycocyanin-IFN-γ (clone XM1.2; BioLegend) was added to the cells and incubated for 20 min. Cells were washed and resuspended in FACS buffer and analyzed using a Becton Dickinson FACS Calibur (BD Biosciences).

Statistical analysis
Significant differences between two groups were evaluated using a two-tailed Student t test (p < 0.05). When more than two groups were compared, one-way ANOVA with a Dunnett’s posttest was used (p < 0.05). All statistical tests were performed using GraphPad Prism 5 for Mac OS X (GraphPad, La Jolla, CA).

Results

Reduced bacterial clearance and Ag-specific CD8 T cell responses in CLP mice
The initial hyperinflammatory phase of sepsis is quickly followed by a sustained immunosuppressive state in patients and clinically...
relevant animal models of sepsis (36, 37), and it has been suggested that this immune suppression is critical to the pathogenesis and subsequent morbidity and mortality in septic patients that acquire secondary infections. Thus, the current study examined the impact of sepsis-induced immunosuppression on the naive CD8 T cell response to the Gram-positive, facultative intracellular bacterium *L. monocytogenes*. Our version of the CLP model of sepsis involves ligation of the distal one third of the cecum and a single puncture to permit release of the cecal contents into the peritoneum. This level of injury reproducibly resulted in the development of many clinical symptoms of sepsis, including a significant loss of weight compared with sham-treated wild-type (WT) B6 mice (Fig. 1A). Importantly, these CLP-treated mice regained all of the lost weight by 7 d after surgery. When we added the secondary infection with attenuated LM-OVA, the sham or untreated WT B6 mice displayed minimal morbidity (as measured by weight loss; Fig. 1B) and were able to control the LM-OVA infection (Fig. 1C). In contrast, CLP-treated WT B6 mice that received the subsequent attenuated LM-OVA infection demonstrated sustained weight loss compared with sham-treated mice that was not regained, and these mice were less able to control the attenuated LM-OVA infection (50- and 1400-fold increase in bacterial titer in the spleen and liver, respectively, compared with untreated or sham-treated mice). These results suggest that the induction of sepsis compromises overall health such that even a secondary heterologous infection with an attenuated bacterial strain cannot be adequately controlled.

Sepsis induces widespread immune cell apoptosis, and, consistent with previous studies (23), we observed a significant reduction in the number of CD8 T cells in WT B6 mice 2 d post-CLP compared with sham-treated mice (Fig. 2A). Similar significant reductions were also seen for CD3+CD4+, CD11c+, and CD19+ cells (Supplemental Fig. 1). We then specifically examined the primary CD8 T cell response 7 d after attenuated LM-OVA infection and found a significant decrease in total number of Ag-experienced [based on CD11a upregulation and OVA257-specific CD8 T cells in the spleen was determined. The total number of Ag-experienced CD11a+CD8 T cells in spleen was determined directly ex vivo by flow cytometry, whereas the number of OVA257-specific CD8 T cells in the spleen was determined by intracellular cytokine staining for IFN-γ after in vitro restimulation with OVA257. C shows representative flow plots measuring the frequency of IFN-γ-producing CD8 T cells with and without OVA257 restimulation in vitro. *p < 0.05 (Student t test).

**FIGURE 2.** Ag-specific CD8 T cell responses to attenuated LM-OVA are reduced in CLP-treated WT B6 mice. WT B6 mice underwent sham or CLP surgery and were then infected with 10^7 CFU attenuated LM-OVA on day 2 postsurgery. A, Total CD8 T cell numbers in the spleen were determined on day 2 postsurgery (i.e., day of infection). B–D. On day 7 postinfection, the number of Ag-experienced (based on CD11a upregulation and OVA257-specific CD8 T cells in the spleen was determined. The total number of Ag-experienced CD11a+CD8 T cells in spleen was determined directly ex vivo by flow cytometry, whereas the number of OVA257-specific CD8 T cells in the spleen was determined by intracellular cytokine staining for IFN-γ after in vitro restimulation with OVA257. C shows representative flow plots measuring the frequency of IFN-γ-producing CD8 T cells with and without OVA257 restimulation in vitro. *p < 0.05 (Student t test).

**FIGURE 3.** CLP-treated Trail^-/- and Dr5^-/- mice have reduced morbidity after LM-OVA infection and retain the ability to control the LM-OVA infection. Sham or CLP-treated B6 WT, Trail^-/-, and Dr5^-/- mice were infected with 10^7 CFU attenuated LM-OVA 2 d postsurgery. A–C, Animal weight was measured daily beginning the day of surgery, and the percent weight loss with respect to the starting weight is shown. The dotted line represents the starting weight of the mice, normalized to 100%, and the arrow indicates the time of infection. Arrow points to the time mice were infected with attenuated LM-OVA. D–F, Sham or CLP surgery was performed on WT, Trail^-/-, and Dr5^-/- B6 mice, and the mice were infected with attenuated LM-OVA on day 2 postsurgery as in A. The bacterial titer in the spleen was then determined on day 3 postinfection. The dotted line represents the limit of detection (l.o.d.).
CLP-treated Trail−/− or Dr5−/− mice retain the ability to control a secondary L. monocytogenes infection and mount CD8 T cell immunity

We recently showed that sepsis-induced apoptosis resulted in the TRAIL-dependent suppression of DTH (23). To determine the extent to which the TRAIL-dependent immune suppression contributed to the sepsis-induced morbidity and inability to control the secondary L. monocytogenes infection, we first examined weight loss in sham- and CLP-treated WT, Trail−/−, and Dr5−/− B6 mice given a secondary infection with attenuated LM-OVA. As in Fig. 1, CLP-treated WT B6 mice showed sustained weight loss compared with sham-treated mice and were unable to control the attenuated LM-OVA infection as well as sham-treated mice (2200-fold increase in L. monocytogenes CFU in the spleen of CLP-treated mice; Fig. 3A, 3D). By comparison, the CLP-treated Trail−/− and Dr5−/− B6 mice were able to recover their body weight back to that of the sham-treated mice (Fig. 3B, 3C) postinfection. Moreover, the splenic LM-OVA burden in CLP-treated Trail−/− and Dr5−/− B6 mice was only slightly increased over sham-treated Trail−/− and Dr5−/− B6 mice (7- and 10-fold, respectively; Fig. 3E, 3F). Data presented in our previous report showed sepsis-induced lymphopenia was TRAIL independent (23), and similar findings were seen in the current study (Fig. 4A). However, there was no decrease in the total number of Ag-experienced CD11ahi (38) or OV A257-specific CD8 T cells in the spleens of CLP-treated Trail−/− and Dr5−/− mice compared with sham-treated mice (Fig. 4B, 4C). Thus, these data suggest that sepsis-induced alterations in morbidity, bacterial control, and CD8 T cell responses following a secondary infection with attenuated LM-OVA require an intact TRAIL/DR5 signaling pathway.

To investigate the extent to which the TRAIL-dependent decrease in ability to control a secondary bacterial infection after sepsis was mouse strain sensitive, we examined the ability of sham- and CLP-treated WT and Trail−/− BALB/c mice to control the secondary L. monocytogenes infection. There was a similar reduction in CD8 and CD4 T cells in CLP-treated WT and Trail−/− BALB/c mice compared with the sham-treated mice (Fig. 5A). Moreover, the CLP-treated WT BALB/c mice had increased splenic LM-OVA titers compared with sham-treated mice, but there were nearly identical LM-OVA titers in the spleens of the sham- and CLP-treated BALB/c Trail−/− CLP mice (Fig. 5B). Thus, these data suggest that there is no strain specificity for the observed TRAIL-dependent alterations in the ability to control a secondary bacterial infection during sepsis.

Therapeutic use of a blocking anti-TRAIL mAb in septic mice decreases weight loss, improves the control of secondary bacterial infection, and restores CD8 T cell responses

The data presented thus far suggest that an intact TRAIL/DR5 pathway correlates with increased morbidity, reduced control of a secondary bacterial infection, and decreased CD8 T cell response in CLP-treated mice. Because the genetic elimination of either TRAIL or DR5 in CLP-treated mice resulted in them behaving similar to sham-treated mice, we determined the extent to which the response of CLP-treated mice to secondary L. monocytogenes infection could be improved through the therapeutic disruption of the TRAIL/DR5 pathway. Thus, CLP-treated WT B6 mice were treated with a blocking anti-TRAIL mAb after the induction of sepsis, but before LM-OVA infection (Fig. 6). Using this approach, the anti-TRAIL mAb-treated mice lost more weight compared with the sham-treated mice, but not as much as the CLP-treated mice (Fig. 6A). However, there was a dramatic reduction in the splenic bacterial burden in the anti-TRAIL mAb-treated mice compared with the CLP-treated mice that did not receive the anti-TRAIL mAb (Fig. 6B). Similarly, the CD8 T cell response was restored in the anti-TRAIL mAb-treated mice (Fig. 6C, 6D). The magnitude of the Ag-specific CD8 T cell response in CLP-treated mice was similar to that seen in CLP-treated mice given an isotype control IgG (data not shown), which is consistent with other reports describing the in vivo use of the anti-TRAIL blocking mAb N2B2 and isotype control Ab (41–43). Together, these results suggest that mAb-mediated disruption of the TRAIL/DR5 pathway had a therapeutic benefit in the septic mice with regard to morbidity, bacterial control, and the CD8 T cell response after a secondary bacterial infection.

Discussion

Sepsis induces widespread immune cell apoptosis that is required for the subsequent immune suppression (23). Consequently, septic patients and animals become very susceptible to subsequent infection that can often result in death (44). In the current study, we used a modified version of the clinically relevant CLP sepsis
model to include a secondary heterologous bacterial infection to study the role of TRAIL in sepsis-induced alterations in bacterial clearance and CD8 T cell immunity. Our results provide further evidence to suggest that TRAIL is a key molecular component in the establishment of sepsis-induced immune suppression. In this particular case, sepsis induced TRAIL-dependent immune suppression that limited the host’s ability to control a secondary bacterial infection and reduced the bacterial Ag-specific CD8 T cell response. Our data also suggest that TRAIL may be a potential therapeutic target during sepsis, because neutralization of TRAIL in septic mice restored the ability of these mice to clear the bacterial infection and mount a normal bacterial Ag-specific CD8 T cell response.

The CLP mouse model is considered the gold standard of sepsis models because it reproduces the pathophysiology of polymicrobial sepsis and intra-abdominal peritonitis seen in humans (34). Early after CLP, mice show many of the symptoms of sepsis that are seen in septic humans, such as hypothermia, tachycardia, and proinflammatory cytokine production. Depending on the severity of the sepsis induced (and the experimental interest of the investigator), death may occur within the first 48 h, or the mice will survive and develop immune suppression. This flexibility in disease severity and ability to study acute or chronic sepsis is simultaneously one of the biggest advantages and disadvantages of the CLP model. We reproducibly have a >95% animal survival rate in CLP-treated mice, which is similar to that reported by others examining the chronic effects of sepsis on the responsiveness of the immune system. A number of reports have combined secondary infection with CLP, and it is clear that sepsis alters the immune response to the infectious pathogen in the preclinical models much like that seen in the clinical situation. For example, CLP-treated mice are highly susceptible to pulmonary Pseudomonas aeruginosa infection and demonstrate increased mortality compared with sham-treated mice (2, 45, 46). One weakness in using P. aeruginosa as the secondary infections agent is that specific T cell responses are not as easily followed as they are when L. monocytogenes is used. The vast amount of information gathered over recent years regarding the T cell response to L. monocytogenes is one reason why we chose to use this bacterial pathogen in our system. L. monocytogenes normally induces a robust CD8 T cell response (25), so the observation that CLP-treated WT mice had a reduced ability to control a secondary attenuated L. monocytogenes infection and reduced CD8 T cell responses compared with sham-treated WT mice or CLP-treated Trail−/− or Dr5−/− mice was very intriguing. Interestingly, a recent report by Delano et al. (46) showed that CLP mice were more resistant to virulent L. monocytogenes infection when compared with sham-treated mice, resulting in increased survival. These data are in direct contrast with the data we show in this report. Possible explanations for these differences, compared with our data, include (but are not limited to): 1) use of attenuated versus virulent L. monocytogenes; 2) day of infection (day 2 versus day 3 post-surgery); 3) infectious dose (107 CFU attenuated L. monocytogenes versus 108 CFU virulent L. monocytogenes); 4) day of assessment of bacterial titer (day 3 versus day 5 post-surgery); and 5) severity of sepsis induction (<5% versus ~15% mortality). Attenuated and virulent L. monocytogenes elicits very distinct immune responses (47). Attenuated L. monocytogenes is replication
deficient such that it enters the cell and does not propagate; however, virulent *L. monocytogenes* replicates and propagates from one cell to another (48). The kinetics of bacterial burden and CD8 T cell responses elicited by attenuated and virulent *L. monocytogenes* are also considerably different (47); note that *L. monocytogenes*-specific CD8 T cells were not directly measured in the Delano et al. (46) study. Given these multiple differences, we are presently unable to reconcile the differences between the data in Delano et al. (46) and our data. Regardless, our study adds the important observation that bacterial Ag-specific CD8 T cell responses are suppressed in septic mice in a TRAIL-dependent manner.

We previously showed that sepsis-induced apoptosis led to suppression of DTH through a TRAIL-dependent mechanism (23), but the similar importance of TRAIL-dependent suppression of CD8 T cell responses during sepsis to a secondary heterologous bacterial infection had not been demonstrated until now. In our previous study, the suppression of DTH in septic mice resulted from the activity of TRAIL-expressing CD8 regulatory T (Treg) cells (23). We have had a long-standing interest in the tolerogenic nature of apoptotic cells, especially the ability of apoptotic cells to influence the induction of T cell-mediated immunity, and the induction of Ag-specific unresponsiveness (or tolerance) in an adult can be achieved experimentally through a variety of approaches (49–53). The use of several different experimental tolerance models has helped us demonstrate that one way the immune system reacts to the plethora of self-Ag derived from a large wave (49–53). The use of several different experimental tolerance cells (23). We have had a long-standing interest in the tolerogenic health of the immune system in regard to the ability to control increased survival after sepsis induction in mice (55–57). Although susceptible to secondary infections. Recent studies have shown that therapeutic disruption of the TRAIL/DR5 pathway would have any benefits during sepsis. We hypothesized the anti-TRAIL of TRAIL in the sepsis-induced unresponsiveness to a secondary heterologous infection and the data showing the restored ability to control the LM-OVA susceptibility to secondary infections. The potential impact of the anti-TRAIL mAb on immune system responses to a secondary heterologous bacterial infection after sepsis, we cannot exclude the possibility that the anti-TRAIL mAb was also inhibiting the death of cells outside of the immune system. Thus, anti-TRAIL mAb blockade of intestinal epithelial cell apoptosis may have contributed to the increased weight gain in this group of CLP-treated mice. Further investigation, which would also include analyses of CLP-treated Trail−/− and Dr5−/− mice, is needed to determine the likelihood of this. Regardless, the data presented in this study suggest TRAIL could be a potential therapeutic target in sepsis to help decrease the susceptibility to potentially life-threatening secondary heterologous infections.

**Disclosures**

The authors have no financial conflicts of interest.

**References**

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