Importance of Fibronectin for *Clostridium perfringens* Adhesion to Host Tissue

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Abstract

*Clostridium perfringens*, a Gram-positive, spore-forming, obligate anaerobe, is the cause of gas gangrene and food poisoning. Colonization of the host requires adhesion of bacterial cells to injured tissue. We review here recent results regarding the role of fibronectin in the interaction of *C. perfringens* cells with host collagens.

Keywords: *Clostridium perfringens*; Fibronectin; Fibronectin-binding proteins

Abbreviations

Fn: Fibronectin; Fbp: Fn-binding protein; BSA: Bovine Serum Albumin; HRP: Horseradish Peroxidase; ELISA: Enzyme-linked Immunosorbent Assay

Mini Review

*Clostridium perfringens*, the cause of gas gangrene and food poisoning, exerts its pathogenicity by secreting toxins [1]. Delivery of toxins follows *C. perfringens* colonization of a wound site, which requires the adherence of bacterial cells to damaged host tissue. We investigated the adhesive interaction of *C. perfringens* cells with collagen, a typical component of connective tissue. Although *C. perfringens* cells exhibited marginal binding to the surfaces of microplate wells coated with collagen type I, the bacterial cells never exhibited binding to wells coated with collagen type II, collagen type III, or gelatin [2]. However, *C. perfringens* showed strong binding to immobilized collagens and gelatin when the proteins were precoated with fibronectin (Fn). This observation implies that the adhesion of *C. perfringens* cells to collagens is an Fn-mediated process.

Fn is a major plasma component, and is also a ubiquitous extracellular matrix protein. Fn plays a pivotal role in many cellular processes, including wound healing, tissue structure formation, and cell migration [3]. Fn is a disulfide-linked homodimeric glycoprotein of approximately 450 kDa, with each Fn monomer consisting of three types of repeating modules: 12 type I, two type II, and 15-17 type III (Figure 1). The N-terminal domain is composed of five type-I modules (I1-I5), which correspond to the sites of binding for most bacterial Fn-binding proteins (Fbps) [4], and an adjacent gelatin-binding domain containing four type-I modules and two type-II modules (I6-I10). The central part of the Fn polypeptide is composed of 15-17 type-III modules, including the region (III10) responsible for binding of host cells. In contrast to both type-I and type-II modules, type-III modules lack intramodular disulfide bonds.

Figure 1: Schematic representation of the modular structure of the Fn protein.

Tissue injury is usually accompanied by bleeding, leading to coating of connective tissue with Fn, which promotes tissue reconstitution. Given the importance of Fn for the binding of *C. perfringens* cells to collagens, two questions arise.
The first question is, what are the Fn receptor(s) on *C. perfringens* cells? To date, we have identified two FBPs, FbpC and FbpD, among the peptidoglycan-associated membrane proteins of *C. perfringens* [5]; the identification of additional FBPs is expected. FbpC (CPE0625; 56kDa) and FbpD (CPE0630; 45 kDa) are encoded with signal peptides and contain putative cell wall-binding repeats; FbpC and FbpD also harbor a peptidase motif and a peptidoglycan hydrolase motif, respectively. Recombinant forms of both FbpC and FbpD exhibited binding to Fn. Moreover, a preliminary experiment suggested that both anti-FbpC and anti-FbpD antibodies partially inhibit the binding of *C. perfringens* cells to gelatin coated with Fn (unpublished data). These observations strongly suggest the involvement of both FbpC and FbpD in the adhesion of *C. perfringens* cells to Fn-coated collagen (represented here by gelatin).

The second question is, what site on the Fn molecule is bound by the bacterial Fn receptors? To elucidate this domain, we employed several anti-Fn monoclonal antibodies that recognize distinct epitopes [6]. Among these reagents, the HB39 monoclonal antibody (IgG isotype), which has been shown to recognize the III9-10 region of Fn, was found to inhibit Fn binding by *C. perfringens* cells. Furthermore, Fn binding by *C. perfringens* cells was competitively inhibited by the recombinant III9 fragment of Fn (rIII9), but not by rIII2-4, rIII5-7, rIII8, or rIII10 fragments. Moreover, *C. perfringens* cells bound weakly, but significantly, to microplate wells coated with either rIII9 or rIII10. In contrast, bacterial cells never bound to wells coated with rIII2-4, rIII5-7, or rIII8. These results suggest that the Fn receptor(s) on *C. perfringens* cells recognize the III9-10 region of Fn.

Because the individual rIII9 and rIII10 fragments showed weak affinity for *C. perfringens* cells, we additionally generated a recombinant III9-10 fragment of Fn and tested this preparation for bacterial adherence (Figure 2). The results showed that *C. perfringens* cells bound immobilized rIII9-10 with avidities similar or higher than that for immobilized Fn. This observation implies that the intermodule linker between III9 and III10 is critical for the binding by the Fn receptors on *C. perfringens* cells.

Taken together, these data indicate that Fn plays an important role in the adherence of *C. perfringens* cells to collagen, a major matrix protein in the connective tissue. At least two Fn-binding proteins, FbpC and FbpD, are involved in the binding of *C. perfringens* cells to Fn. Specifically, *C. perfringens* cells target the III9-10 domain of Fn. Indeed, recombinant FbpC and FbpD both appear to bind to the rIII9-10 region of Fn (unpublished data).

Although most bacterial FBPs have been shown to bind the N-terminal of Fn [4], several Fn receptors that bind to the III9-10 region of Fn have been identified, including the Yersinia pestis Ail protein [7] and the Pasteurella multocida PM1665 protein [8]. Ail and PM1665 reportedly bind to III9 and III9-10, respectively. Notably, Ail is critical for the delivery of cytotoxic Yop proteins to host cells and is a major virulence factor for *Y. pestis* pathogenesis [9]. It will be of interest to determine whether III9-10-mediated Fn binding by *C. perfringens* cells affects the toxin secretion and/or virulence of this bacterium.

The III10 region contains a unique peptide sequence, RGD that interacts with integrins on the host cell surface [10]. Thus, it is possible that Fn-bearing *C. perfringens* cells can escape from integrin-mediated binding to host cells, contributing to evasion of immune surveillance. Further work will be needed to assess the biological significance of Fn binding by *C. perfringens* in the context of the pathogenesis of clostridial gas gangrene and food poisoning.

![Figure 2: Binding of *C. perfringens* cells to immobilized full-length or truncated Fn, as detected by ELISA. The individual wells of a microplate were coated (at 1 µg/well) with either Fn, rIII9-10, rabbit anti-*C. perfringens* IgG antibody, or BSA. Biotinylated *C. perfringens* cells were added to each well and plates were incubated for 1 h at room temperature. After washing, HRP-conjugated streptavidin was added and incubated. HRP activity was measured by colorimetric assay, with detection via absorbance at 405 nm (A405), indicative of the number of bound bacterial cells. A405 values are presented as mean ± SD (n=5). **p <0.01 vs. BSA (one-way ANOVA followed by Bonferroni multiple comparison test).](image)

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