Adhesin-dependent binding and uptake of \textit{Salmonella enterica} serovar Typhimurium by dendritic cells

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\textit{Salmonella enterica} serovar Typhimurium can be internalized by immature dendritic cells (DCs). The interacting host and bacterial molecules initiating this process remain uncharacterized. The objective of this study was to investigate whether specific fimbriae are involved in the early step of binding and uptake of \textit{Salmonella} by DCs. Type 1 fimbriated \textit{S. enterica} serovar Typhimurium or recombinant \textit{Escherichia coli} expressing the type 1 fimbriae showed a significantly greater ability to attach to murine bone-marrow-derived DCs than non-fimbriated bacteria. The FimH adhesin was required for efficient interactions with DCs, since fimbriated \textit{fimH} mutants were impaired in both binding and internalization. Finally, the internalization involved a FimH-dependent process but did not require \textit{sipE}, a gene essential for \textit{Salmonella}-mediated invasion of mammalian epithelial cells. Collectively, these data suggest that the bacterial interaction of DCs through the type 1 fimbrial adhesin FimH is sufficient to target \textit{S. enterica} serovar Typhimurium for cellular uptake.

\section*{INTRODUCTION}

\textit{Salmonella enterica} serotypes are correlated with different types of human and animal disease syndromes such as bacteraemia, typhoid fever and enterocolitis. \textit{S. enterica} serotype Typhimurium is a highly evolved pathogen that has adapted to colonizing and surviving for long periods in humans, mice and other animal hosts. The virulence is partially attributed to its capacity to invade host cells.

After oral infection, \textit{S. enterica} serotype Typhimurium colonizes the small intestine and preferentially enters Peyer’s patches M cells (Jepson & Clark, 2001). Thanks to the \textit{Salmonella} type III secretion system (T3SS) encoded by \textit{Salmonella} pathogenicity island 1 (SPI-1) (Galan, 2001; Lostroh & Lee, 2001), \textit{Salmonella} induce their uptake by non-phagocytic cells, such as intestinal epithelial cells. How important epithelial cell invasion is \textit{in vivo} for crossing the intestinal barrier remains questionable, SPI-1 deletion mutants being still enteroinvasive (Murray \& Lee, 2000). SPI-1 facilitates M cell invasion (Penheiter \textit{et al.}, 1997), but is not essential for entry by this route (Clark \textit{et al.}, 1998; Jepson \& Clark, 2001). The invading \textit{Salmonella} cells that have translocated through the intestinal epithelial layer reach the subepithelial compartment where they interact most efficiently with dendritic cells (DCs) that reside under M cells or in M cell pockets (Hopkins \textit{et al.}, 2000). As a third uptake mechanism, CD18-expressing phagocytes were shown to transport \textit{Salmonella} from the gastrointestinal tract to the bloodstream, suggesting a paracellular route of entry (Vazquez-Torres \textit{et al.}, 1999; Vazquez-Torres \& Fang, 2000). Such direct uptake has been visualized with intraepithelial DCs that send tight-junction-forming dendrites to the intestinal lumen (Niess \textit{et al.}, 2005; Rescigno \textit{et al.}, 2001), suggesting that CD18-expressing phagocytes are most probably DCs. DCs are found in the peripheral tissues, mainly as immature DCs that have high phagocytic and endocytic abilities. Immature DCs are widely spread in the lamina propria of the gut and can be recruited at the subepithelial site of infection through proinflammatory molecules, particularly chemokines expressed by the infected enterocytes (Didierlaurent \textit{et al.}, 2002; Sierro \textit{et al.}, 2001). Both host molecules and microbe-associated molecular pattern, such as lipopolysaccharide, peptidoglycan and flagellin, then participate in the maturation process of DCs. Maturing DCs lose their phagocytic properties and upregulate surface molecules, such as chemokine receptor

Abbreviations: DC, dendritic cell; FCS, fetal calf serum; PE, R-phyceroerythrin.
CCR7 that initiates their migration to the mesenteric lymph node (Cheminay et al., 2002), as well as major histocompatibility complex and co-stimulatory molecules (e.g. CD80, CD86 and CD40) that serve to activate naïve T cells (Wick, 2002, 2003). Thus unique maturation steps empower DCs to become most effective antigen-presenting cells in their role as specialized messengers linking innate to adaptive immunity.

Although the uptake of *S. enterica* serotype Typhimurium in DCs remains an intensive field of study, very little is known about the bacterial molecules involved in the earliest recognition steps. *S. enterica* serotype Typhimurium produces a variety of fimbrial adhesins that have been shown to be involved in bacterial binding in murine intestines (van der Velden et al., 1998). Fimbriae are surface-exposed hair-like structures that consist of helically arranged protein subunits. Their adhesive properties are most frequently mediated by a minor tip subunit. The fimbrial subunits are typically encoded by a gene cluster, together with export, assembly and regulatory proteins. The current study investigated whether *Salmonella* fimbriae are involved in the initial interaction between bacteria and DCs. Our data provide evidence that the type 1 fimbriae, thanks to their adhesive subunit FimH, mediate the attachment of *S. enterica* serotype Typhimurium to DCs, leading to a significantly increased SPI-1-independent bacterial internalization process.

### METHODS

**Bacterial strains and plasmids.** *S. enterica* serotype Typhimurium and *Escherichia coli* strains and plasmids used in this study are detailed in Table 1. Bacteria were routinely grown on LB medium supplemented with the appropriate antibiotics. All the transfectants were prepared by high-voltage electroporation. Static broth cultures were performed at 37 °C for 48 h to enrich for type 1 fimbriae. Expression of type 1 fimbriae was determined with antibodies, and expression of FimH and mannose-sensitive binding was screened by yeast cell (*Saccharomyces cerevisiae*) agglutination (Hancock et al., 1998). *Salmonella* type 1 fimbriae were also expressed as recombinant organelles on the surface of *E. coli* strains ORN103 or AAEC189, both lacking their own mannose-sensitive type 1 fimbrial genes (Table 1; designated *fim* or *pil*). *E. coli* ORN103(pKT303) was essentially non-fimbriated, suggesting a polar effect on the transcription of downstream regulatory genes (Table 1). Therefore, a new *fimH* deletion mutant was constructed according to the following strategy. Long PCR was carried out to amplify all the type 1 fimbrial genes encoding the structural and biogenesis proteins (*fimA–fimF*) using pISF101 as the template, according to the manufacturer’s protocol (Expand Long Template PCR; Roche Applied Science). The oligonucleotides were derived from the 5' end of *fimA* and a sequence complementary to 3' end of *fimE*. *NdeI* and *BamHI* sites were introduced into the upper and lower primer, respectively

### Table 1. Strains and plasmids used in this study

<table>
<thead>
<tr>
<th>Strain or plasmid</th>
<th>Parental strain and relevant properties*</th>
<th>Source or reference</th>
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</thead>
<tbody>
<tr>
<td><strong>S. enterica</strong> serotype Typhimurium**</td>
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<tr>
<td>AJB3</td>
<td>p425 (NalR), derived from SR-11</td>
<td>Bäumler et al. (1996a, c); Lockman &amp; Curtiss (1992b)</td>
</tr>
<tr>
<td>AJB4</td>
<td>p4253 (NalR), derived from SR-11 Δfim</td>
<td>Bäumler et al. (1996a, c); Lockman &amp; Curtiss (1992b)</td>
</tr>
<tr>
<td>ST100</td>
<td>Typhimurium C5 * sipB::Tn5 (KmR)</td>
<td>Hermant et al. (1995); Hersh et al. (1999)</td>
</tr>
<tr>
<td>DMS1508</td>
<td>AJB3 * sipB::Tn5</td>
<td>This study</td>
</tr>
<tr>
<td>DMS1509</td>
<td>AJB4 * sipB::Tn5</td>
<td>This study</td>
</tr>
<tr>
<td>LBH4</td>
<td>LB5010 * fimH::KmR, derived from LT2</td>
<td>Hancock et al. (1998)</td>
</tr>
<tr>
<td><strong>E. coli</strong></td>
<td></td>
<td></td>
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<tr>
<td>BL21(DE3)</td>
<td>F *ompT hsdS6 (rK mK) gal dcm (DE3)</td>
<td>Novagen, EMD Biosci.</td>
</tr>
<tr>
<td>SE5000</td>
<td>MC4100 recA (Fim−)</td>
<td>Silhavy et al. (1984)</td>
</tr>
<tr>
<td>ORN103</td>
<td>thr1 let6 thi1 (argF-lac) U169 xyl7 ara13 mtl2 gal6 rpsL *fluA2 minA minB recA13 Δ(pilABCDFE hyp)</td>
<td>Maurer &amp; Orndorff (1987)</td>
</tr>
<tr>
<td>AACEC189</td>
<td>MM294 Δlac recA endA Δfim (rK mK)</td>
<td>Blomfield et al. (1991)</td>
</tr>
<tr>
<td><strong>Plasmids</strong></td>
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<tr>
<td>pISF101</td>
<td>pACYC184 <em>fimAICDFHYZYWU</em> (CmR)</td>
<td>Thankavel et al. (1999)</td>
</tr>
<tr>
<td>pKT303</td>
<td>pISF101 ΔfimH (polar effects on downstream <em>fim</em> genes)</td>
<td>Thankavel et al. (1999)</td>
</tr>
<tr>
<td>pET16b</td>
<td>Expression vector with a T7 promoter</td>
<td>Novagen</td>
</tr>
<tr>
<td>pAZ22</td>
<td>pET16b <em>fimAICDF</em></td>
<td>This study</td>
</tr>
<tr>
<td>pAZ25</td>
<td>pAZ22 ΔfimH (non-polar 33 bp deletion)</td>
<td>This study</td>
</tr>
<tr>
<td>pAZ26</td>
<td>pACYC184 <em>fimAICDF</em></td>
<td>This study</td>
</tr>
<tr>
<td>pISF204</td>
<td><em>fimHEZ</em> in ApR plasmid</td>
<td>Hancock et al. (1998)</td>
</tr>
<tr>
<td>pISF211</td>
<td>pISF204 ΔfimH</td>
<td>Hancock et al. (1998)</td>
</tr>
<tr>
<td>pKKGFP</td>
<td>pKK223.3 GFP (ApR)</td>
<td>Hopkins et al. (2000)</td>
</tr>
<tr>
<td>pGFPMut3.1</td>
<td>pUC GFPMut3.1 (ApR)</td>
<td>BD Biosciences, Clontech</td>
</tr>
<tr>
<td>pAZ8</td>
<td>pLG339 GFPMut3.1 (TcR)</td>
<td>This study</td>
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*A: Ap, Ampicillin; Cm, chloramphenicol; Km, kanamycin; Nal, nalidixic acid; Tc, tetracycline.*
E. coli cloned as a agglutinated yeast cells. The fimbrial gene cluster of pAZ25 was fim and self-ligation to obtain pAZ25. The deletion was confirmed by checking for the loss of an internal CiaI site. Plasmid pAZ24 or pAZ25 was introduced into BL21(DE3) by electroporation, and fim and fimH gene expression was checked after IPTG induction. Both transformants produced type 1 fimbriae, but only BL21(DE3)[pAZ25] agglutinated yeast cells. The fimbrial gene cluster of pAZ25 was cloned as a BglII–BamHI DNA fragment into the BamHI site of pACYC184, in the opposite orientation to the tetracycline resistance gene, to obtain pAZ36. E. coli AEIC189(pAZ36) was shown to express fimbriae constitutively without agglutinating yeast cells.

**Preparation of antiserum to type 1 fimbriae.** The *Salmonella* type 1 fimbriae expressed by *E. coli* SE5000(pJS101) were purified as described by Dodd & Eisenstein (1982). SDS-PAGE and Coomassie blue staining confirmed the preparation’s purity. Antifimbrial antibodies were raised in rabbits (Cocalico Biologicals) with subsequent serum ELISA titres of fimbrial antibodies were raised in rabbits (Cocalico Biologicals) with subsequent serum ELISA titres of 1 : 640. Flow cytometry binding and uptake assays. Bacterial and fimbrial adhesion to DCs, as well as bacterial uptake by DCs, was examined by flow cytometry. Bacteria transformed with a GFP-expressing plasmid and other appropriate compatible plasmids (Table 1) were grown statically. Initial experiments were undertaken with high-copy-number pKKGFP-transformed *S. enterica* serotype Typhimurium. However, the percentage of fluorescent bacteria grown statically for 2 days was frequently under 50 %, suggesting plasmid loss. Thus, later experiments took advantage of the low-copy-number and stable plasmid construct pAZ8. Although the fluorescence intensity of *Salmonella*-associated DCs was lower with pAZ8- than with pKKGFP-containing bacteria, all bacteria associated to DCs were detectable. Bacteria were collected by centrifugation, washed and resuspended in 4 % PBS. Bacterial numbers were estimated by OD600 and checked by colony counts (c.f.u. ml⁻¹). DCs were harvested and cell viability (≥90 %) was tested by Trypan Blue staining. Bacteria and DCs were incubated at 4 °C for various times at an m.o.i. of 50 with gentle end-to-end agitation. Unbound bacteria were removed from DCs by three to five washes in PBS. DCs were further incubated with anti-CD11c-PE antibodies for 30–60 min in the dark at 4 °C with gentle end-to-end agitation. After three to five washing steps, cells were resuspended in 300–500 µl cold RPMI 1640 for flow cytometry (FACScan or FACS Calibur with CellQuest pro software; BD Biosciences). For the binding inhibition assay, bacteria were incubated with DCs in the presence of 100 mM 2-methyl mannose for 1 h. To study the binding of type 1 fimbriae to DCs, purified fimbriae of strain AJB3 were biotinylated with sulfo-NHS-LC-Biotin (Pierce), according to the manufacturer’s instructions. Biotinylated fimbriae were incubated in 2 % FCS-PBS with DCs at 4 °C for 1 h. PE-conjugated streptavidin was used to probe the biotinylated fimbriae that remained attached to DCs after three to five washes.

The uptake of bacteria by DCs was also examined by flow cytometry. Bacteria and DCs were incubated for 30 min at 37 °C on a rotating wheel (m.o.i. = 50), washed three to five times and the cells fixed with 4 % paraformaldehyde. The cells were then divided into two groups, one group being permeabilized with 0.5 % Triton X-100 at room temperature for 10 min. Cells were then labelled with rabbit anti-*Salmonella* O antigen factor 4 antiserum (Dífco, BD Diagnostic Systems), sequentially followed by biotin-conjugated goat anti-rabbit IgG and FITC-conjugated streptavidin or Alexa Fluor488 anti-rabbit IgG (Invitrogen, Molecular Probes). The DCs were probed with PE-conjugated anti-CD11c IgG. The non-permeabilized cells were used to by washing three times with RPMI1640 medium containing 50 µg gentamicin ml⁻¹. The cells were further incubated for 1 h in RPMI1640 medium containing 50 µg ml⁻¹ gentamicin at 37 °C. DCs were then washed three times with PBS, lysed with 0.5 % Triton X-100 in PBS for 10 min and live *Salmonella* counted by a standard plating technique. The colony numbers for the wild-type strain AJB3 were normalized to 100 % for comparisons with the numbers of the other *Salmonella* strains. The human HeLa epithelial cell line (ATCC CCL-2) was cultured in Dulbecco’s modified Eagle medium (Gibco) supplemented with 10 % FCS and 1 mM non-essential amino acids (Gibco) at 37 °C in an atmosphere containing 5 % CO2. Monolayers were prepared by seeding 2 x 10⁵ cells into each well of a 24-well tissue culture plate and incubating for 48 h at 37 °C in a humidified atmosphere of 5 % CO2 in air. Bacteria were added to the cells at an m.o.i. of 20 : 1 to 200 : 1. Bacteria were incubated with cell monolayers for 1 h at 37 °C in 5 % CO2 before being washed three times with PBS. Extracellular bacteria were killed by incubating the cells in medium containing 50 µg gentamicin ml⁻¹ for an additional 90 min. After three PBS-washing steps, the cell monolayers were lysed with cold 0.1 % Triton X-100 and viable counts were determined as described above.

**Flow cytometry binding and uptake assays.** Bacterial and fimbrial adhesion to DCs, as well as bacterial uptake by DCs, was examined by flow cytometry. Bacteria transformed with a GFP-expressing plasmid and other appropriate compatible plasmids (Table 1) were grown statically. Initial experiments were undertaken with high-copy-number pKKGFP-transformed *S. enterica* serotype Typhimurium. However, the percentage of fluorescent bacteria grown statically for 2 days was frequently under 50 %, suggesting plasmid loss. Thus, later experiments took advantage of the low-copy-number and stable plasmid construct pAZ8. Although the fluorescence intensity of *Salmonella*-associated DCs was lower with pAZ8- than with pKKGFP-containing bacteria, all bacteria associated to DCs were detectable. Bacteria were collected by centrifugation, washed and resuspended in 4 % PBS. Bacterial numbers were estimated by OD600 and checked by colony counts (c.f.u. ml⁻¹). DCs were harvested and cell viability (≥90 %) was tested by Trypan Blue staining. Bacteria and DCs were incubated at 4 °C for various times at an m.o.i. of 50 with gentle end-to-end agitation. Unbound bacteria were removed from DCs by three to five washes in PBS. DCs were further incubated with anti-CD11c-PE antibodies for 30–60 min in the dark at 4 °C with gentle end-to-end agitation. After three to five washing steps, cells were resuspended in 300–500 µl cold RPMI 1640 for flow cytometry (FACScan or FACS Calibur with CellQuest pro software; BD Biosciences). For the binding inhibition assay, bacteria were incubated with DCs in the presence of 100 mM 2-methyl mannose for 1 h. To study the binding of type 1 fimbriae to DCs, purified fimbriae of strain AJB3 were biotinylated with sulfo-NHS-LC-Biotin (Pierce), according to the manufacturer’s instructions. Biotinylated fimbriae were incubated in 2 % FCS-PBS with DCs at 4 °C for 1 h. PE-conjugated streptavidin was used to probe the biotinylated fimbriae that remained attached to DCs after three to five washes.
determine the extracellular bound bacteria, while the permeabilized cells were interpreted as representing all the DC-associated bacteria, namely both the extra- and intracellular Salmonella.

**Statistical analysis.** Groups of data were compared with the unpaired Student’s *t*-test, using an unequal variance *t*-test when needed, as determined by the F-test (Zar, 1974). Probability (*P*) values of less than 0.05 indicated that the groups were significantly different.

**RESULTS**

**Adhesion of type 1 fimbriated *S. enterica* serotype Typhimurium to DCs**

To characterize interaction of DCs with bacteria, cells were incubated at 37 °C, followed by extensive washing, Giemsa staining and analysis by light microscopy. *S. enterica* serotype Typhimurium grown statically interacted better with DCs than bacteria grown overnight or in exponential growth phase under aerated conditions (data not shown). Since *Salmonella* produce large amounts of type 1 fimbriae when grown statically, we hypothesized that the type 1 fimbriae were involved in bacterial association to DCs. The interactions of *S. enterica* serotype Typhimurium AJB3 and its non-fimbriated *fim* derivative AJB4 with DCs were compared over time (Fig. 1). The association of wild-type bacteria with DCs was low or not detectable at 5 min and increased gradually over time. The interaction of the fimbriated bacteria was about three- and fivefold more effective at 30 and 60 min, respectively, compared to the *fim* mutant (Fig. 1b). Type 1 fimbriated Salmonella bound significantly better to cytochalasin D-treated DCs at 30 min [0.72 ± 0.10 *Salmonella* *Fim* + per DC versus 0.10 ± 0.03 *Salmonella* *Fim* − per DC (means ± SE, *P* < 0.01)] and 60 min (2.97 ± 0.54 *Salmonella* *Fim* + versus 0.20 ± 0.06 *Salmonella* *Fim* −, *P* < 0.01), indicating that the interaction was independent of actin polymerization and phagocytosis. To specifically assess binding of bacteria to DC, flow cytometry assays were undertaken at 4 °C. Treatment at 4 °C impairs any internalization events and was observed to slow down DC aggregation. Bacteria-associated DCs were identified by using GFP-expressing bacteria and PE-anti-CD11c-labelled DCs. Moreover, to exclude any effect of the *Salmonella* type III secretion system SPI-1 in attachment, we constructed fimbriated and non-fimbriated bacteria deficient for SipB. After a 30 min bacteria/DC incubation, the type 1 fimbriated strain associated with most DCs (92 %), whereas no significant binding was detected with the non-fimbriated bacteria (Fig. 1c). Taken together, the flow cytometry data supported the microscopic observation that *S. enterica* serotype Typhimurium type 1 fimbriae are critical for bacterial adhesion to DCs.

**Fimbriae-specific binding to DCs**

To confirm that the binding of bacteria to DCs was mediated by type 1 fimbriae, experiments were performed with an *E. coli* *fim* operon deletion mutant (ORN103) that was complemented with either plasmid pISF101 harbouring *Salmonella* type 1 fimbrial genes or pKT303, which is

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**Fig. 1.** Type 1 fimbriae-dependent binding of *S. enterica* serotype Typhimurium to DCs. (a) Giemsa-stained DCs and *S. enterica* serotype Typhimurium AJB3 and AJB4 (*fim*) incubated at 37 °C for 60 min at an m.o.i. of 100. Bars, 40 μm. (b) Bacteria were incubated with DCs at 37 °C for 5, 30 and 60 min. The numbers of bacteria associated with DCs were counted microscopically. The bars represent the mean ± SE values for AJB3 (filled bars) and AJB4 (shaded bars) of three experiments. There were statistically significant differences for the 30 and 60 min values (*P* < 0.01). (c) Binding of type 1 fimbrial *S. enterica* serotype Typhimurium DMS1508 (*sipB*). Bacteria were transformed with the GFP-expressing plasmid pAZ8. Only CD11c + -gated cells (DCs) are shown. DCs and bacteria were incubated for 30 min at an m.o.i. of 50 at 4 °C. The histograms represent control non-infected DCs (dotted line), DCs with DMS1509(pAZ8) (*fim sipB*) (thin grey line) or DMS1508(pAZ8) (*sipB*) (thick black line). The data are from one experiment and represent reproducible data from three experiments.
pISF101 with a deletion that eliminates fimbriation. *E. coli* expressing the *Salmonella* type 1 fimbriae interacted significantly better with DCs than the non-fimbriated *E. coli* after 30 or 60 min incubation at 37 °C (Fig. 2a). Binding assays with GFP-expressing *E. coli* showed that 93% of the DCs were associated with fimbriated *E. coli* (Fig. 2b). In contrast, DCs incubated with *E. coli* (pKT303) did not bind any bacteria (Fig. 2b). Biotinylated purified type 1 fimbriae were found to bind in a dose-dependent manner to DCs (data not shown). Altogether, type 1 fimbrial expression was needed for efficient bacterial adhesion to the DCs.

**FimH-dependent binding to DCs**

The type 1 fimbriae of *Salmonella* bind to mannosylated glycoproteins on mammalian cells, and this interaction is mediated by their minor adhesive subunit FimH (Hancox et al., 1998). To determine whether the adhesin FimH mediated bacterial binding to DCs, *fimH* mutants producing wild-type strain levels of fimbriae were studied. Whether *S. enterica* serotype Typhimurium *fimH* bound to DCs was evaluated by flow cytometry. As described by Hancox et al. (1998), *S. enterica* serotype Typhimurium LBH4, which has an insertional inactivated *fimH* gene, produced type 1 fimbriae and agglutinated yeast cells only when complemented with pISF204 carrying *fimHFZ*. When complemented with pISF211, which contains *fimFZ*, only type 1 fimbriation was restored. Most of LBH4(pISF204) bound to DCs, as shown by flow cytometry, whereas the binding of LBH4 or LBH4(pISF211) was not significant and similar to the bacteria-free control (Fig. 3, top panel). The role of FimH in recognizing DC was also investigated with recombinant *E. coli* AAEC189(*fim*) complemented with plasmids pISF101(*fim*+) or pAZ36(*fimH*). Both *E. coli* AAEC189(pISF101) and AAEC189(pAZ36) produced similar amounts of type 1 fimbriae, but only the former bacteria agglutinated yeast cells and bound significantly to DCs, as shown in Fig. 3 (bottom panel). Moreover, the binding of type 1 fimbriated *E. coli* ORN103(pISF101) (Fig. 4) or *S. enterica* serotype Typhimurium AJB3 (data not shown) to DCs was impaired by α-methyl mannoside. Nearly the whole population of DCs (93%) associated with *E. coli* ORN103(pISF101) in the absence of the inhibitor, whereas most DCs (76%) were free of bacteria in the presence of α-methyl mannoside. This result demonstrated that α-methyl mannoside efficiently blocks fimbriae-mediated bacterial binding to DCs, and thus that the type 1 fimbriae of *S. enterica* serotype Typhimurium most probably recognize one or more mannosylated glycoprotein receptor(s) on the DCs. This result also confirmed that the mannose-specific adhesin FimH was needed for DC recognition by *Salmonella* type 1 fimbriae.

**Fimbriae-mediated uptake by DCs**

We investigated whether type 1 fimbriae-dependent binding of *Salmonella* to DCs influences intracellular uptake. Both strains AJB3 and AJB4(*fim*) express SPI-1 T3SS that mediates cellular uptake by mammalian cells. To determine whether SPI-1 was involved in internalization by DCs, a *sipB* mutant that expressed type 1 fimbriae (DMS1508) was included in the study. Standard invasion assays (gentamicin...
resistance) indicated that in comparison to the wild-type strain AJB3 (relative invasion of 100%), significantly fewer AJB4 (fim) cells were taken up by the DCs (5.5 ± 4.3% relative to AJB3, mean ± SE of three independent experiments). In contrast, the uptake of DMS1508 (sipB) (67 ± 6% of AJB3 invasion) was significantly less affected than the uptake of AJB4. These results indicated that SPI-1 does not affect the uptake of *Salmonella* by DCs as much as the type 1 fimbriae do. In contrast, DMS1508 (sipB) was internalized significantly less efficiently by HeLa cells when compared to strain AJB3 (2.7 ± 0.6% of AJB3 invasion), suggesting that *Salmonella* invasion of epithelial cells relied more on SPI-1 than *Salmonella* invasion of DCs. Strain AJB4 (fim) was also significantly less invasive with HeLa cells (0.6 ± 0.3% of AJB3 invasion). This result was anticipated, since AJB4 (fim) had been previously shown to be significantly less adhesive and, as a consequence, less invasive than its parental strain AJB3 (Bäumler *et al.*, 1996c). The cellular uptake of *Salmonella* was further analysed by flow cytometry using O-antigen-specific antibody to label the bacteria and DC permeabilization to track the bacterial localization. Indeed, when DC membranes were permeabilized, both extra- and intracellular bacteria were detected by the antigen-specific antibody. The data are presented in Table 2 as the means of

Fig. 3. FimH determines *S. enterica* serotype Typhimurium binding to DCs. DCs and bacteria were incubated at 4°C for 30 min, at an m.o.i. of 50. Only CD11c<sup>+</sup>-gated cells (DCs) are shown. Top panel: complementation of *S. enterica* serotype Typhimurium LBH4 (fimH) with plasmids carrying (pISF204) or lacking (pISF211) fimH. Bacteria were transformed with GFP-expressing plasmid pAZ8. The histograms represent control non-infected DCs (dotted line), or DCs with LBH4(pAZ8) (thin black line), LBH4(pISF211, pAZ8) (grey line) or LBH4(pISF204, pAZ8) (thick black line). The former three histograms overlapped. Bottom panel: recombinant *E. coli* AAEC189 expressing non-adhesive or adhesive *S. enterica* serotype Typhimurium type 1 fimbriae from plasmid pAZ36 or pISF101, respectively. The histograms represent control non-infected DCs (dotted line), DCs with *E. coli* AAEC189(pAZ36, pAZ8) (thin grey line) or DCs with AAEC189(pISF101, pAZ8) (thick black line).

Fig. 4. Inhibition of FimH-mediated bacterial adhesion to DCs with α-methyl mannoside. Bacteria were transformed with the GFP-expressing high-copy-number plasmid pKKGFP. DCs and bacteria were incubated as described in the legend to Fig. 3. The histograms represent control DCs stained with the appropriate isotype subclass (dotted line), or DCs incubated with recombinant *E. coli* ORN103 expressing adhesive *S. enterica* serotype Typhimurium type 1 fimbriae from plasmid pISF101, in the presence (grey thin line) or absence (black thick line) of 100 mM α-methyl mannoside.
four experiments, one typical experiment being shown in Fig. 5. It should be noted that all the presented data are based on apparent numbers, since DCs associated with one (or a few) bacterium might have escaped detection. Among the DCs that were associated with the fimbriated \textit{Salmonella} AJB3 (66.7\% of all DCs), a majority contained solely intracellular bacteria (Table 2). In contrast, only 10.4\% of DCs were associated with the non-fimbriated \textit{Salmonella} AJB4. There were also fewer DCs with only intracellular AJB4 bacteria (5.1\% of all DCs). Most DCs (71.7\%) were associated with the fimbriated \textit{sipB} mutant DMS1508, and a majority of these DCs carried \textit{Salmonella} exclusively intracellularly. DC association of the non-fimbriated \textit{fim} \textit{sipB} double mutant (DMS1509) was similar to that found for the non-fimbriated \textit{fim} strain AJB4 (data not shown). Taken together, these data showed that type 1 fimbriae-mediated \textit{Salmonella} association with DCs was the major determining factor for bacterial internalization.

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\begin{tabular}{|l|c|c|c|c|}
\hline
Strain & Total DCs counted* & Total DCs with associated \textit{Salmonella} (%) & DCs with extracellular \textit{Salmonella} (%) & DCs with intracellular \textit{Salmonella} (%) \\
\hline
AJB3 & 23 600 & 66.7 ± 10.4 (\(P < 0.01\))§ & 22.5 ± 4.9 (\(P < 0.05\))§ & 44.2 ± 8.7 (\(P < 0.01\))§ \\
DMS1508 & 26 945 & 71.7 ± 3.1 (\(P < 0.0001\))§ & 30.4 ± 5.1 (\(P < 0.05\))§ & 41.3 ± 3.4 (\(P < 0.0001\))§ \\
AJB4 & 28 228 & 10.4 ± 2.4 & 5.3 ± 3.0 & 5.1 ± 0.9 \\
\hline
\end{tabular}
\caption{Apparent percentage of DCs with intra- and extracellular \textit{Salmonella} (flow cytometry data)}
\end{table}

\*DCs defined as cells positive for expression of CD11c.
\‡Permeabilized DCs with either internal, external, or both internal and external \textit{Salmonella}.
\§DCs carrying only intracellular \textit{Salmonella} (calculated as associated minus extracellular bacteria).
\¶\textit{t}-Test comparisons with AJB4 data; there were no statistically significant differences between the AJB3 and DMS1508 data.

\begin{figure}
\centering
\includegraphics[width=\textwidth]{Fig5.png}
\caption{Type 1 fimbriae-dependent internalization of \textit{S. enterica} serotype Typhimurium by DCs. Strain AJB3 (c, d), AJB4 (\textit{fim}) (e, f) or DMS1508 (\textit{sipB}) (g, h) were incubated with DCs for 30 min at an m.o.i. of 50 (37 °C). (a) and (b) represent mock-infected DCs. The DCs either were (b, d, f, h) or were not (a, c, e, g) permeabilized. Bacteria were labelled with anti-O antigen antibodies, followed by FITC-conjugated secondary antibodies. Only CD11c\(^{+}\)-gated cells (DCs) are shown. The percentages represent the ratios of DCs carrying extracellular bacteria (a, c, e, g) or both extracellular and intracellular bacteria (b, d, f, h). The data are from one experiment and represent reproducible data from four experiments.}
\end{figure}
DISCUSSION

Following oral ingestion, *S. enterica* serotype Typhimurium reaches the intestines, where it first makes contact with M cells, enterocytes and DC. There is evidence that DCs sending luminal dendrites or DCs within Peyer’s patches interact closely with *Salmonella* (Niess *et al.*, 2003; Rescigno *et al.*, 2001; Salazar-Gonzalez *et al.*, 2006). Although studies have identified that specific fimbrae preferentially target the *Salmonella* towards M cells or intestinal epithelial cells (Bäumler *et al.*, 1996a, b), the bacterial ligand initiating the interaction with DCs is not known. This study showed that the type 1 fimbiae are extremely efficient at mediating the binding and internalization of *S. enterica* serotype Typhimurium into murine DCs. Whether binding directs bacterial uptake by creating the necessary conditions for other interactions to occur or by triggering bacterial internalization itself remains to be determined.

Microscopic analysis and flow cytometry showed that type 1 fimbriated *S. enterica* serotype Typhimurium, in contrast to an isogenic *fim* mutant, binds to bone-marrow-derived DCs, suggesting specificity of the fimbiae for one or more cell receptor(s). The binding specificity was confirmed with recombinant *E. coli* expressing *S. enterica* serotype Typhimurium type 1 fimbiae and with isolated fimbiae, which bound to DCs in a dose-dependent manner. The fimbral structure consists of different types of subunits, all being encoded on the *fim* gene cluster, together with the chaperone-usher system that coordinates the export and assembly of the subunits (Clegg & Swenson, 1994). The helically arranged structural subunit FimA forms most of the polymerized fimbral shaft, whereas the binding determinant for mannosylated proteins is a minor structural subunit designated FimH (Hancox *et al.*, 1998). Here, type 1 fimbiae-mediated *Salmonella* binding to DCs was shown to be mediated by the FimH adhesin. Moreover, this interaction was dependent on mannose residues. Strikingly, FimH-mediated binding was also required for efficient bacterial internalization. Under our experimental conditions, the SPI-1-encoded mammalian cell invasion machinery, did not significantly affect *S. enterica* serotype Typhimurium binding to DCs and was dispensable for DC invasion.

Type 1 fimbiae have previously been shown to confer *S. enterica* serotype Typhimurium adhesion to a variety of mammalian epithelial cells such as human HEP-2, HeLa, and small intestinal and bladder epithelial cell lines (Bäumler *et al.*, 1996c; Boddicker *et al.*, 2002; Hancox *et al.*, 1998; Thankavel *et al.*, 1999), and to contribute to the bacterial invasion of some of these cells (Bäumler *et al.*, 1997; Ernst *et al.*, 1990; Horiuchi *et al.*, 1992; van der Velden *et al.*, 1998). More recently, *S. enterica* serotype Typhimurium colonization of the murine intestine by biofilm formation was shown to require type 1 fimbiae carrying an adhesive allelic FimH (Boddicker *et al.*, 2002). However, unlike epithelial cell internalization, which requires an active bacterial invasion machinery, the observed uptake of *S. enterica* serotype Typhimurium by DC was SPI-1-independent, as described recently by others (Kiama *et al.*, 2006; Petrovská *et al.*, 2004; van der Velden *et al.*, 2003). A recent in vitro study with a macrophage cell line showed that the route of entry of SPI-1-induced, non-opsonized and complement- or IgG-opsonized *Salmonella* influenced their vacuolar surroundings and intracellular gene expression (Drecktrah *et al.*, 2006). *Salmonella* uptake by murine DCs is greatly enhanced by serum complement (Eriksson *et al.*, 2003). What mediated *Salmonella* binding and uptake remained unknown. The novel finding of this study is that the type 1 fimbiae-mediated binding of *S. enterica* serotype Typhimurium to DCs concomitantly targets the bacteria to the intracellular compartment of these professional migratory cells. The potential effect of fimbral lectins on the phagocytic process of DCs was not evaluated in other studies (Drecktrah *et al.*, 2006; Eriksson *et al.*, 2003; Petrovská *et al.*, 2004; van der Velden *et al.*, 2003). DCs are known to recognize and take up bacteria and bacterial molecules via different groups of receptors, including C-type lectin and pattern recognition receptors (Geijtenbeek *et al.*, 2004). One or more unidentified mannosylated DC glycoprotein(s) might represent a new class of phagocytic DC receptors that interact with bacterial lectins, such as the type 1 fimbiae of *Salmonella*. Whether this lectinophagocytic interaction (Ofek & Sharon, 1988) directs uptake by a distinct mechanism of bacterial entry and triggers the formation of a new vacuolar compartment, as described for type 1 fimbriated *E. coli* with mast cells or macrophages (Baorto *et al.*, 1997; Shin *et al.*, 2000), is a question that requires further studies.

SPI-1 gene products are involved in the invasion of intestinal epithelial cells (Galan, 2001) and participate in M cell uptake (Penheiter *et al.*, 1997). However, studies have suggested that SPI-1 is not essential for crossing the intestinal barrier in vivo (Clark *et al.*, 1998; Jeppson & Clark, 2001; Murray & Lee, 2000), and variable use of the SPI-1 proteins has been observed between host species and *Salmonella* serotypes (Morgan *et al.*, 2004). Here, *Salmonella* uptake by murine DCs was mainly determined by FimH-mediated processes and not by its SPI-1-mediated invasive attributes. Although SPI-1 gene expression might not be optimal under the growth conditions used, recent microarray data showed that SipB and other SPI-1 genes are transcribed in stationary phase (Adkins *et al.*, 2006). Moreover, since the SPI-1 apparatus and plasma membrane ruffling are activated within 2 min of macrophage exposure to *S. enterica* serotype Typhimurium grown to stationary phase (Alpuche-Aranda *et al.*, 1994), it is unlikely that our interpretation of the data was significantly affected by the experimental conditions used for the adhesion assays presented. Nevertheless, it remains possible that SPI-1 plays some role in *Salmonella* uptake in vivo.

That *Salmonella enterica* serotype Typhimurium express their type 1 fimbiae in vivo was clearly shown in an earlier study which compared the bacterial colonization of murine...
organisms two days after oral administration of a wild-type strain together with its isogenic fim mutant (Lockman & Curtiss, III, 1992a). Higher numbers of the former strain were isolated from reticulo-endothelial tissues such as the spleen and liver. In contrast, more fim mutants were found in the blood. This observation is consistent with the more efficient uptake of fim+ Salmonella by DCs, together with the previously described extraintestinal systemic distribution of Salmonella by CD18-expressing phagocytes (Vazquez-Torres et al., 1999) that were later postulated to be mainly DCs (Niess et al., 2005; Rescigno et al., 2001). In spite of the increased uptake of type 1 fimbriated S. enterica serotype Typhimurium by DCs, such strains were previously found to be less virulent than isogenic fim mutants in mice, as determined by their LD50 (Lockman & Curtiss, III, 1992b; van der Velden et al., 1998). How the type 1 fimbriae attenuate S. enterica serotype Typhimurium in mice is not clear. Whether the targeting of Salmonella towards DCs amplifies an antibacterial inflammatory response in vivo, as expected from in vitro studies (Marriott et al., 1999; Yrlid et al., 2000), or whether S. enterica serotype Typhimurium-activated resident DCs induce a non-inflammatory response with IL-10 secretion in mice, as observed with epithelial-cell-conditioned human DCs in vitro remains to be determined (Rimoldi et al., 2005). Moreover, DC targeting is expected to affect the adaptive immune response (Buono et al., 2005). In any case, the amplified interaction of type 1 fimbriated S. enterica serotype Typhimurium with DCs might play an important role in attenuating Salmonella, as observed with isogenic fim” and fim” strains (Lockman & Curtiss, III, 1992b; van der Velden et al., 1998). By acting as ‘anti-virulence’ factors (Foreman-Wykert & Miller, 2003), type 1 fimbriae might be beneficial to the intestinal establishment of S. enterica serotype Typhimurium as a persistent inhabitant of immunocompetent adult mice or carriers.

In this study, we showed that the expression of type 1 fimbriae optimized Salmonella uptake by DCs. Efficient bacterial internalization depended on the mannose-specific adhesin FimH and did not require SipB, an essential gene of the T3SS of Salmonella involved in epithelial cell invasion. The role of type 1 fimbriae-mediated binding and uptake of Salmonella by DCs and the importance of these interactions for infection remain to be investigated in the mammalian host.

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