

Iota Toxin, S Toxin and CDT: Members of the Same Class of Clostridial Binary Toxins in Feces of Humans and Other Animals

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Abstract: Some strains of *Clostridium perfringens*, *Clostridium spiroforme* and *Clostridium difficile* produce binary toxins known respectively as iota toxin, S toxin and CDT. Each toxin consists of two unlinked polypeptides (e.g. CDTa and CDTb) that only together have biological activity. Taking an historical perspective, we review the development and early use of assays employing the specific neutralization of a biological activity for the detection and quantification of binary toxin. The survey moves on to more recent immunological assays and culminates with a discussion of the relevance of binary toxin, especially CDT, in feces.

Keywords: *Clostridium perfringens*, *Clostridium difficile*, *Clostridium spiroforme*, iota, CDT, binary toxin, enterotoxin, diarrhea.

INTRODUCTION

Iota toxin and its close relatives, S toxin and CDT, are binary toxins produced by *Clostridium perfringens* Type E, *Clostridium spiroforme* and *Clostridium difficile*, respectively. Essential properties of the toxins and their corresponding genes have been reviewed in the past [1]. Each toxin is produced as two unrelated separate gene products, one an enzyme, the other a cell-binding / membrane translocation factor. In the case of iota toxin, the two components are called iota a (ia) and iota b (ib). This convention is followed for S toxin and CDT, thus their respective components are designated as Sa and Sb, plus CDTa and CDTb. The binding component of binary toxins is secreted as a propeptide that requires proteolytic activation to its mature, active form. Activation from pro-ib to ib and from pro-Sb to Sb occurs in culture fluids, but pro-CDTb is not activated *in vitro* which likely reflects a paucity of serine-type proteases produced by *C. difficile* in culture, different cleavage sites, or both. Exogenous trypsin is required for the activation of pro-CDTb [2]. Mature binding component rapidly oligomerizes on the surface of a target cell to form a heptameric, donut-shaped channel through which the enzyme enters the cell (following endosome acidification) and mono-ADP-ribosylates globular actin, thus killing the cell by disrupting its cytoskeleton and particularly filamentous actin formation [1, 3]. Neither component by itself has toxic activity, though individually each retains its respective binding and enzymatic properties.

Clostridial binary toxins are biologically active in various animal models. They kill mice when injected intraperitoneally and are dermonecrotic when injected intradermally

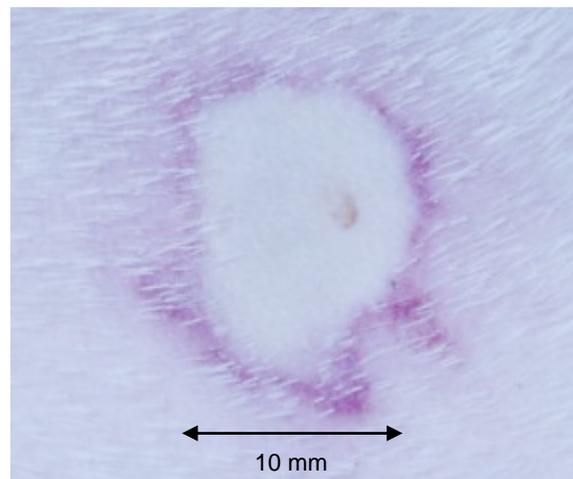
into the flanks of guinea pigs (Fig. 1) [4]. The binary toxins are enterotoxic and cause fluid to accumulate in ligated ileal loops of rabbits (Fig. 2) [5], an assay originally developed by workers studying cholera toxin and its activity. Clostridial binary toxins are readily cytotoxic to cultured cells [6]. All of these activities can be neutralized by antiserum to one or both of the binary components. Additionally, each component is also cross neutralized by antiserum to the individual components of the other binary toxins. Thus anti-iota toxin neutralizes S toxin of *C. spiroforme* and anti-Sb will neutralize *C. perfringens* iota toxin, etc. Such results suggest a high degree of structural similarity between the iota and iota-like toxin components produced by various clostridial species, which are all involved in enteric diseases of humans and other animals. Furthermore, it has been shown by different groups that chimeric toxins (i.e. Sb + ia, ib + Sa, etc.) formed from mixing heterologous components are biologically active [7].

In this short review, we will compare and contrast the assays used to detect binary toxins. An historical approach will be taken that progresses from assays for the biological activities of *C. perfringens* Type E iota toxin, the first clostridial binary toxin to be discovered, through today and the very latest immunological assays for *C. difficile* CDT. We will discuss what the presence of the binary toxins in feces means for the host.

1943-1980: CLOSTRIDIUM PERFRINGENS TYPE E IOTA TOXIN, MOUSE LETHALITY AND DERMONECROSIS

Iota toxin was first described as a cause of diarrhea in week-old calves [8]. Symptomatic animals had sterile stool filtrates that were toxic to mice when given by an intraperitoneal injection. Toxigenic isolates of *C. perfringens* were recovered from the same material and homologous antitoxin was subsequently raised. This antitoxin neutralized the toxic activity in fecal samples and culture fluids of *C. perfringens*. The toxin that caused this activity was named

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Sa (0.3 μg) and Sb (2.7 μg) in 0.3 mL injected intradermally

Fig. (1). Dermonecrosis lesion on the shaven flank of a guinea pig caused by *C. spiroforme* S toxin after 24 h.



Sa (1 μg) and Sb (9 μg) injected in 1 mL into each positive rabbit ileal loop

Fig. (2). Enterotoxic activity of *C. spiroforme* S toxin in rabbit ligated ileal loops.

iota toxin [9], and it became one of the four major toxins (i.e. alpha, beta, epsilon and iota) produced during log-phase growth that are now used to classify *C. perfringens* isolates as Type A, B, C, D or E. Thus, iota-producing strains were classed as Type E, and, while some Types make several toxins, only Type E isolates make iota toxin [4]. In 1986, Stiles and Wilkins [10, 11] showed iota to be a binary toxin, consisting of ia (the enzymatic mono-ADP-ribosyltransferase) and ib (cell binding/membrane translocation component). Stiles and his colleagues [12] went on to show that ib forms oligomers on the cell surface through which ia passes into the cell, where it disrupts actin monofilament formation.

For about 40 years, neutralizing the mouse-killing activity of iota toxin was both the means to detect iota toxin in feces and identify Type E isolates that make toxin *in vitro*.

A full typing assay for all the major toxins of *C. perfringens* was simply and comprehensively described by Sterne and Batty [4]. Their protocol is still used and has remained unchanged in anything but the finest of details. The set of four antitoxins, including Type E antitoxin needed for the assay, were sold in the 1970s by the Wellcome Research Laboratories, Beckenham, England. This antitoxin gradually became unavailable, starting in the 1980s. No new reliable commercial source has emerged since. In veterinary circles, particularly when mouse lethality is used to detect iota toxin, neutralization is nowadays accomplished with a research-use only reagent. If a mouse-killing, filterable toxin is detected and no neutralization is attempted (say from the want of a suitable antitoxin), PCR assays for ia, ib or both are used to make a diagnosis. However, such genetic-based diagnosis say nothing about the actual toxin production *in situ* that is, ultimately, responsible for disease.

There have been few diagnostic studies of the levels of iota toxin required for activity, and none connect toxin levels with disease severity. Obviously, when diarrhea is diagnosed by the mouse assay, a sample must contain at least 1 mouse lethal dose (MLD). This though is probably a gross underestimation. In a titration of highly purified ia and ib, 0.025 nmol of each (1.25 µg and 1.88 µg, respectively) represented 1 MLD when injected intraperitoneally [10]. Lower amounts were not always toxic (e.g. 0.013 nmol of each component killed only half the mice). 0.62 pmol of each (i.e. 40-fold less than 1 MLD) was not lethal to mice yet remained dermonecrotic in guinea pigs; 0.31 pmol had no activity in either assay. When the ib level was held constant but in excess (2.8 pmol, 200 ng) and ia was titrated, dermonecrosis was still present at ia levels of 0.088 pmol (4 ng). The cytotoxicity and enterotoxicity of purified iota toxin have not been properly titrated to date, or at least not reported in the literature.

On occasions, iota toxigenic isolates are found in the feces of healthy livestock [8]. It is possible that, on occasions, this represents a carrier state in which there was no *in vivo* production at all in these animals and that is why the animals were asymptomatic. Equally, extremely low levels (less than 1 MLD) of iota toxin may have been present, but being that low, the animals were unaffected. In any event, these examples of discrepant findings (no toxin, only toxigenic bacteria) will be no better served by PCR (no toxin, only potentially toxigenic bacteria). Of course with spore formers like clostridia, the amplification of a particular toxin gene may represent nothing more than transient spores, not colonization, with vegetative cell division equating to a persistent carrier state.

1978 TO 1986: CLOSTRIDIUM SPIROFORME S TOXIN AND MOUSE LETHALITY, DERMONECROSIS, ILEAL LOOPS AND CYTOTOXICITY

The S toxin of *C. spiroforme* was the second binary toxin to be characterised. It is closely related to *C. perfringens* iota toxin [5, 14]. S toxin is the cause of a mainly post-weaning diarrhea of rabbits that is often fatal [15, 16]. There are cases of unintended antibiotic-associated *C. spiroforme* diarrhea in rabbits [17]. Following clindamycin treatment, rabbits have been used to show *C. spiroforme* fulfilled Koch's postulates [18] and to study the course of the disease over time [19]. More recently S toxin-producing *C. spiroforme* has been isolated from horses with diarrhea (Unpublished data).

S toxin has all the same activities as iota toxin. It is mouse lethal, dermonecrotic, enterotoxic, and cytotoxic [5, 6, 20, 21]. Mouse killing and dermonecrosis - both modified from protocols in Sterne and Batty [4] - have routinely been the assays used to make a laboratory diagnosis. Thus, the initial reports of *C. spiroforme* diarrhea in rabbits were made by neutralizing the mouse-killing activity with antitoxin to *C. perfringens* Type E. Before the first reports of S toxin production by *C. spiroforme* from diarrheic rabbits, this cross neutralization led to the understandable but erroneous conclusion that *C. perfringens* Type E was the source of the toxin [16, 21-25]. However, in all of these cases, *C. perfringens* Type E was never isolated from diseased animals yet a curiously-coiled bacterium (*C. spiroforme*) was evident in only the sick, but not healthy, animals.

To date, there is very limited data on biological activity levels of purified Sa and Sb, and we know of only one quantitative assessment of S toxin levels in diarrheic feces. Thus, in a titration of Sa and Sb in combination, 100 ng of each represented 1 MLD [26]. This was approximately ten-fold lower than that reported for iota toxin [10, 11]. An important question then becomes, in a titration just what should be the ratio of ia to ib? Should ia and ib be titrated in combination or individually, but still remain as a mix with only one component changing in concentration? If the latter, what should the level be for the untitrated component? These considerations apply equally to all assays involving biological activity of binary toxins

Carman & Borriello [19] titrated the mouse lethal activity of S toxin in 5 rabbits experimentally infected with *C. spiroforme*. They injected mice with serial two-fold dilutions of filtrates from digesta collected at fixed time intervals via a cecal cannula. The bacterium was detected within 5 to 10 h of challenge, and about 5 to 10 h before S toxin had become detectable. Diarrhea began 15 to 20 h or so post challenge. In 3 animals, diarrhea and S toxin appeared simultaneously at 19 h. The mouse lethal titers from digesta of these 3 animals were respectively 1/1, 1/1, and 1/4, suggesting that 1 to 4 MLD/mL may approximate 1 enterotoxic dose of S toxin. If this is so, it is hardly surprising that a mix of Sa (1 µg) and Sb (9 µg) injected into rabbit ileal loops was enterotoxic [5], as it would have represented 10 or possibly more MLDs in Popoff's hands [26].

1986 TO PRESENT: CLOSTRIDIUM DIFFICILE CDT AND ELISAS

While studying the enzymatic activities of toxigenic *C. difficile*, Popoff and his group [27] found an ia-like ADP-ribosyltransferase in the culture fluids of *C. difficile* 196, an historical isolate from a human with diarrhea. The same group [28] went on to show, using immunoblots and cytotoxicity, the production of enzymatic and binding components for a binary toxin that they called CDT and found it to be very similar to both iota and S toxins at the functional, genetic and immunological levels. They also showed that in *C. difficile*, expression of cytotoxic levels of CDT was approximately forty-fold lower than that for *C. perfringens* iota toxin, a finding since confirmed by others [2]. A low level of transcription, not loss of specific activity from synthesized toxin components, explained the shortfall in CDT activity [27]. In fact while *cdtR* has been shown to regulate CDT expression in *C. difficile* [29], PCR and bioinformatics reveal that an equivalent *cdtR* is absent from both *C. spiroforme* and *C. perfringens* (Unpublished data). Each of the three binary toxin producers has an upstream sequence distinct from the others, pointing to possibly three distinct mechanisms of regulation. In many ways this is a remarkable twist on evolution, as at the protein level these clostridial binary toxins are very similar in various ways.

In about 2000, the effects of changes in the use of fluoroquinolone antibiotics began to manifest themselves in hospitals as an emerging resistance among *C. difficile* isolates to moxifloxacin. Though many different strains had become resistant, one in particular, ribotype 027, was identified as the dominant epidemic strain in several outbreaks worldwide [30, 31]. Previous outbreaks of *C.*

difficile had often been associated with the development of resistance to a particular antimicrobial agent. There have been other *C. difficile* outbreaks linked to clindamycin resistance, and others linked to resistance to cephalosporins. While resistance to widely used fluoroquinolone may help explain the recent frequency of cases, it probably has less to do with the parallel rise in the severity of *C. difficile* 027 disease reported by some but not others. Any link between severity of symptoms and 027 infections thus requires another explanation. There is the deregulation of toxins A and B expression during early log-phase growth in the test tube that is assumed to translate to increased toxin production *in vivo*. This is probably caused not by the widely reported deletions in the down regulator, *tcdC*, but by the less well-known stop codon resulting from a point mutation. Then there is CDT.

Ribotype 027 isolates carry *cdtA*, *cdtB*, and the regulatory gene, *cdtR*. All 3 are required for a functional CDT locus. Some non-CDT producing isolates carry a ghost locus, consisting of the entire *cdtR* but only parts of *cdtA* and *cdtB*. Most researchers have used the PCR technique to show the carriage of *cdtA*, *cdtB*, or both. Though many authors have, using PCR, identified many isolates that carry the CDT locus, very few have shown *in vitro* production of CDT by ribotype 027 isolates. When production is assayed, it is generally through immunoblotting and guinea pig dermonecrosis whenever proof of biological activity was needed. Titrations and firm quantitation of CDT in feces have not been reported.

Geric and her colleagues [2] studied CDT from IS 58, an otherwise non-toxic isolate that lacks genes for the large clostridial toxins A and B. IS 58 made low levels of CDT that were enterotoxic and required exogenous trypsin for activation, possibly because the peptidase cleavage site differs between pro-CDT_b versus pro-S_b and pro-I_b. Additionally, the protease activities of *C. difficile* may be inadequate compared to those in *C. perfringens* Type E and *C. spiroforme* cultures. Whether or not activation occurs *in vivo*, perhaps via pancreatic or microbial proteases provided by other bacterial flora, is not known. Possibly then, the want of activation coupled with a low level of expression may help explain why clindamycin-treated hamsters do not have diarrhea despite colonization by IS 58 [2].

In an unpublished study, TechLab used an enzyme immunoassay to detect CDT_b, the binding component of CDT, in antibiotic-associated diarrheal samples from humans and from which *C. difficile* was cultured. By comparison with a standard curve, the estimated levels of fecal cdt_b in 12 of 19 positive samples exceeded 100 ng/mL. Thus, these 12 samples contained a level of CDT_b that, had it been S_b, would have been lethal to mice [26] and exceeded the level of S_b in cecal digesta from clearly diseased, diarrheic rabbits [19]. So, even if CDT does not have a direct role in *C. difficile*-induced diarrhea, it may reach levels that contribute more indirectly to the overall clinical picture, in a complementary way not seen with otherwise non-toxicogenic isolates like IS 58.

CONCLUSIONS

As our knowledge of binary toxins has grown, techniques for their detection have become less dependent upon

neutralization of biological activity and more so upon immunoassays, though none of the toxins are yet well served by a commercial assay. The role of Iota and S toxins in causing diarrhea is beyond dispute, but the minimum doses needed for activity are not known. A similar role for CDT remains unproven, though it is a distinct possibility as fecal levels from cases of human disease sometimes match the levels of S toxin that occur in diarrheic animals. The correlation of CDT amounts with disease severity in humans should be addressed in future studies.

Footnote Added During Preparation

Shwan *et al.* [32] recently offered a new explanation of how CDT might enhance the virulence of *C. difficile*. They suggested that CDT induces the formation of microtubules that protrude from the intestinal epithelial cells. These microtubules form a dense mesh, rich in capture proteins, that wrap and embed *C. difficile* and effectively increase its attachment to and persistence on the host's epithelium.

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