CLOSTRIDIUM DIFFICILE BINARY TOXIN (CDT) IN ANTIBIOTIC ASSOCIATED DIARRHEA Adam L. Stevens¹; Matthew W. Lyerly¹; Megan F. Hiltonsmith¹; Bradley G. Stiles^{2,3}; Tracy D. Wilkins¹; Robert J. Carman^{1*} ¹TechLab, Inc., Blacksburg, VA 24060, USA; ²Toxinology Division, Department of Immunology and Molecular Biology, United States Army Medical Research Institute of Infectious Diseases, Fort Detrick, MD 21702, USA; ³Wilson College, 1015 Philadelphia Avenue, Chambersburg, PA 17201, USA

Abstract

We used an EIA, calibrated with purified recombinant antigen, to quantify cdtB which acts as the binding component of Clostridium difficile binary toxin (CDT). This work was done with feces from 80 confirmed human cases of C. difficile diarrhea and culture fluids of isolates recovered from 53 of the same fecal samples. By using PCR, we ribotyped isolates and probed each for the three genes (cdtR, cdtA, cdtB) that make up the CDT locus. CDT is best known as a toxin associated, though not exclusively so, with the ribotype 027 isolates that are central to the recent worldwide rise in the incidence of C. difficile diarrhea. We found CDT(+) isolates in 50/80 (63%) fecal samples; cdtB was present in only 38% of that 50. In contrast, 90% of all CDT(+) isolates we recovered made cdtB protein in broth culture. Ribotype 027 isolates were 86% of all CDT(+) isolates. Six other ribotypes accounted for the 7 non-027, CDT(+) isolates; 6 of the 7 made cdtB in vitro. 25/80 (31%) isolates carried the ghost CDT locus, i.e. an intact cdtR but only fragments of cdtA and cdtB. 5/80 (6%) lacked all or any part of the CDT locus. No CDT- isolate was recovered from an EIA negative stool sample. Levels of cdtB in the fecal samples were on average nearly twenty-fold greater than the corresponding isolates achieved in vitro. We saw, however, no direct relationship between toxin concentrations in feces and in vitro cultures. Over half of the EIA positive samples contained >100 ng cdtB/mL, a level that reached in stools matches other closely related clostridial binary toxin diarrheas. This information suggests that when there is 100 ng/mL or more of cdtB present, the level of CDT may be sufficient to cause diarrheic symptoms, possibly complementing the known enteric activities of Toxins A and B.

Background

Clostridium difficile is an important nosocomial pathogen associated with about 20% of all antibiotic-associated diarrheas. C. difficile is also the most common causative agent of pseudomembranous colitis, a potentially life threatening form of disease in the colon. The main virulence factors of C. difficile are two toxins, A and B, which are large single-chain proteins 308 and 270 kDa, respectively. It has been shown that in addition to Toxins A and B several C. difficile strains produce a third, binary toxin, CDT. CDT is composed of two independent, unlinked polypeptides, CDTa and CDTb. CDTa is an enzyme, an actin-specific ADP-ribosyl transferase. Though enzymatically active by itself, CDTa is only toxic when internalized via a CDTb pentamer docked on the surface of host enterocytes. CDTb forms the oligomer even in the absence of CDTa. Only together do CDTa and CDTb have cytotoxic and enterotoxic activity.

At genetic, immunologic and toxicologic levels CDT is very closely related to two other clostridial binary toxins, iota toxin of *C. perfringens* type E, and iota-like toxin of C. spiroforme. Expression in C. perfringens and C. spiroforme is 10 or more-fold greater than in *C. difficile*, suggesting a different regulation for *C.* difficile. In fact, cdtR, a gene absent from the plasmid in C. perfringens that carries the iota toxin genes or from the iota-like gene locus on the chromosome of C. spiroforme, regulates CDT production by C. difficile. Production of CDT by C. difficile is positively regulated by CDTr, encoded by cdtR.

We screened samples from patients with *C. difficile* diarrhea for the presence of CDT using an ELISA. We isolated C. difficile from these samples and screened the isolates for the CDT gene locus using PCR and for the in vitro production of CDTb.

Aims

- To assay stool from patients diagnosed with C. difficile diarrhea for the presence of CDTb, the binding component of *C. difficile* binary toxin.
- To recover C. difficile from patient stool and to assay these isolates for the three genes, *cdtA*, *cdtB* and *cdtR*, that make up the CDT locus.
- To assay recovered isolates for the *in vitro* production of CDTb.

Materials and methods

Patient feces : Already existing, anonymous and unlinked fecal sa patients diagnosed by the presence of GDH and toxins A or B or bo C. difficile diarrhea were used.

Bacterial isolates: Feces were plated onto Cycloserine cefoxitin fro a selective and differential medium for the selective growth of C. dif inoculated plates were incubated anaerobically at 37 C for 48 to 72 colonies were subcultured and stored in chopped meat broth until re Bacterial DNA: DNA was extracted using the QIAGEN DNA Mini k summary, 24 h brain heart infusion broth cultures were centrifuged pellets digested with a lysozyme buffer. After incubation, proteinase additional lysis buffer was added, followed by a second incubation. was centrifuged and DNA was washed and collected by alcohol pre QIAamp spin columns. Collected DNA was stored at -20°C until rea analysis.

PCR: See table below.

ELISA for CDTb: A prototype ELISA using monoclonal and monos polyclonal antibodies against CDTb was used. Feces were assave added dilution being needed. In vitro cultures were concentrated 1 before testing. This was necessary since, as we and others have p noted, levels of detectable CDT in broth medium are 10 or 20 fold lo the equivalent levels in levels in C. perfringens Type E and C. spirot the event, about 10 or more fold lower than the levels found in fecal from which isolates were recovered. Concentrations of CDTb were by comparison with a curve prepared with rCDTb standards.

PCR conditions and primers for detection of cdt locu Forward Primer | Reverse Primer (5' The Gene amplified (5' to 3') to 3') AGTCTATAATTAT AGTTGCGCAGCAT cdtR TGTTAAATAATTC GCATCTAAATCTG (regulatory gene) TTC GTAAAG cdtA TGAACCTGGAA AGGATTATTTACT (ADP ribosyl AAGTGGATG GGACCATTTG transferase) cdtB GATTCACAGCTA TATAGTCTGACCA (binding

ATGTAACTA ACTATTTC component) Primers for the cdtR gene were taken from Carter et al (2007). Prim cdtA gene were taken from Stubbs et al. (2000). Primers for cdtB v generated from the complete sequence using the Primer3 web-base

References

selection program.

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Stubbs S, Rupnik M, Gibert M, et al. 2000. FEMS Microbiol Lett 186: 307-12.

| CONTRACT SHE | | Table 1. CDTb and <i>cdtb</i> in <i>C. difficile</i> diarrhea | | | | | | Results |
|--|---|---|-----------|--------|---------|------------|---------------|------------|
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| ndividual | | 4 | _ | _ | _ | _ | _ | Sampl |
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| cles | Fig. 1. Agarose gel analysis of PCR | 35 | _/+ | + | + | + | + | have b |
| lo sec | amplicons. Lane 1 is an isolate negative for | 36 | _/+ | + | + | + | + | possib |
| 1 min | <i>cdtR</i> . Lanes 2 - 5 represent positive | 37 | + | + | + | + | + | perfrin |
| 1 min | isolates. Lane 6 is a negative control | 38 | + | + | + | + | + | |
| cycles | isolates and lanes 7 and 8 are positive | 39 | + | + | + | + | + | Isolate |
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fecal samples from patients diagnosed with *C. difficile* diarrhea or the presence of CDTb using a prototype immunoassay.

7 (13% of 53) were all negative in the CDT assay. The recovered ed all three genes of the CDT locus. None made CDT in vitro.

21 (26%) were all negative in the CDT assay. The recovered ed cdtR but were negative in the PCR assay for cdtA and cdtB. CDT in vitro.

o 34 (26%) were all negative in the CDT assay despite all the plates carrying the entire CDT locus and expressing CDTb in vitro. CDTb however, required the concentration of culture fluids by 10 or

ssay samples 35 and 36 (4%) gave positive elevated signals well ckground seen with true negatives. Their isolates carried the entire They made CDTb in vitro but once more it was only detected after

is 37 to 53 (30%) had CDTb in them, 5 of them at levels over 1 μ g/ shown). Their isolates carried the entire CDT locus. They made but again it was only detected after concentration.

nunoassay detected CDTb in about 2/3rds of the samples that hight be expected to contain CDTb. Its minimum level of detection her improved with refinement, though it may be as much a question n levels or prior metronidazole or vancomycin therapy as s in the assay. Even so, CDTb concentrations in at least 5 samples ame order as iotaB in an enterotoxic dose of purified iota toxin ingens Type E (Carman et al., 1986).

isolates carrying the full CDT locus produced CDT, though at only els compared with other binary toxin producers. Low levels of CDTb eported before (Popoff et al., 1988, Geric et al., 2006) and it may ect two or more regulatory mechanisms among C. difficile, C. ype E and C. spiroforme.

ying the 5' end of *cdtA* and the 3' end of *cdtB* but not the central quence have been known for some time (Chang & Song, 2001). d to carry ghost genes. All known examples have recently been o carry cdtR (Carter et al., 2007). In this study these are by the isolates from samples 8 to 21. They predictably did not Tb; none of the corresponding fecals was positive in the

for CDT in *C. difficile* disease caused by isolates producing neither but only CDT has been questioned (Geric et al., 2006), though levels of CDTb we saw may warrant reconsideration.

nuing our work by improving the immunoassay. We are also ne complementation of toxin A and B by the addition of CDT (of ern with the epidemic Ribotype 027 isolates that produce all three xins) and collecting data on the relationship between CDTb levels ee of intestinal inflammation indicated by lactoferrin levels.

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