Interactions with fibronectin attenuate the virulence of *Streptococcus pyogenes*

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Fibronectin-binding surface proteins are found in many bacterial species. Most strains of *Streptococcus pyogenes*, a major human pathogen, express the fibronectin-binding protein F1, which promotes bacterial adherence to and entry into human cells. In this study, the role of fibronectin in *S. pyogenes* virulence was investigated by introducing the protein F1 gene in an *S. pyogenes* strain lacking this gene. Furthermore, transgenic mice lacking plasma fibronectin were used to examine the relative contribution of plasma and cellular fibronectin to *S. pyogenes* virulence. Unexpectedly, protein F1-expressing bacteria were less virulent to normal mice, and virulence was partly restored when these bacteria were used to infect mice lacking plasma fibronectin. Dissemination to the spleen of infected mice was less efficient for fibronectin-binding bacteria. These bacteria also disseminated more efficiently in mice lacking plasma fibronectin, demonstrating that plasma fibronectin bound to the bacterial surface downregulates *S. pyogenes* virulence by limiting bacterial spread. From an evolutionary point of view, these results suggest that reducing virulence by binding fibronectin adds selective advantages to the bacterium.

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Introduction

Streptococcus pyogenes (group A streptococcus) is a significant human pathogen. This Gram-positive bacterium causes common and mostly uncomplicated suppurative infections such as pharyngitis, impetigo, and erysipelas, and also nonsuppurative sequelae (acute rheumatic fever, poststreptococcal glomerulonephritis, and reactive arthritis) and severe inva-

sive infections including necrotising fasciitis, sepsis, and a toxic shock-like syndrome (for references, see the review by Cunningham, 2000). *S. pyogenes* virulence is based on a multitude of secreted and surface-associated components. An important group of virulence factors is the cell wall-attached surface proteins. In this group, M proteins, elongated coiled-coil proteins that bind a variety of human proteins (see the review by Fischetti, 1989), are responsible for the ability of *S. pyogenes* to survive and multiply in human blood. Several other streptococcal surface proteins have been identified and characterised. For many of these, isogenic mutants are less virulent in animal models of infection. Mutants deficient in the M protein (Ashbaugh *et al*, 1998), the C5a peptidase that cleaves and inactivates C5a of the complement system, thereby disturbing chemotaxis and neutrophil migration (Wexler *et al*, 1985), and GRAB that binds the plasma proteinase inhibitor α_2 -macroglobulin (Rasmussen *et al*, 1999), are all less virulent in animal challenge studies. Mutants lacking the fibronectin (Fn)-binding surface proteins SOF (Courtney *et al*, 1999), Fba, or FbaB (Terao *et al*, 2001, 2002) also show reduced virulence.

Fibronectins are dimeric glycoproteins that exist in human blood plasma (plasma Fn; pFn) at a concentration of 300 μ g/ml (Hynes, 1990), and in the extracellular matrix (cellular Fn; cFn). cFn is produced locally in tissues by a variety of cells, whereas hepatocytes are the sole source of pFn (Owens and Cimino, 1982; Mosher, 1989; Hynes, 1990; Schwarzbauer, 1991). Binding of Fn by *S. pyogenes* mediates bacterial adhesion to human cells (Abraham *et al*, 1983). Several *S. pyogenes* surface proteins are implicated in this binding, including M and M-like proteins (Schmidt *et al*, 1993; Frick *et al*, 1995; Cue *et al*, 1998), FBP54 (Courtney *et al*, 1999), PFBP (Rocha and Fischetti, 1999), glyceraldehyde-3 phosphatase dehydrogenase (Pancholi and Fischetti, 1992), protein F2 (Jaffe *et al*, 1996), SOF, Fba and FbaB (Terao *et al*, 2001, 2002), and protein F1/SfbI (Hanski and Caparon, 1992; Talay *et al*, 1992). The binding of Fn by protein F1 has been studied extensively *in vitro*. Protein F1 binds Fn via two distinct domains: one repeat-containing domain (RD) in the COOH-terminal region with homology to Fn-binding regions in *Staphylococcus aureus* and *Streptococcus dysgalactiae*, and a second more NH₂-terminally located domain (upstream Fn-binding domain; UFB) that binds Fn with higher affinity (Sela *et al*, 1993). These domains bind to different parts of Fn (Ozeri *et al*, 1996; Talay *et al*, 2000). A mutant deficient in protein F1 is defective in adhesion (Hanski and Caparon, 1992), and nonadhesive bacteria devoid of protein F1 became adhesive when complemented with protein F1 (Hanski *et al*, 1992). Moreover, the protein F1 homologue SfbI blocks *S. pyogenes* adhesion when added in solution (Talay *et al*, 1992). Thus, protein F1 is both required and sufficient for effective adhesion in these systems. More recently, it was demonstrated that protein F1 is also important for the entry of *S. pyogenes* into human cells (Jadoun *et al*, 1997; Molinari *et al*, 1997), although Fn-independent internalisation

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(Molinari *et al*, 2000) may also occur. Cellular invasion by *S. pyogenes* is blocked by antibodies against β_1 -integrins (Cue *et al*, 1998; Ozeri *et al*, 1998) and by an $\alpha_5\beta_1$ -integrin antagonist (Cue *et al*, 2000). Streptococcal internalisation occurs *in vivo* as well as *in vitro*, and several studies suggest that invasion of host cells by *S. pyogenes* creates an *in vivo* reservoir of bacteria that could account for recurrent infection (Österlund and Engstrand, 1995; Kiska *et al*, 1997; Österlund *et al*, 1997; Cockerill *et al*, 1998; Neeman *et al*, 1998).

Adhesion is an initial and important step in bacterial infections, but so far no animal studies have been published regarding the contribution of protein F1 to the virulence of *S. pyogenes*. Furthermore, due to the lack of appropriate animal models of infection, the effect of pFn and/or cFn on bacterial virulence has not been studied. Recently, we generated mice lacking pFn using the *Cre-loxP* recombinant technology (Sakai *et al*, 2001). In the present work, such animals and *S. pyogenes* strains lacking Fn-binding activity or expressing Fn-binding protein F1 were used to investigate the role of Fn in *S. pyogenes* virulence. Unexpectedly, interactions between protein F1 and Fn were found to downregulate virulence.

Results

Generation and characteristics of AP1 bacteria expressing protein F1

To generate AP1 bacteria with a high binding capacity for Fn, the pLZ12 self-replicating plasmid without (pLZ12) or with (pPTF8) the insert encoding protein F1 were electroporated into AP1 bacteria. The transcomplemented bacteria with or without protein F1 showed the same growth rate at chloramphenicol (Cm) concentrations up to 6 μ g/ml (data not shown). Nontransformed, parental AP1 bacteria failed to grow at 1.5 μ g Cm/ml. Furthermore, the strains with or without protein F1 had similar growth rates in fresh human blood and bound fibrinogen to the same degree (data not shown). In these strains, the M1 protein is responsible for fibrinogen binding and the binding curves demonstrated that very similar amounts of M1 protein were expressed by the two strains. AP1(pPTF8), but not AP1 or AP1(pLZ12) bacteria, bound significant amounts of radiolabelled human Fn (Figure 1A). Moreover, AP1(pPTF8) bacteria expressing protein F1, but not AP1(pLZ12) bacteria, absorbed Fn from mouse serum (Figure 1B), showing that protein F1-expressing bacteria bind Fn also in a complex mixture of proteins.

To determine whether binding of Fn to the bacterial surface would allow secondary binding of other serum proteins, or whether the absence of pFn would result in interactions between protein F1 and other serum proteins, AP1(pPTF8) and AP1(pLZ12) bacteria were incubated with serum from normal (control) and pFn-null mice. Serum proteins bound to the bacteria were eluted, separated by SDS-PAGE under reducing conditions, and silver stained (Figure 2A). A protein with a molecular mass corresponding to pFn was absorbed from normal mouse serum by AP1(pPTF8) bacteria (Figure 2A, STAIN). Western blotting verified that the protein band corresponded to Fn (Figure 2A, BLOT). Furthermore, this protein band was not present in the other eluted samples (Figure 2A, STAIN). Apart from Fn, no qualitative differences were observed between the different lanes. The results de-

Figure 1 AP1 bacteria expressing protein F1 bind human and mouse Fn. (A) The binding of human 125 I-labelled Fn to serial log dilutions (from 2×10^9 cfu's) of AP1, AP1(pLZ12), and AP1(pPTF8) bacteria is shown. The error bars indicate $\pm$ s.d. (B) AP1(pLZ12) or AP1(pPTF8) bacteria were incubated with 10% mouse serum, and bound proteins were eluted with 0.1 M glycine-HCl, pH 2.0. The proteins were separated by SDS-PAGE (reducing conditions) and subjected to Western blot analysis using anti-Fn antibodies. Human Fn, 20 or 100 ng, was run as a control.

Figure 2 Silver staining of absorbed serum proteins. (A) AP1(pLZ12) or AP1(pPTF8) bacteria were incubated with serum from control (pFn⁺) or pFn-null (pFn⁻) mice. Bound proteins were eluted with 0.1 M glycine-HCl, pH 2.0, and separated by SDS-PAGE (reducing conditions) on two identical gels. One gel was silver stained (STAIN), and the other was used for Western blotting (BLOT). A band with the size of Fn (indicated by the arrow) is present in the material eluted from the AP1(pPTF8) bacteria incubated with serum from control mice (pFn⁺), but not in the other lanes. Western blotting with anti-Fn antibodies confirmed that this band was Fn. (B) Serum proteins from pFn-null mice (pFn⁻) or control mice (pFn⁺) were separated by SDS-PAGE and subjected to Western blotting with anti-Fn antibodies.

monstrate that AP1(pPTF8), but not AP1(pLZ12) bacteria, bind pFn, and suggest that the binding of Fn does not interfere with binding of other serum proteins. Moreover, no additional protein bands were visible when AP1(pPTF8) was incubated with pFn-null serum, indicating that the absence of pFn does not promote interactions between protein F1 and other serum proteins.

Effect of fibronectin on *S. pyogenes* adhesion and internalisation

The effect of pFn on bacterial adhesion and entry into mammalian cells was studied *in vitro* by performing adhesion and internalisation experiments with the pharyngeal epithelial cell line Detroit 562 or human fibroblasts. First, the density of cFn on the surface of these cells was determined by ELISA (Figure 3). In these experiments, anti-Fn antibodies were applied in different dilutions to pharyngeal epithelial cells or fibroblasts. For comparison, wells were coated with different concentrations of purified Fn, followed by the application of anti-Fn antibodies. The results indicate a four- to eight-fold higher density of Fn on the surface of fibroblasts as compared to Detroit 562 cells. To evaluate the importance of pFn for adhesion and internalisation, experiments were subsequently performed with serum from control mice (control serum) or from pFn-null mice (pFn-null serum).

When protein F1 was expressed on the surface of AP1 bacteria, the adherence to epithelial cells in the presence of control serum increased by 65% (95% confidence interval 40–90%) (Figure 4A). Importantly, the absence of pFn reduced the adhesion of bacteria expressing protein F1 to epithelial cells (Figure 4A). In the presence of control serum, the entry of AP1(pPTF8) into epithelial cells was increased almost three-fold (95% confidence interval 1.5–4) compared to AP1(pLZ12). When the bacteria and cells were incubated in pFn-null serum, internalisation of the two strains was similar (Figure 4B). The results show that *S. pyogenes* adhesion to and entry into epithelial cells increase when protein F1 is expressed on the bacterial surface. Furthermore, both effects were only observed in the presence of pFn, demonstrating that protein F1 *per se* is not sufficient. Finally, the data show that protein F1-independent adhesion and internalisation is not influenced by the presence of pFn.

To examine the relative importance of pFn for bacterial adhesion to and internalisation into cells expressing large amounts of cFn, AP1(pPTF8) or AP1(pLZ12) bacteria were incubated with human foreskin fibroblasts in the presence of either control or pFn-null serum. Adhesion of AP1(pPTF8) bacteria to fibroblasts was increased three-fold (95% confidence interval 2.5–3.4) compared to AP1(pLZ12) (Figure 4C) in control serum. Interestingly, a similar increase was obtained in pFn-null serum, which was in contrast to the results

Figure 3 Cellular Fn is more abundant on the surface of fibroblasts than on pharyngeal epithelial cells. Different dilutions of anti-Fn antibodies were applied to wells coated with epithelial cells or fibroblasts, followed by peroxidase-labelled secondary antibodies. The results are from one representative experiment with triplicate samples and the error bars indicate $\pm$ s.d. For comparison, the wells were coated with different concentrations of purified Fn, followed by anti-Fn antibodies and secondary antibodies.

Figure 4 Effect of Fn on *S. pyogenes* adhesion and internalisation. Adherence and internalisation experiments were performed with AP1(pLZ12) or AP1(pPTF8) bacteria. The cells used were a human pharyngeal cell line (Detroit 562) (A, B) and human foreskin fibroblasts (C, D). Experiments were performed in the presence of serum from control mice (black bars) or serum from pFn-null mice (grey bars). Values are the means of four experiments with triplicate samples, given as a percentage of the mean adhesion or internalisation of AP1(pLZ12). Error bars $\pm$ s.e.m. and *P*-values are indicated.

seen with epithelial cells (Figure 4A). Internalisation of AP1 bacteria expressing protein F1 was almost four-fold increased in control serum when compared to AP1 bacteria lacking protein F1 (Figure 4D). In the presence of pFn-null serum, however, internalisation of AP1(pPTF8) was similar to AP1(pLZ12). These findings indicate that efficient internalisation, but not adhesion, depends on pFn bound to the bacterial surface by protein F1.

S. pyogenes bacteria expressing protein F1 are less virulent

Unlike most strains of *S. pyogenes*, the AP1 strain is virulent to mice (Björck *et al*, 1989) and it was therefore used in an animal model simulating invasive *S. pyogenes* disease. A potential problem when introducing the protein F1 gene on a plasmid in the AP1 strain concerns its regulation. However, it has been published that other M1 strains, despite their lack of *prtF*, still carry *RofA*, the activator of the gene (Granok *et al*, 2000). To investigate the virulence of AP1(pPTF8) relative to AP1(pLZ12) bacteria *in vivo*, 10^5 or 10^6 bacteria were injected intraperitoneally (i.p.) into sex- and age-matched groups of control mice. While 81% (13 of 16) of the mice injected with AP1(pLZ12) died within 8 days after injection, only 44% (17 out of 39) mice injected with AP1(pPTF8) died within the same period (Figure 5A). This difference is statistically significant ($P = 0.016$) and suggests that protein F1 attenuates virulence. When 10^6 bacteria were injected, 50% (three of six) of the AP1(pPTF8)-injected mice died, compared to 100% (six of six) of the mice injected with AP1(pLZ12) (data not shown). Although not statistically significant ($P = 0.18$), the results could indicate that binding of Fn also at this high bacterial load influences virulence.

As Fn-binding bacteria (AP1(pPTF8)) were less virulent to control mice, the effect of pFn on bacterial virulence was investigated using pFn-null mice. In all, 10^5 AP1(pPTF8)

Figure 5 I.p. challenge of mice with *S. pyogenes*. (A) Control mice were injected i.p. with 10^5 AP1(pLZ12) or AP1(pPTF8) bacteria. The mice were monitored daily, and the percentage of surviving mice at various days after injection is shown. Significantly more mice infected with AP1(pPTF8) were alive at the end of the experiment ($P=0.017$). (B) Survival rate in percent when control and pFn-null mice were injected i.p. with 10^5 AP1(pPTF8) bacteria. At the end of the experiment, 56% of the control mice were alive as compared to 32% of the pFn-null mice ($P=0.07$).

bacteria were injected i.p. into such animals. While 68% (15 of 22) of the pFn-null animals died within 8 days after injection, 44% (17 of 39) of the control mice died during the same time period (Figure 5B). This difference is not statistically significant ($P=0.07$), but the results could suggest that the reduced virulence of AP1(pPTF8) bacteria in control mice (Figure 5A) is partially restored in pFn-null mice. As pFn-null mice still express cFn in all tissues and protein F1 can interact with cFn (Ozeri *et al*, 1996; Molinari *et al*, 1997), the data support the notion that cFn prevents full restoration of virulence in pFn-null mice.

Fn-binding AP1 bacteria disseminate less efficiently to the spleen of infected mice

When AP1 bacteria are injected i.p. into mice, bacteria spread to the spleen within a few hours after administration (data not shown). To investigate whether bacterial Fn binding and pFn could affect the dissemination process, and hence could be responsible for the reduced virulence, a system was designed where different variants of AP1 bacteria were injected into control mice or pFn-null mice. To ensure that the effect of protein F1 on virulence was due to the Fn-binding capacity of the protein, plasmid constructs were made containing the protein F1 gene without the RD and UFBD domains (Figure 6A). These constructs were electroporated into AP1 bacteria, generating AP1(pPTF8 Δ BD). AP1(pPTF8 Δ BD) bacteria showed the same growth rate as AP1(pLZ12) and AP1(pPTF8) bacteria at Cm concentrations up to 6 μ g/ml (data not shown). Also, AP1(pPTF8 Δ BD) bacteria only bound background levels of radiolabelled Fn (Figure 6B) and did not absorb Fn from mouse serum (Figure 6C). Finally, AP1(pPTF8 Δ BD) bacteria behaved as AP1(pLZ12) bacteria in experiments analysing the adherence to and internalisation into fibroblasts (Figure 6D and E) and epithelial cells (data not shown). These data verify earlier findings that protein F1 interacts with Fn through the RD and the UFBD domains, and demonstrate that these domains are responsible for the contribution of protein F1 to plasma- and cellular Fn-mediated adhesion and internalisation.

For the dissemination experiments, control mice or pFn-null mice were injected ip with AP1(pLZ12), AP1(pPTF8), or AP1(pPTF8 Δ BD) bacteria. The mice were killed 24 h later to

Figure 6 Characterisation of *S. pyogenes* mutant bacteria expressing protein F1 lacking the Fn-binding domains. (A) A plasmid with a truncated protein F1 gene, lacking the regions coding for the upstream Fn-binding domain (UFBD) and the repeat-containing domain (RD), was generated by performing inverse PCR on the plasmid pPTF8. The resulting plasmid, designated pPTF8 Δ BD, was electroporated into AP1 bacteria, generating AP1(pPTF8 Δ BD) bacteria. (B) The binding of human 125 I-labelled Fn to serial log dilutions (from 2×10^9 cfu's) of AP1(pPTF8) and AP1(pPTF8 Δ BD) bacteria is shown. The error bars indicate $\pm$ s.d. (C) AP1(pLZ12), AP1(pPTF8), or AP1(pPTF8 Δ BD) bacteria were incubated with 10% mouse serum, and bound proteins were eluted with 0.1 M glycine-HCl, pH 2.0. The proteins were separated by SDS-PAGE (reducing conditions) and subjected to Western blot analysis using anti-Fn antibodies. (D, E) The adherence to and internalisation into human fibroblasts was determined for AP1(pLZ12), AP1(pPTF8), and AP1(pPTF8 Δ BD) bacteria. Experiments were performed in the presence of serum from control mice (black bars) or serum from pFn-null mice (grey bars). Values are the means of three experiments with triplicate samples, given as a percentage of the mean adhesion or internalisation of AP1(pLZ12). Error bars $\pm$ s.e.m. are indicated.

determine the number of colony-forming units (cfu) in their spleens (Figure 7). Control mice infected with AP1(pPTF8) bacteria expressing protein F1 had significantly less bacteria in the spleens than control mice infected with AP1(pLZ12) or AP1(pPTF8 Δ BD) bacteria. Furthermore, a significantly lower number of AP1(pPTF8) bacteria as compared to AP1(pLZ12) or AP1(pPTF8 Δ BD) bacteria were isolated from the spleens of pFn-null mice. Finally, the number of AP1(pPTF8) bacteria in the spleens of pFn-null mice was significantly higher than in control mice. Combined, these results suggest that protein F1 reduces the spreading from the peritoneal cavity to the spleen by interacting with both pFn and cFn, and that the reduction is dependent on the presence of the Fn-binding domains of protein F1.

S. pyogenes adherence to and entry into mouse fibroblasts lacking cFn

To further test whether cFn can modulate *S. pyogenes* adherence to and entry into mammalian cells, embryonic

Figure 7 Bacterial dissemination to the spleen. Control mice (open symbols) or pFn-null mice (black symbols) were injected i.p. with AP1(pLZ12), AP1(pPTF8), or AP1(pPTF8ΔBD). The mice were killed 24 h after injection and the total number of cfu's in the spleen was determined for each mouse. The lines represent the median value in each group. *P*-values were determined with the Mann-Whitney *U*-test.

Figure 8 *S. pyogenes* adhesion and internalisation experiments using murine cells lacking cFn. Adhesion and internalisation experiments were performed with mouse fibroblasts lacking the Fn gene and AP1(pLZ12), AP1(pPTF8), or AP1(pPTF8ΔBD) bacteria (A, B). The experiments were performed in the presence of serum from control mice (black bars) or serum from pFn-null mice (grey bars). Values are the means of four experiments with triplicate samples, given as a percentage of the mean adhesion or internalisation of AP1(pLZ12). Error bars $\pm$ s.e.m. and *P*-values are indicated.

fibroblasts were isolated from floxed Fn mice, immortalised with the SV40 large T antigen and subsequently treated with adenovirus transducing Cre to delete both Fn genes (Fn^{-/-}). Western blotting revealed high cFn expression in floxed cells and confirmed the complete absence of cFn expression upon Cre-mediated deletion (data not shown). Initial experiments demonstrated that bacterial adhesion and entry of control mouse fibroblasts and human foreskin fibroblasts were similar. In control serum, adhesion of AP1(pPTF8) bacteria to Fn^{-/-} fibroblasts was increased by 78% (95% confidence interval 46–110%) compared to AP1(pLZ12) and by 65% (95% confidence interval 35–90%) as compared to AP1(pPTF8ΔBD). In contrast, there was no difference in adhesion between the three bacterial strains when pFn-null serum was used (Figure 8A). Similarly, in the presence of control serum, internalisation of AP1(pPTF8) was increased over three-fold compared to AP1(pLZ12) (95% confidence

interval 2.9–3.5) and AP1(pPTF8ΔBD) (95% confidence interval 3.0–3.5), but there were no significant differences between the bacterial strains in experiments performed in pFn-null serum (Figure 8B). Taken together, these data suggest that both cFn and pFn participate in protein F1-mediated adhesion, whereas protein F1-mediated entry depends on pFn only.

Discussion

Fn-binding proteins are important for the adhesion and cellular entry of several major bacterial pathogens, suggesting that these molecules could play a role in virulence. However, virulence is a highly polygenic property and it is difficult to design models, especially animal models, by which the significance of a putative virulence factor can be explicitly investigated. The aim of the present study was therefore to establish an experimental system where Fn-dependent adhesion and internalisation could be studied *in vitro*, and be directly correlated to virulence *in vivo*. This was achieved by utilising a combination of well-defined *S. pyogenes* mutant strains, different cell lines, and transgenic mice.

Binding of Fn is crucial for *S. pyogenes* adhesion and in most *S. pyogenes* strains protein F1 is the major Fn-binding protein. As shown here, the introduction of protein F1 in a strain with low Fn-binding activity resulted in a 65% increase of adhesion to epithelial cells. This effect was dependent on the presence of pFn, suggesting that binding of pFn promotes adherence. Most studies of *S. pyogenes* adhesion have used epithelial cells (for references, see Courtney *et al*, 2002), cells that the bacteria will encounter when, for instance, colonising the throat. However, viral infections that often precede *S. pyogenes* pharyngitis will damage the epithelial cell layer and expose subepithelial cellular and matrix structures. Moreover, in wounded skin or deep tissue infections, *S. pyogenes* will come across numerous host cell types and matrix components. This study shows that the adhesion of *S. pyogenes* to fibroblasts does not require serum. The cFn produced by these cells, present both as an intercellular matrix and at the cell surface, is apparently sufficient to promote bacterial adhesion. AP1(pLZ12) bacteria lacking protein F1 and AP1(pPTF8ΔBD) bacteria expressing protein F1 without the Fn-binding domains adhered similarly to fibroblasts and to epithelial cells, whereas the adherence of bacteria expressing intact protein F1 increased about three-fold. The same increase was recorded when normal fibroblasts and pFn-null serum were used, but not with fibroblasts lacking cFn. Thus, a majority of Fn-binding bacteria will adhere to fibroblasts also in the absence of pFn, most likely through binding to cFn via the RD and the UFBD domains of protein F1.

The ability of *S. pyogenes* to invade human epithelial cells was first described by LaPenta *et al* (1994), a property that has been correlated with the failure to eradicate the bacteria with penicillin allowing for recurrent infection (Neeman *et al*, 1998). In the present work, protein F1 was found to enhance internalisation in a pFn-dependent manner. This should be due to interactions between pFn, bound to protein F1 at the bacterial surface, and Fn-binding integrins (Ozeri *et al*, 1998). Moreover, even though protein F1-mediated adhesion to fibroblasts also occurred in the absence of pFn, the increased entry into fibroblasts by protein F1-expressing bacteria was

found to require pFn. At least during inflammatory and exudative phases of *S. pyogenes* infection, pFn will be available to the bacteria and facilitate entry into human cells.

In the present study, expression of protein F1 was found to reduce the virulence of *S. pyogenes*. This finding was quite unexpected, as the introduction of protein F1 into the AP1 strain did not affect growth rate or phagocytic killing, measured as the ability of the bacteria to survive in human blood. Moreover, a number of previous investigations have shown that most surface proteins of *S. pyogenes* enhance virulence (for references, see Navarre and Schneewind, 1999). LaPenta *et al* (1994) showed a correlation between high internalisation rates and invasive disease, and *S. pyogenes* mutants that are deficient in the Fn-binding proteins SOF, Fba, and FbaB were all reported to show attenuated virulence (Courtney *et al*, 1999; Terao *et al*, 2001, 2002). However, it is possible that functions other than Fn binding could contribute to their role in virulence. Thus, SOF is also a lipoproteinase (Courtney *et al*, 1999) and Fba limits C3b deposition on the bacterial surface through binding of Factor H and Factor H-like protein-1 (Pandiripally *et al*, 2002). Moreover, a recent study demonstrated that Fn bound to the protein F1 homologue Sfb1 mediates interactions with collagens (Dinkla *et al*, 2003). These interactions were found to reduce phagocytosis, but their influence on virulence was not investigated. There are, however, previous studies that support our conclusion that Fn binding and internalisation decrease virulence. Epidemiological data revealed a negative correlation between Fn binding and severity of disease (Natanson *et al*, 1995), and *S. pyogenes* isolates from patients with systemic disease were reported to internalise poorly (Jadoun *et al*, 1998; Molinari and Chhatwal, 1998). Moreover, *S. pyogenes* expressing a virulence-promoting capsule do not internalise as efficiently as non-capsulated bacteria (Cywes and Wessels, 2001), and a recent study reported that a mutant of *Staphylococcus aureus* deficient in Fn binding was more virulent than Fn-binding wild-type bacteria (McElroy *et al*, 2002).

Protein F1 can interact with both pFn and cFn (Okada *et al*, 1997). As mentioned above, invading *S. pyogenes* bacteria are likely to encounter cFn present in the extracellular matrix and at the surface of many cell types. As the infection proceeds and an inflammatory response is induced, plasma proteins, including pFn, will extravasate due to increased vascular permeability. In this phase, or if bacteria disseminate to the bloodstream, interactions with pFn will dominate. In order to evaluate the relative contributions of pFn and cFn to *S. pyogenes* virulence during the course of infection, the Fn-binding AP1(pPTF8) strain was injected into pFn-null mice. These animals have a deletion of the Fn gene only in the hepatocytes where pFn is produced, while the production of cFn is normal in all other tissues (Sakai *et al*, 2001). Consequently, protein F1-expressing bacteria will interact exclusively with cFn in these mice. Although not statistically significant, there was a tendency that mice without pFn were more susceptible to lethal infections with Fn-binding bacteria. However, in these animals, the bacteria expressing protein F1 are still not as lethal as the bacteria lacking protein F1, when injected into normal mice. This difference could be explained by interactions between protein F1-expressing bacteria and cFn.

To further elucidate the relative importance of protein F1 and its binding to pFn and cFn, mice received a bacterial

injection i.p. and were killed 24 h later. For these experiments, AP1(pPTF8ΔBD) bacteria expressing a truncated version of protein F1 lacking the Fn-binding domains (Figure 6A) were also used. Counting the number of bacteria recovered from the spleen revealed a significant difference in dissemination efficiency, depending on the presence of intact protein F1 with retained Fn-binding capacity. The finding that bacterial dissemination is reduced when protein F1 is expressed is in concordance with the challenge experiments (Figure 5A). Furthermore, pFn-null mice infected with protein F1-expressing bacteria had a significantly higher number of bacteria in their spleens than control mice infected with protein F1-expressing bacteria. Thus, pFn bound to *S. pyogenes* limits bacterial dissemination in this model. Increased internalisation, resulting in a local retention of the infection, could explain this observation. The virulence of protein F1-expressing bacteria was only partially restored in pFn-null mice, and in the absence of pFn *S. pyogenes* expressing protein F1 still disseminated less efficiently than bacteria lacking protein F1 or the Fn-binding domains of protein F1. The results obtained with control and cFn-null fibroblasts indicate that this is due to the RD and UFBD domains of protein F1 interacting with cFn. However, binding to cFn did not increase internalisation, suggesting that interactions between bacteria and cFn retain bacteria locally and reduce the number of bacteria that can spread, even though they are not internalised. As mentioned above, phagocytic killing was not influenced by protein F1, which further supports the notion that the reduced virulence of Fn-binding *S. pyogenes* is the result of bacteria being immobilised by host cells and tissues.

In cases of severe *S. pyogenes* infections, including streptococcal toxic shock syndrome (STSS), strains of the M1 serotype are the most frequent (Stevens, 2000). A recent study suggests that the formation of complexes between M1 protein released from the bacterial surface, and fibrinogen, could play a pathogenic role in these severe cases by inducing vascular leakage (Herwald *et al*, 2004). In relation to the present investigation, it is noteworthy that M1 strains commonly lack the protein F1 gene (Natanson *et al*, 1995), indicating that this could also contribute to their virulence. Invasive *S. pyogenes* infections and STSS are connected with high mortality rates, which have led to an increased demand for alternative therapeutic strategies. Specific inhibition of host–bacterial interactions, such as adhesion and internalisation mediated by Fn-binding, has been suggested as a treatment of these serious infections. However, the present finding that expression of protein F1 in a highly virulent M1 strain results in decreased virulence suggests that therapeutic intervention directed toward this protein might actually increase the risk of a fatal outcome. *S. pyogenes* can be regarded as a relatively unsuccessful parasite that can survive and grow only in humans, where it probably also elicits a too powerful inflammatory response (Kotb *et al*, 2002). It could be that this pathogen, by the acquisition of Fn-binding activity, is striving to decrease its virulence and establish a more peaceful and evolutionary advantageous co-existence with its human host.

Materials and methods

Bacterial strains, growth conditions, and plasmids

The *S. pyogenes* AP1 strain is from the AP collection, Institute of Hygiene and Epidemiology, Prague, Czech Republic. Plasmids

pLZ12 (carrying Cm and kanamycin (Km) resistance genes) and pPTF8 (pLZ12 also carrying the protein F1 gene) (Hanski and Caparon, 1992) were purified from DH5 α strains and electroporated into the AP1 strain, thus generating Cm- and Km-resistant clones AP1(pLZ12) and AP1(pPTF8). For generation of AP1(pPTF8 Δ BD), the Fn-binding domains of protein F1 determined previously (Sela *et al*, 1993) were replaced with an HA tag (influenza A virus haemagglutinin epitope) as follows: pPTF8 was used as template in an 'inside-out' PCR reaction with primers, prtF Δ BD-up (CCGGAATTCAGACTACCCGCTAGAGGTGATTGG) and prtF Δ BD-dn (CCGGAATTCACCCATAACGATGTTCAGATTACGCTTTCAGT GAAACGGCGACTGTTG). Digestion of the PCR fragment with *Eco*RI (sites underlined), followed by re-ligation, resulted in an in-frame replacement of the Fn-binding domain of PrtF (G370–G582) with the HA epitope (Y P Y D V P D Y A, the DNA sequence in prtF Δ BD-dn is indicated in bold). The fidelity of the construct was verified by sequencing. The resulting plasmid pPTF8 Δ BD was used to transform *Escherichia coli* (DH5 α) and *S. pyogenes* (AP1), and similar levels of prtF1 gene expression from AP1(pPTF8) and AP1(pPTF8 Δ BD) bacteria were confirmed with Northern blotting using a prtF1-specific probe (CCTGGATATAATATTGGACTCGTTAT CATGACTTGAGAGTAAATTTAAATGGGAGTCCG), hybridising with the 5' part of the gene.

S. pyogenes strains were grown in Todd-Hewitt broth (Difco, Detroit, MI, USA) with 0.2% yeast extract (Difco) (THY) in 5% CO₂ at 37°C. For growth of the transformed strains, 500 μ g/ml of Km and 3 μ g/ml of Cm were added. *E. coli* strains were grown in Luria Bertoni broth (10 g tryptone (Difco), 10 g NaCl, and 5 g yeast extract (Difco)/l), supplemented with 25 μ g Km/ml and 15 μ g Cm/ml.

Binding assays

Binding assays were performed using human Fn (Sigma) radiolabelled with IODO-BEADS (Pierce). Free ¹²⁵I was separated from labelled protein on a PD10 column (Amersham Pharmacia Biotech). In the binding assays, bacteria were grown overnight, washed, resuspended in PBS, and bacterial concentrations were determined by measuring optical density at 620 nm. Bacteria were diluted to appropriate concentrations in PBS with 0.05% Tween 20 and 0.02% azide, and incubated with the radiolabelled protein for 30 min. After centrifugation, the radioactivity of the pellets was determined and expressed as a percentage of added radioactivity, deducing binding of the protein to the polypropylene tubes.

Serum absorption experiments were performed by culturing bacteria overnight, washing once, and incubating 2×10^9 bacteria in 100 μ l of PBS and 100 μ l of serum for 30 min. The bacteria were washed five times with PBS and bound proteins were eluted with 0.1 M glycine-HCl (pH 2.0). The supernatant was collected and neutralised with 1 M Tris. Protein samples were separated by SDS-polyacrylamide gel electrophoresis (PAGE) (Neville, 1971), containing 8 or 12% of acrylamide, and blotted onto an Immobilon-PTM polyvinylidene difluoride filter. The filter was subjected to immunodetection with polyclonal rabbit anti-Fn antibodies diluted 1:5000 (DAKO, Denmark). Horseradish peroxidase-conjugated goat anti-rabbit antibodies (BIO-RAD) were used as secondary antibodies, and immunodetection was achieved by chemiluminescence (Nesbitt and Horton, 1992). Alternatively, the gel was stained using Commassie Brilliant blue or silver staining (Heukeshoven and Dernick, 1988).

Adhesion and internalisation assays

AP1(pLZ12), AP1(pPTF8), or AP1(pPTF8 Δ BD) were grown overnight, washed once in PBS, and diluted in minimal essential medium with Earle's salt (MEM) (Life Technologies) with or without supplemented serum, to a final concentration of approximately 5×10^5 bacteria/ml. Foetal bovine serum (FBS), serum from control mice, or serum from pFn-null mice, was used to supplement MEM in the assays (similar results were obtained with FBS supplementation and with serum from normal mice). Human pharyngeal epithelial Detroit 562 cells (ATCC CCL-138) or human foreskin fibroblasts were grown in 24-well tissue plates (Costar) in MEM supplemented with antibiotics, glutamine, and 10% FBS. Near-confluent monolayers of cells were washed three times in PBS and three times in unsupplemented MEM, to remove antibiotics and serum. Subsequently, 1 ml bacterial suspension was added to each well and the plate was incubated for 2 h with 5% CO₂ at 37°C, followed by five washes with PBS and release of bacteria as described previously (Berry and Paton, 1996). Alternatively, to

determine the number of internalised bacteria, the wells were washed and incubated a further 2 h in MEM supplemented with 100 μ g gentamycin and 5 μ g of penicillin to kill extracellular bacteria, and then washed three times with PBS to remove antibiotics. Bacteria were released as above and viable counts were performed. The number of adherent and internalised bacteria were related to the size of the inoculum, and expressed as a percentage of the number of adherent or internalised AP1(pLZ12) bacteria, respectively. All calculations for statistical significance were carried out using an unpaired Student's *t*-test, two-tailed.

ELISA

Human pharyngeal epithelial Detroit 562 cells or fibroblasts were grown to confluence in microtitre plates (NUNC/NONCLONTM Surface, NUNC, Denmark). The wells were washed six times in PBST and blocked using PBST with 2% (w/v) bovine serum albumin (BSA) for 30 min at 37°C. As a control, wells were coated with purified Fn in different concentrations. Primary rabbit anti-Fn antibodies (Sigma) were applied to the wells in two-fold dilutions (from 1:500) and the plates were incubated at 37°C for 1 h. The wells were washed and horseradish peroxidase-conjugated goat anti-rabbit antibodies (BIO-RAD) were applied. After a 1 h incubation at 37°C, the wells were washed and substrate solution (0.1% diammonium-2,2-azino-bis (3-ethyl-2,3-dihydrobenzthiazoline)-6-sulphonate (ABTS), 0.012% (v/v) H₂O₂ in 100 mM citric acid, 100 mM NaH₂PO₄, pH 4.5) was added. The change in optical density at 415 nm was determined after 10 min.

Generation of fibroblast-like cells lacking fibronectin

To establish fibroblast-like cells, primary cells of E13.5 Fn (flox/flox) mouse embryos were immortalised by the transfection of SV-40 large T antigen, cloned, and several immortalised and clonal fibroblast-like cell lines were generated. Subsequently, two clonal lines were treated with a cre-transducing adenovirus to delete the floxed Fn genes. The deletion of Fn alleles in both cre-treated clones was confirmed by PCR and the lack of Fn protein expression by immunoprecipitation using metabolically labelled conditioned media of these cells.

Animal challenge studies

The construction and characterisation of the pFn-deficient mice was described elsewhere (Sakai *et al*, 2001). Briefly, mice carrying a floxed Fn gene and expressing Cre under control of the interferon and polyinosinic-polycytidic acid (pI-pC)-inducible Mx promoter (Mx-Cre+) were generated. After i.p. injection with pI-pC, pFn levels were undetectable by Western analysis, whereas production of Fn in other tissues was only marginally affected. The pFn-deficient mice showed no overt phenotype or abnormalities up to 8 months of age. The absence of pFn was confirmed by Western analyses in the animals used in this study, and these mice are referred to as pFn-null mice. Mice carrying a floxed Fn gene (Fn(flox/flox)) were used as controls after injection with pI-pC.

Bacteria were grown to early logarithmic growth phase (OD₆₂₀ ~ 0.35), harvested, washed in PBS, diluted in PBS to an appropriate concentration, and kept on ice until injection. Colony counts of the inoculum verified that identical numbers of bacteria were injected. The mice were treated twice daily with 300 μ g Cm (a nontoxic dose) throughout the experiment in order for the bacteria to retain the transfected plasmids. This corresponds to a peak plasma concentration of approximately 3 μ g/ml (Joiner *et al*, 1981). The first Cm injection took place 2 h before injection of the bacteria. The mice were monitored twice daily for 8 days after injection and the time of death for each mouse was recorded. Post mortem, the spleens were removed and homogenised, in order to recover disseminated bacteria. To verify that the bacteria retained their plasmids throughout the infection, the homogenate was plated on THY plates with or without Km and Cm. This was performed on several mice, and on all occasions the number of bacteria was unaffected by the presence of antibiotics. Colonies from plates without antibiotics were reinoculated with antibiotics. This was repeated with bacterial colonies from several different mice, for a total of 100 colonies, and all colonies grew on antibiotic-containing plates. After 8 days, surviving mice were killed. Statistical analyses for differences in the number of dead animals between groups were carried out using Fisher's exact test.

Dissemination to the spleen

Bacteria were grown to early logarithmic phase ($OD_{620} \sim 0.35$), harvested, washed in PBS, diluted in PBS to appropriate concentrations, and kept on ice until injection. After i.p. injection of the bacteria, the mice were killed after 24 h, the spleens were removed, homogenised in 1 ml of PBS, and viable counts were performed. The mice were treated with Cm (300 µg/ml) 2 h prior to injection and 12 h after injection. Statistical analyses for differences between groups were carried out by using the Mann–Whitney *U*-test.

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