Systemic Dissemination of *Clostridium difficile* Toxins A and B Is Associated With Severe, Fatal Disease in Animal Models

Jennifer Steele,1,a Kevin Chen,1,a Xingmin Sun,1 Yongrong Zhang,1,2 Haiying Wang,1,3 Saul Tzipori,1 and Hanping Feng1

1Tufts Cummings School of Veterinary Medicine, North Grafton, Massachusetts; 2School of Bioengineering, East China University of Science and Technology, Shanghai; and 3School of Bioscience and Biotechnology, South China University of Technology, Guangzhou

(See the editorial commentary by Johnson, on pages 353–4.)

**Background.** *Clostridium difficile* infection (CDI) can cause a wide range of disease, from mild diarrhea to fulminant systemic disease. The incidence of systemic CDI with fatal consequence has increased rapidly in recent years.

**Methods.** Using an ultrasensitive cytotoxicity assay, we measured *C. difficile* toxin A (TcdA) and *C. difficile* toxin B (TcdB) in sera and body fluids of piglets and mice exposed to *C. difficile* to investigate the relationship between the presence of toxins in body fluids and systemic manifestations of CDI.

**Results.** We found that both TcdA and TcdB disseminate systemically, with toxins present in the sera and body fluids of infected animals, and toxemia is significantly correlated with the development of systemic CDI. The systemic administration of neutralizing antibodies against both toxins blocked the development of systemic disease in mice. We measured cytokine concentrations in the sera of mice and piglets with systemic and nonsystemic CDI and found that proinflammatory mediators were considerably elevated in animals with systemic CDI.

**Conclusion.** Our study demonstrates the existence of a strong correlation between toxemia and the occurrence of systemic disease, supporting the hypothesis that systemic CDI is most likely due to the toxicity of TcdA and TcdB and the induction of proinflammatory cytokines by the toxins.

*Clostridium difficile* is an important nosocomial pathogen and the most commonly diagnosed cause of antibiotic-associated diarrhea and colitis [1]. *C. difficile* infection (CDI) can cause a wide range of disease, from mild diarrhea to fulminant systemic disease. Since 2000, outbreaks of CDI with greater morbidity and mortality have occurred in many developed countries [2]. CDI is thought to be caused by 2 glucosylating toxins secreted by the bacterium *C. difficile* toxin A (TcdA) and toxin B (TcdB) because strains lacking the toxins are avirulent. Recent studies show that both toxins can cause disease in hamsters [3], but their relative roles in the disease are unclear. *C. difficile* infection commonly results in severe gastrointestinal pathology, including pseudomembranous colitis and toxic megacolon, but may also result in a variety of observed systemic complications. Reported systemic complications in patients with CDI include ascites [4, 5], pleural effusion [6, 7], cardiopulmonary arrest [8, 9], hepatic abscess [10], abdominal compartment syndrome [11], acute respiratory distress syndrome [12], multiple organ dysfunction syndrome [13], and renal failure [14]. Although bacteremia has been identified as a cause of systemic manifestations of CDI in some cases, it is not always present, and other mechanisms by which *C. difficile* causes these systemic effects are not well understood. The 2 glucosylating toxins produced by the bacteria are likely involved, and a greater understanding of the systemic effects of *C. difficile* infection and why
they occur in some patients, but not others, is important because these effects are often life-threatening in nature.

Animals infected by *C. difficile* often experience systemic manifestations of disease, in addition to typical gastrointestinal signs. Naturally infected piglets sometimes develop respiratory distress and hydrothorax, as well as ascites [15, 16]. Administration of TcdB to zebrafish causes cardiac damage [17], a finding that places more importance on understanding the systemic role of these toxins in mammals. An obstacle to investigations of the relationship between systemic disease and toxemia in CDI has been the lack of a sensitive detection assay for toxins in blood or tissue fluid. We recently reported the development of an ultrasensitive immunocytotoxicity (ICT) assay, which can detect toxin concentrations as low as 1 pg/mL in samples of the serum and body fluids of piglets with CDI [18, 19]. Using this assay, we investigated the relationship between the occurrence of systemic disease and toxemia in gnotobiotic piglet and mouse models of CDI.

**MATERIALS AND METHODS**

**Animals and Inoculation**

Gnotobiotic piglets were delivered via cesarean section into a sterile isolator and maintained in groups of 2-4 inside isolators for the duration of the study, as previously described [18]. We orally inoculated a total of 79 piglets, ranging in age from 2 days to 7 weeks, with $10^3$–$10^9$ *C. difficile* spores using the NAP1/027/BI strain UK6 [20]. For 1 group of 8 piglets, we collected blood daily after inoculation, and for all piglets, we collected blood at the time of euthanasia. We performed a necropsy on each piglet following euthanasia or death, assessed gross gastrointestinal and systemic lesions, and collected body fluids, including pleural and abdominal effusion, if present.

We maintained 6- to 9-week-old C57BL/6 mice (Jackson Laboratory) in pathogen-free facilities in groups of 5. We orally inoculated mice with $10^6$ *C. difficile* spores using the NAP1/027/BI strain UK1 [20] after a series of antibiotic treatments, as previously described by Chen et al [21]. We collected blood daily after inoculation to assess the progression to toxemia.

We collected tissues, including small intestine, large intestine, mesenteric lymph nodes, liver, kidney, spleen, lung, and heart, from piglets and mice at the time of necropsy for histopathologic examination. We handled and cared for all animals according to Institutional Animal Care and Use Committee guidelines.

All piglets and mice were monitored for the development of clinical signs of CDI several times daily after inoculation with *C. difficile*. Based on the assessment of signs, we classified animals as developing either nonsystemic CDI or systemic CDI. For both piglets and mice, nonsystemic CDI refers to the presence of localized gastrointestinal illness only, evidenced by development of diarrhea, without any systemic clinical signs. Systemic CDI refers to the development of systemic signs of illness in addition to diarrhea. In piglets, clinically observable systemic signs include lethargy, depression, weakness, and dehydration, progressing to complete anorexia and dyspnea. In mice, systemic CDI is evidenced by depression, unkempt haircoat, hunched posture, weakness, and dehydration, progressing to anorexia and abdominal distension. Both piglets and mice were euthanized either when systemic clinical signs were progressively worse over the course of several daily checkpoints or when the animal was found moribund or dyspneic. Animals that never developed any systemic signs of illness were euthanized at a predetermined experimental endpoint, generally 7–10 days after inoculation.

**Immunocytotoxicity Assay**

To determine the presence of toxins in serum samples, we used the ultrasensitive ICT assay using both the real-time cell electronic sensing system (RT-CES, Roche Applied Science) as previously described by our laboratory [18], as well as traditional methods of observing cell rounding. In brief, mRG1-1 cells, which express the FcγRI-α chain, were added to wells of microelectrode-imbedded microplates (E plates) or standard 96-well plates. We added the A1H3 mouse anti-TcdA monoclonal antibody generated by our laboratory [22] to wells with cells. A1H3 increases the sensitivity of cells to TcdA, allowing detection of low concentrations of TcdA in samples to the 1–10 pg/mL range [18]. TcdB is more cytotoxic than TcdA and can be detected ~ 10 pg/mL without an enhancing antibody. Mouse and piglet serum samples were added to cells at a 1:10 final dilution. We used goat anti-TcdA and anti-TcdB serum (TechLab) in the assays to neutralize the activity of both toxins. Rabbit anti-TcdA and alpaca anti-TcdB sera were used for neutralizing TcdA and TcdB, respectively. Recombinant TcdA and TcdB [23] were used as positive controls and standards. After addition of samples, toxins, and antibodies, we connected E plates to the RT-CES and collected cytotoxicity data that was measured via cell index for 12–24 hours. We used a phase-contrast microscope to assess the percentage of cell rounding on standard plates hourly for ~8 hours and then after overnight incubation.

**Rac1 Glucosylation**

We evaluated piglet and mouse serum samples, which were toxin positive in the ICT assay, for glucosyltransferase activity by assessing glucosylation of the Rho GTPase Rac as described previously [23]. In brief, mRG1-1 cells were seeded in 24-well plates and incubated until the cells reached confluence. Cells and A1H3 only were used as negative controls, and 10 ng/mL of TcdA with A1H3 was used as a positive control. We used the polysera described previously to neutralize either both toxins or each toxin separately. We harvested cells when toxin control wells reached 100% cell rounding, and we performed immunoblotting. We used antibodies that specifically recognize the nonglucosylated form of Rac1 (clone 102, BD Bioscience) and anti-β-actin (clone AC-40, Sigma) as the primary antibodies.
and horseradish peroxidase–conjugated anti-mouse immunoglobulin G (Amersham Biosciences) as the second antibody.

**In Vivo Neutralizing Antibody Treatment**

We orally inoculated C57BL/6 mice with 10^6 C. difficile UK1 spores after antibiotic treatment, as previously described by Chen et al [21]. We treated mice 4 hours after inoculation with C. difficile, with alpaca polyclonal antisera against TcdA and TcdB. The antiserum for each toxin were generated separately, and mice were dosed based on the neutralizing activity of the sera with a single intraperitoneal injection of the mixed anti-TcdA and anti-TcdB sera. We injected control mice with an equal amount of nonimmune sera. All mice were monitored for development of clinical signs of CDI and progression to systemic CDI and mice that became moribund were euthanized.

**Cytokine Measurement**

In piglet serum samples, cytokine concentration was determined for interleukin 1β (IL-1β), interleukin 4 (IL-4), interleukin 6 (IL-6), interleukin 12 (IL-12), tumor necrosis factor α (TNF-α), transforming growth factor β, and interferon γ (IFN-γ) using porcine cytokine quantification kits (Invitrogen and R&D). In total, 69 piglet serum samples were analyzed, including 29 from animals with systemic CDI and 40 from animals with nonsystemic CDI. For mice, a total of 19 serum samples from 69 mice were analyzed, including 7 from animals with systemic CDI, 7 from animals with nonsystemic CDI, and 5 from animals treated with alpaca polyclonal antisera against TcdA and TcdB described previously. The cytokine concentration for IFN-γ, IL-12, IL-10, TNF-α, IL-6, KC (CXCL1), and IL-1β was determined, following the manufacturer’s directions, with murine cytokine quantification kits (Invitrogen and R&D).

**Statistical Analysis**

We compared categorical variables, such as presence of toxemia and systemic CDI, for each group using the Fisher exact test. We used the t test to analyze continuous variables, such as cytokine measurements. All analyses were performed using SPSS statistical software version 16.0.

**RESULTS**

**Both TcdA and TcdB Are Liberated Into Serum and Body Fluids**

Using the ICT assay [18], we found that 12 of 43 piglet serum samples with CDI (27.9%) and 23 of 69 mouse serum samples with CDI (33.3%) were positive for toxin. Additionally, 6 of 13 pleural fluid samples (46.2%) and 11 of 14 ascites samples (78.6%) collected from piglets and 5 of 5 pleural fluid samples (100%) and 2 of 2 ascites samples (100%) collected from mice were positive for toxin (Table 1). The concentrations of TcdA or TcdB detected in the sera and fluids ranged from 1 pg/mL to 10 ng/mL, and were determined either by the cell index curve or percentage of cell rounding. Figure 1 depicts the appearance of cells in the ICT assay using a standard 96-well plate. Serum samples collected from a piglet and mouse with systemic CDI caused typical cell rounding (Figure 1B and 1E), similar to that of TcdA as a control (Figure 1A). Preinoculation serum or the toxemic samples mixed with neutralizing anti-TcdA and TcdB antibodies had no ability to cause cell rounding (Figure 1C, D, F), indicating that the cell rounding is indeed caused by the toxins.

To determine which of the 2 toxins were present in the samples, neutralizing antibodies against individual toxins were added to sample wells in the ICT assay. Figure 2A illustrates representative data from the assay with serum from 1 piglet with systemic CDI. Cells in the control well displayed normal attachment and growth (Figure 2A) with a rapid rise in the cell index, and cells with 10 ng/mL of TcdA displayed retarded growth with a minimal increase in cell index (Figure 2A). The cell index of the well containing the piglet serum remained at baseline levels throughout the 24-hour period (Figure 2A). Addition of anti-TcdA and anti-TcdB antibodies to the serum
sample fully neutralized the toxicity, indicating that it is the presence of the toxins in the serum causing cytotoxicity (Figure 2A). Addition of only anti-TcdB antibody did not improve the cell index (Figure 2A), and addition of only anti-TcdA antibody resulted in a marginal increase in the cell index (Figure 2A), indicating that both toxins were present in the serum in cytotoxic concentrations. Similarly, in Figure 2B, sera from 1 representative mouse with systemic CDI and 1 mouse with nonsystemic CDI were evaluated for the presence of toxin using the ICT assay. Only serum from the mouse with systemic CDI caused a retarded growth curve (Figure 2B), and addition of antibodies against TcdA and TcdB fully neutralized the toxic effect of the serum (Figure 2B).

To further verify the presence of clostridial glucosylating toxins, we evaluated the glucosylation of Rac1 in cells in the presence of the serum samples. When TcdA or TcdB is present in a sample, the Rac1 in mRG1-1 cells incubated in wells with the sample will be glucosylated by the toxin and thus will be undetectable by the anti-Rac1 antibody, which recognizes only the nonglucosylated protein [23]. In control wells containing only cells or cells plus neutralizing antibodies alone, Rac1 was readily detectable (Figure 3A, lane 1, 2), but when toxin was incubated with the cells as a positive control, Rac1 was not detected (Figure 3A, lane 3, 4). Mouse presera and sera collected 1 day postinoculation were negative for toxin based on the presence of Rac1, but samples collected from mice 2 days postinoculation, with development of systemic CDI, were positive (Figure 3A, lanes 5–9). When anti-TcdA and TcdB antibodies were added to the samples and incubated with cells, toxin activity was completely neutralized, and Rac1 levels were similar to that of controls (Figure 3A, lane 10). When only anti-
Toxemia Is Associated With the Development of Systemic Manifestations of CDI

In both the mouse and piglet models of CDI, the severity of disease ranged from mild, localized gastrointestinal illness to severe, systemic, and often fatal disease. To examine whether the presence of toxin in the systemic circulation is associated with systemic manifestations of CDI, we divided the animals into 2 groups: systemic CDI and nonsystemic CDI. Of the 43 piglet serum samples tested, 18 were from animals with systemic CDI and 25 were from animals with nonsystemic CDI (Table 1). Twelve of 18 (66.6%) serum samples from piglets with systemic CDI had detectable levels of toxin. In piglets with nonsystemic CDI, however, toxin was not present in the serum at detectable levels (Table 1). The association of toxemia in piglets with the occurrence of systemic CDI is highly significant (P < .0001). Similarly, in mice, 27 of 61 serum samples tested were from animals with systemic CDI and 34 were from animals with nonsystemic CDI. Of 27 animals with systemic CDI, 23 (85.2%) had detectable levels of toxin, but none of the animals with nonsystemic CDI had detectable toxin levels (Table 1). The association of toxemia with systemic CDI in mice is also highly significant (P < .001). We found no evidence of bacteremia or bacteria in the lungs of piglets or mice with systemic CDI ([19] and data not shown). As evidenced by the results from a representative piglet and mice (Figure 3), toxemia coincides with the development of systemic CDI, and neither TcdA nor TcdB was detectable in the sera of animals before the onset of systemic clinical signs of CDI.

Systemic Neutralizing Antibodies Block Toxemia and Systemic Disease

Because both TcdA and TcdB are liberated into circulation in animals with toxemia, we examined whether neutralizing the 2 toxins would eliminate systemic manifestations of disease and thus reduce severity. Mice treated with systemic neutralizing antibodies against TcdA and TcdB (antitoxins) were completely protected from the development of toxemia, systemic CDI, and death. Of the nonimmune serum-treated control mice, 40% developed fatal systemic CDI (Figure 5A) with systemic clinical signs of CDI such as weight loss, anorexia, dehydration, lethargy, dyspnea, and poor body condition. The antitoxin-treated mice, on average, lost <6% of their body weight and quickly recovered body weight compared with control mice (Figure 5B).
cytokine production. For piglet sera, we divided samples into systemic-CDI (n = 29) or nonsystemic-CDI (n = 40) groups, and the mean cytokine concentrations were compared for significant differences between groups with the *t* test for each cytokine. In piglet sera, IL-1β, IL-4, and IL-6 concentrations were significantly different between groups, with *P* values of .03, .04, and .001, respectively (Figure 6A). Piglets developing systemic CDI had significantly greater levels of the inflammatory cytokines IL-1β and IL-6, but significantly lower levels of IL-4, an anti-inflammatory cytokine (Figure 6A). Likewise, mice with systemic CDI also had elevations in the concentrations of several cytokines measured. CXCL1 (KC) and IL-6 were most elevated in mice with systemic CDI, with a trend for elevations also noted for IFN-γ, IL-1β, IL-10, and IL-12 (Figure 6B).

**DISCUSSION**

*C. difficile* is an important nosocomial pathogen in developed countries [29]. The increased morbidity and mortality associated with CDI, which began in the early 2000s with the emergence of NAP1/027/BI strains, has reached epidemic proportions [29]. In the United States, some 500 000 cases of CDI, with approximately 15 000–20 000 deaths, are reported annually [29, 30]. Systemic complications of CDI are often life threatening, and although large-scale studies on systemic CDI in humans are unavailable, there exists a wealth of case reports describing a variety of extracolonic complications [31–33]. Although suspected, a linkage between systemic CDI and toxemia has not yet been confirmed with direct evidence in humans. We conducted experiments in mice and piglets, aiming to define the features that distinguish systemic CDI from nonsystemic CDI.

We have previously reported the detection of toxins in sera and body fluids of systemically ill animals with CDI [18, 19], which suggested a potential association with systemic involvement. In contrast, in piglets and mice with CDI that exhibit diarrheal symptoms with no apparent systemic manifestations, toxins are undetectable in the sera or body fluids.
Recent reports describing the relative importance of TcdA and TcdB in animal models of CDI have suggested that TcdB was likely linked with systemic disease [17, 34]. Our findings show that both toxins are present in the serum at cytotoxic concentrations, observations consistent with a recent report that TcdA and TcdB are implicated in the pathogenesis of systemic CDI [3].

Our work clearly shows that TcdA and TcdB reach systemic circulation following colonization of the intestinal tract by the bacteria, and when they do, systemic disease ensues. Systemic abnormalities in these animals included pleural effusion, ascites, and cardiopulmonary lesions, all observations that have been found, which included cranial-ventral consolidation of the lung lobes, pleural effusion, and diffuse petechiation of the lung lobes, which may be due to a direct toxic effect.

Because TcdA and TcdB are both proinflammatory [35], we believe that toxemia may have induced a profound inflammatory reaction that contributed to the severity of disease. To investigate one aspect of the inflammatory response, we examined pig and mouse serum samples for the presence of various pro- and anti-inflammatory cytokines. We found that in both pigs and mice, proinflammatory cytokine levels were elevated in animals with systemic CDI compared with those in animals with nonsystemic CDI. In both species, significant elevations were found for levels of IL-1β and IL-6, both of which are proinflammatory mediators. Interestingly, a significant elevation in the anti-inflammatory cytokine IL-4 was observed in piglets with no apparent systemic manifestations of disease. These differences between the 2 groups warrant further investigation, as the proinflammatory response in CDI may contribute to the development of systemic disease. Our findings in both animal-infection models of CDI support the importance of inflammatory mediators in the C. difficile immune response and the potential implications for inflammation-induced tissue damage systemically.

Our findings have implications for understanding the pathogenesis of CDI, the development of novel vaccines, and immunotherapy. Knowing that the toxins can be liberated into circulation and subsequently can cause systemic inflammation and tissue damage illustrates the need to not only target the bacteria with effective antibiotics, but also to prevent or reduce the impact of toxin present in circulation in severe cases. Development of vaccines and monoclonal and polyclonal antibodies directed against the 2 toxins will help prevent the development of systemic CDI. Our observations warrant a close examination of toxemia in humans with systemic CDI, in addition to treating the infection within the gastrointestinal tract.

Notes

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References


Figure 6. Piglet and mouse serum cytokine concentrations. A, Serum from piglets (n = 89) was grouped according to disease severity, as systemic (black bar, n = 29) or nonsystemic (open bar, n = 40) Clostridium difficile infection (CDI); B, Serum samples from 7 mice with systemic CDI (black bar), 1 mouse with nonsystemic CDI (open bar), or 5 mice treated with anti–Clostridium difficile toxin A and anti–C. difficile toxin B polysera (anti AB, gray bar) were run separately in cytokine assays. Columns represent the mean cytokine concentration with standard error bars for each cytokine. Statistical differences were determined by the t test. *P < .05; **P < .01. Abbreviations: IFN-γ, interferon-γ; IL, interleukin; TGF-β, transforming growth factor β; TNF-α, tumor necrosis factor α.