

The normal chain length distribution of the O antigen is required for the interaction of *Shigella flexneri* 2a with polarized Caco-2 cells

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ABSTRACT

Shigella flexneri causes bacillary dysentery in humans. Essential to the establishment of the disease is the invasion of the colonic epithelial cells. Here we investigated the role of the lipopolysaccharide (LPS) O antigen in the ability of *S. flexneri* to adhere to and invade polarized Caco-2 cells. The *S. flexneri* 2a O antigen has two preferred chain lengths: a short O antigen (S-OAg) regulated by the WzzB protein and a very long O antigen (VL-OAg) regulated by Wzz_{pHS2}. Mutants with defined deletions of the genes required for O-antigen assembly and polymerization were constructed and assayed for their abilities to adhere to and enter cultured epithelial cells. The results show that both VL- and S-OAg are required for invasion through the basolateral cell membrane. In contrast, the absence of O antigen does not impair adhesion. Purified LPS does not act as a competitor for the invasion of Caco-2 cells by the wild-type strain, suggesting that LPS is not directly involved in the internalization process by epithelial cells.

Key words: Shigella, epithelial cells, O antigen

INTRODUCTION

Species of the genus Shigella cause bacillary dysentery in humans, an acute intestinal infection characterized by bloody diarrhoea due to an intense inflammatory reaction accompanied by destruction of the colonic mucosa (Schroeder and Hilbi, 2008). An essential step to establish the disease is the invasion of the intestinal epithelial cells followed by bacterial multiplication and spread into adjacent cells. In the initial phase of infection, Shigella is transcytosed across M cells reaching the submucosa, and then free bacteria invade epithelial cells from the basolateral side (Schroeder and Hilbi, 2008). The invasive capacity of Shigella depends on proteins encoded in the large virulence plasmid (Parsot, 2009). This region contains more than 30 genes organized into two clusters: the mxi-spa locus that encodes the components of a type three secretion system (T3SS), and the ipaBCDA locus that encodes effector proteins that are translocated through the T3SS and cause cytoskeletal rearrangements that promote bacterial ingestion (Parsot, 2009).

The expression of *Shigella* virulence genes is controlled by environmental conditions prevalent in the large intestine (Dorman and Porter, 1998); however, full activation of the secretion system is triggered upon contact of *Shigella* with the host cells (Parsot, 2009). Thus, early interactions of bacteria with epithelial cells must take place before invasion can occur. Nonetheless, the nature of these interactions is not completely understood.

Lipopolysaccharide (LPS) has been implicated in the adherence and/or invasion of epithelial cells by several Gramnegative bacteria (Hoare et al., 2006; Kanipes et al., 2004; Klena et al., 2005; Zaidi et al., 1996). LPS consists of the inner lipid A portion, the core oligosaccharide, and the O polysaccharide or

O antigen (OAg), which is exposed to the cell surface (Raetz and Whitfield, 2002). The LPS of *S. flexneri* 2a has O-antigen molecules with two preferred chain lengths, a short or S-OAg composed on average of 17 repeated units (RU) that is regulated by a chromosomally-encoded WzzB protein, and a very long or VL-OAg of about 90 RU that is regulated by Wzz_{pHS2}, which is encoded in plasmid pHS2.

The role of LPS in the invasion process of Shigella is unclear. Early reports using spontaneous or transposoninsertion mutants lacking O antigen or with truncated cores indicated that neither a complete core nor the O antigen is required for S. flexneri entry into HeLa cells (Okamura et al., 1983; Okamura and Nakaya, 1977; Rajakumar et al., 1994; Sandlin et al., 1995). More recently, a study conducted with mutants that were identified by signature-tagged mutagenesis indicated that a strain lacking O antigen was able to invade HeLa cells; however that work also showed that glycosylation of the O antigen enhances bacterial invasion, suggesting that a polymeric O antigen is necessary to this process (West et al., 2005). Only one study has addressed the role of LPS in the interaction of S. flexneri with polarized epithelial cells. Using mutants obtained by Tn10 insertion, Köhler et al. (2002) found that adhesion to polarized human T84 colonic epithelial cells requires the presence of polymerized O antigen.

In this study, we have investigated the role of the O antigen in the ability of *S. flexneri* to adhere to and invade polarized Caco-2 cells, using a series of mutants with defined, non-polar deletions of the genes required for the assembly and polymerization of the O antigen. Our results indicate that polymerized O antigen is necessary for internalization and, moreover, a normal chain length distribution of the O antigen is required for the invasive process.

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MATERIAL AND METHODS

Bacterial strains, plasmids, media and growth conditions

Table I summarizes the properties of the bacterial strains and plasmids used in this study. Bacteria were grown in Luria-Bertani medium (LB) (10 g/L Bacto tryptone, 5 g/L Bacto yeast extract, 5 g/L NaCl). Solid cultures were done in tryptic soy agar (TSA, Difco) containing 0.01% (w/v) Congo red (TSA-CR) at 30 °C to ensure the presence of the virulence plasmid in the constructed strains (Schuch and Maurelli, 1997). Ampicillin (100 μ g/mL), kanamycin (50 μ g/mL), and tetracycline (10 μ g/mL) were added when appropriate.

Mutagenesis of LPS biosynthesis genes

Mutagenesis was performed by the Red/Swap method to create chromosomal mutations by homologous recombination using PCR products (Datsenko and Wanner, 2000). Shigella flexneri 2a strain 2457T containing pKD46, which expresses the λ Red recombinase system, was transformed with PCR products that were generated using as template plasmid pKD4, which contains the FRT-flanked kanamycin-resistance gene (aph). Each primer pair also carried 30 bases that were homologous to the edge of the gene targeted for disruption. The sequences of the oligonucleotide primers used in this study are available on request. The kanamycin-resistant transformants were replica plated in the absence of antibiotic selection at 37 °C and finally assayed for ampicillin sensitivity to confirm loss of pKD46. To obtain non-polar deletion mutants, the antibiotic resistance gene was removed by transforming the gene replacement mutants with pCP20, which encodes the FLP recombinase that mediates site specific recombination between the FRT sites

(Cherepanov and Wackernagel, 1995). Correct insertional gene replacements and the deletion of the antibiotic gene cassettes were confirmed by PCR.

Cloning of the wzzB and wzz $_{pHS2}$ genes

DNA fragments containing the S. flexneri 2457T $wzz_{\rm pHS2}$ (NC_002773) and wzzB (GeneID:1077593) genes and their promoter regions were amplified by PCR. The amplicons were cloned into pDA12 (Aubert et al., 2008). The sequences of the primers used for the cloning experiments are available upon request.

LPS purification

Large scale LPS preparations from *S. flexneri* wild type and the O-antigen mutants wzz_{pHS2} ::aph and wzzB were obtained by the method of Darveau and Hancock (1983). The quality of the purified LPS samples was confirmed by Tricine-SDS-PAGE as described below. To rule out the presence of proteins, LPS samples were analyzed by SDS-PAGE and proteins were quantified using the Coomassie Plus protein assay kit (Pierce).

LPS analysis

Proteinase K-digested whole-cell lysates were prepared as described previously (Hitchcock and Brown, 1983), and LPS was separated on 12% (w/v) acrylamide gels using a Tricine-sodium dodecyl sulfate (SDS) buffer system (Lesse et al., 1990). Gel loadings were normalized so that each sample represented the same number of cells as previously described (Bravo et al., 2008). Gels were silver stained by a modification of the procedure of Tsai and Frasch (1982).

TABLE IStrains and plasmids used in this study

Strain or plasmid	Relevant properties ^a	Source or reference
Shigella flexneri 2a		
2457T	wild type	ISP ^b
MSF1107	2457T wzz _{pHS2} ::aph, Kan ^R	This study
MSF1033	2457T <i>wzzB</i>	This study
MSF1749	2457T wzy	This study
MSF1210	2457T waaL	This study
MSF252	2457T mxiH::aph, Kan ^R	This study
Plasmids		
pKD46	bla P _{BAD} gam bet exo pSC101 oriTS	(Datsenko and Wanner, 2000)
pKD4	bla FRT aph FRT PS1 PS2 oriR6K	(Datsenko and Wanner, 2000)
pCP20	bla cat cI857 _PR flp pSC101 oriTS	(Cherepanov and Wackernagel, 1995)
pDA12	cloning vector, oripBBR1, mob-, Pdhfr, Tet ^R	(Aubert et al., 2008)
pDAPF2	$wzz_{ m pHS2}$ cloned into pDA12	This study
pDABF3	wzzB cloned into pDA12	This study

^a Kan^R, kanamycin resistant.

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^b ISP, Institute of Public Health, Santiago, Chile.

Type three-secretion system induction

The secretion of Ipa proteins was induced by supplementing bacterial cultures with 20 μ g/mL Congo red, as described by Bahrani et al. (1997). After the induction, supernatants were concentrated and the protein samples were analyzed by SDS-PAGE and silver stained.

Mammalian cell cultures

Caco-2 cells (American Type Culture Collection HTB-37), human colonic carcinoma cells, were maintained in high-glucose DMEM supplemented with 10% (v/v) fetal bovine serum (FBS) and non-essential amino acids (GIBCO BRL). Caco-2 polarized monolayers were obtained as follows: 3 µm pore size polycarbonate filter inserts (Millicell, Millipore) were coated with type A Gelatin (SIGMA) for 6 h before cell seeding. Then, 3×10^5 cells were seeded onto the inverted filter inserts and allowed to attach overnight. After that, inserts were placed upright in 24-well culture plates (basolateral side of monolayer upward) and maintained for 7 days in 5% CO₂ and 95% air, changing medium every 2 days. Electrical resistance was measured every day using a dual-voltage Ohmmeter clamp until a steady-state transepithelial cell resistance of approximately 400 to 800 Ω /cm was reached.

Adhesion and invasion assays

Bacteria were grown aerobically to an optical density at 600 nm of 0.3, washed twice with phosphate-buffered saline (PBS), suspended in DMEM supplemented with 10% (v/v) FBS and added to polarized Caco-2 cells monolayers at a multiplicity of infection of approximately 100. Adhesion assays were performed by co-incubating the monolayers with bacteria for 1 h at 4 °C. Then, each well was rinsed three times with 100 µL of ice-cold PBS and cells were lysed by incubation with 100 µL of sodium deoxycholate (0.5% w/v in PBS) for 10 min. LB (900 µL) was added, and each sample was vigorously mixed to recover adherent bacteria, which were quantified by plating for colony-forming units (CFU) on TSA-CR plates. Adhesion was calculated as follows: percent adhesion = $100 \times (number of$ cell-associated bacteria/initial number of bacteria added). For invasion assays, after addition of bacteria to the monolayers, the culture plates were centrifuged at 180 x g for 10 min and then incubated for 30 min at 37 °C. Cells were washed three times with pre-warmed PBS and incubated for an additional 1 h in DMEM supplemented with 10% FBS and containing gentamicin (50 µg/mL) to kill extracellular bacteria. Cells were then washed three times with PBS and lysed with sodium deoxycholate (0.5% w/v in PBS). Intracellular bacteria were diluted and plated on TSA-CR plates. Invasion was calculated as the number of bacteria resistant to gentamicin/initial number of bacteria added. Invasion and adherence of the wildtype strain was arbitrarily defined as 100%.

For competition assays, purified LPS preparations (100 μ L) at different final concentrations in PBS were added to polarized Caco-2 cells and incubated for 1 h at 37 °C. Then, PBS was removed and invasion assays were performed as described above

Data were obtained from at least three independent experiments performed in triplicate and were expressed as means ± standard errors. The statistical significance of

differences in the data was determined using the one-way analysis of variance (ANOVA) test and the Dunnett's post-test.

RESULTS

Construction and LPS phenotypes of S. flexneri strains with defined mutations in O-antigen genes

Defined, non-polar deletions of genes responsible for assembly of the O antigen were constructed. The disrupted genes included: (i) $wzz_{\rm pHS2'}$ encoding the chain length regulator that determines polymerization of the VL-OAg (>90 RU), (ii) wzzB, encoding the chain regulator responsible for polymerization of the S-OAg (11-17 RU), (iii) wzy, encoding the O-antigen polymerase, and (iv) waaL, which encodes the O-antigen ligase; a protein catalyzing the link between O-antigen chain and the lipid A-core region (Schnaitman and Klena, 1993). Fig. 1A shows the LPS patterns of the parental and LPS mutant strains. As previously reported (Carter et al., 2007), strain wzz_{pHS2}::aph (MSF1107) produced O-antigen chains lacking VL-OAg (lane 2), whereas the wzzB mutant strain (MSF1033) did not exhibit S-OAg molecules (lane 3). Mutant wzy (MSF1749) showed a semi-rough phenotype with a very prominent band corresponding to core substituted with one O-antigen unit (lane 4). Finally, the waaL mutant (MSF1210) produced LPS with a complete core region and devoid of O antigen (lane 5).

To evaluate whether the LPS defects of the mutants affected the function of the T3SS, exponential cultures of each strain were incubated with Congo red to induce secretion of the effector proteins, as described in Materials and Methods. A mutant defective in type-three secretion (*mxiH*) was also analyzed as negative control. The parental strain and all the O antigen mutants showed the characteristic electrophoretic profile of Ipa proteins (Parsot et al., 1995), indicating a functional T3SS (Fig. 2).

Polymeric O antigen with normal chain-length distribution is required for invasion, but not for adhesion, to Caco-2 cells

To address the role of the O antigen in the initial interaction of *S. flexneri* with epithelial cells, we compared the abilities of mutants and the wild-type 2457T to adhere to and enter Caco-2 cells as a polarized colonic epithelial cell model.

We first investigated the role of O antigen in adhesion. The experiments were done under conditions in which invasion was inhibited by reduced temperature (4 °C) as described in Material and Methods. As shown in Fig. 1B, all the mutants exhibited the same or higher levels of adhesion to Caco-2 cells compared to the parental strain. These results indicate that the presence of O antigen is not required for adhesion of *S. flexneri* to epithelial cells.

The ability of the O-antigen mutants to invade polarized epithelial cells was assayed. A *mxiH* mutant, which lacks one structural protein of the needle complex of the T3SS, was used as negative control. As shown in Fig. 1C, mutants in the wzz_{pHS2} or wzzB genes were significantly less invasive than the wild-type strain. This phenotype was also observed for the wzy and waaL mutants. We concluded from these data that entry of *S. flexneri* into epithelial cells from the basolateral surface requires not only polymerized O antigen but also, most importantly, a modal chain-length distribution of O-antigen molecules contributed by both WzzB and Wzz_{DHS2}.

To ensure that the defects in O-antigen chain length were responsible for the diminished invasive capacity of mutants wzz_{pHS2} and wzzB, each mutant was complemented with the corresponding intact gene cloned into the low copy number plasmid pDA12 (Table I). The complemented strains showed wild-type LPS profiles (Fig. 3A). The invasion abilities of the complemented mutants were compared with the parental strain transformed with the vector pDA12. Both strains were able to enter Caco-2 cells as well as the wild type. The complemented wzzB showed higher invasion levels, but the differences with the wild type levels of invasion were not statistically significant (Fig. 3B).

LPS does not inhibit S. flexneri invasion of polarized Caco-2 epithelial cells

To elucidate whether O-antigen molecules are required for the first interaction with epithelial cells, LPS from the parental, $wzz_{\rm pHS2}$ and wzzB strains was purified and used as a competitor in the invasion assay. As shown in Fig. 4, pretreatment of the monolayers with LPS purified (2 and 5 $\mu g/mL$) from any of the three strains did not affect the invasion ability of the parental strain. The same results were obtained when higher amounts of competitor LPS (100 and 1000 $\mu g/mL$) were added (data not shown). These results demonstrated that LPS does not compete for internalization of *S. flexneri*

into epithelial cells by the basolateral membrane, supporting our finding that LPS is not required for adhesion, but rather it would be needed in a later step of the invasion process.

DISCUSSION

The role of LPS O antigen in the initial interaction of *S. flexneri* with epithelial cells is not yet well defined. Earlier studies carried out with HeLa cells indicated that O antigen is not required for *S. flexneri* invasion (Okamura et al., 1983; Okamura and Nakaya, 1977; Rajakumar et al., 1994; Sandlin et al., 1995). These studies were conducted using mutants that were not rigorously characterized genetically, obtained either spontaneously or by transposon insertions. In any case, secondary mutations and/or polar effects on genes located downstream the mutation site in the *wba* operon cannot be ruled out.

In this work, we dissected the contribution of O antigen to epithelial cell invasion by *S. flexneri* using mutants with defined deletions of genes for O antigen assembly and polymerization. We found that polymeric O antigen is required for invasion of polarized Caco-2 cells by the basolateral side. Our results are not in accordance to those of Köhler et al.

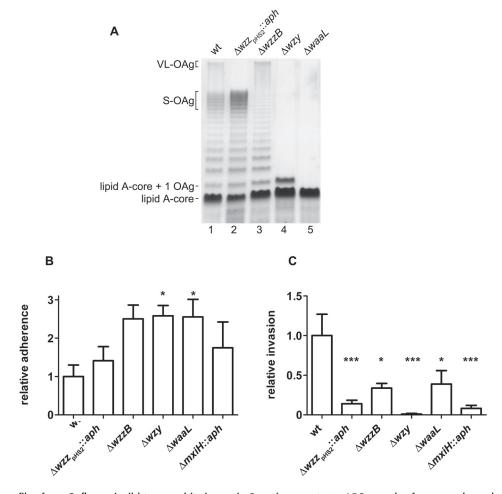


Figure 1: (A) LPS profiles from *S. flexneri* wild type and its isogenic O-antigen mutants. LPS samples from equal numbers of bacterial cells were loaded in each lane and were analyzed by Tricine/ SDS-polyacrylamide gel electrophoresis on a 12% acrylamide gel followed by silver staining. (B) Adhesion and (C) invasion of *S. flexneri* wild type and mutants to polarized Caco-2 cells. The assays were performed in triplicate on at least three independent occasions. Averages \pm standard errors (error bars) are shown. Values that are significantly different from those of the wild type are indicated by * (p<0.05) or *** (p<0.001). The strains are 2457T (wt), MSF1107 (Δwzz_{pHS2} ::aph), MSF1033 ($\Delta wzzB$), MSF1749 (Δwzy), MSF1210 ($\Delta waaL$), MSF252 ($\Delta mxiH$::aph).

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(2002), who reported that the presence of O antigen was not required for the invasion process itself, but it was essential for adhesion. In that study, invasion assays were performed by incubating the monolayer with bacteria at 37 °C for 90 min, then gentamicin was added and incubation proceeded

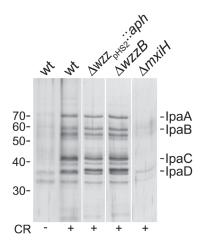


Figure 2: Secretion of Ipa proteins by *S. flexneri* wild type and mutants. Secretion was induced by incubating bacteria with Congo red (CR). Proteins in culture supernatants were electrophoresed and silver stained. The positions of Ipa proteins are indicated. Numbers indicate positions and sizes (in kDa) of standard proteins.

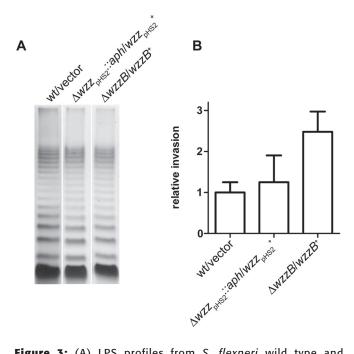


Figure 3: (A) LPS profiles from *S. flexneri* wild type and complemented O-antigen mutants (B) Invasion of polarized Caco-2 epithelial cells by *S. flexneri* wild type and complemented O-antigen mutants. The assays were performed in triplicate on at least three independent occasions. Averages ± standard errors (error bars) are shown. Values were not significantly different from those of the wild-type *S. flexneri*. The strains are 2457T/pDA12 (wt/vector), MSF1107/pDAPF2 (Δwzz_{pHS2}::aph/wzz_{pHS2}⁺) and MSF1033/pDABF3 (ΔwzzB/wzzB⁺).

for additional 90 min. After lysis of the monolayer, CFU were counted. To estimate the number of surface-associated bacteria, the invasion values were subtracted from the CFU obtained after the initial 90 min of incubation without gentamicin. So, it is possible that during incubation with gentamicin, intracellular bacteria proliferated. Thus the invasion index was over estimated, lowering the adherence values. In consequence, these experimental conditions could explain why the authors attributed to an impaired adhesion a defect in invasion.

Interestingly, we found that polymerized O antigen with a normal chain length distribution, that is, containing S-OAg and VL-OAg, is required for polarized-cell invasion by *S. flexneri* (Fig. 1). Mutants with deletions of either *wzzB* or *wzz*_{pHS2} were significantly less invasive than the parental isolate. When these mutants were transformed with plasmids containing the corresponding intact gene, the invasive capacity was restored to wild-type levels (Fig. 3).

Our results suggest that no direct interaction occurs between O-antigen molecules and a possible ligand on the epithelial cell surface to promote invasion (Fig. 4). The invasion defects of the O-antigen mutants could be attributable to a general compromise of the outer membrane integrity. As the loss of bacterial cell envelope integrity often results in increased permeability and sensitivity to detergents, cationic peptides, antibiotics and other toxic compounds (Nikaido and Vaara, 1985), bacteria were tested for sensitivity to SDS, polymyxin B, EDTA, and novobiocin by the method described by Maloney and Valvano (2006). No differences were observed between any of the mutants and the wild-type strain (data not shown), suggesting that the cell envelope of the LPS mutants remains fully functional. Nonetheless, a normal LPS may be necessary for the assembly or exposure of other surface structure that mediates the entry process. In support of this notion, it has been reported that the O antigen contributes

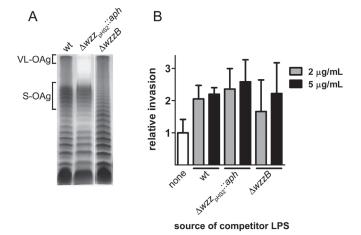


Figure 4: Effect of pre-treatment of Caco-2 cells with purified LPS on bacterial invasion. (A) LPS profiles of samples purified from *S. flexneri* wild type and its isogenic O-antigen mutants. (B) Competition assays. Caco-2 cells were pre-treated with 2 or 5 mg/mL of LPS obtained from each indicated strain before the addition of wild-type bacteria. Invasion assays were performed in triplicate on at least three independent occasions. Averages ± standard errors (error bars) are shown. Values were not significantly different from that of non treated Caco-2 cells.

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to stabilize the outer membrane by the interaction between adjacent O-antigen chains (Nikaido, 2003), and LPS also interacts with outer membrane proteins allowing their correct localization and function (Goldberg and Theriot, 1995).

In *S. flexneri*, mutants that are devoid of VL-OAg, but not of S-OAg, are highly sensitive to serum (Hong and Payne, 1997). These findings, together with the results of this study suggest that, while the presence of VL-OAg would be optimal for extracellular survival, a normal chain length distribution (S-OAg and VL-OAg) is required for internalization of bacteria into polarized epithelial cells.

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REFERENCES

- AUBERT DF, FLANNAGAN RS, VALVANO MA (2008) A novel sensor kinase-response regulator hybrid controls biofilm formation and type VI secretion system activity in *Burkholderia cenocepacia*. Infect Immun 76: 1979-1991.
- BAHRANI FK, SANSONETTI PJ, PARSOT C (1997) Secretion of Ipa proteins by *Shigella flexneri*: inducer molecules and kinetics of activation. Infect Immun 65: 4005-4010.
- BRAVO D, SILVA C, CARTER JA, HOARE A, ÁLVAREZ SA, BLONDEL CJ, ZALDÍVAR M, VALVANO MA, CONTRERAS I (2008) Growth-phase regulation of lipopolysaccharide O-antigen chain length influences serum resistance in serovars of *Salmonella*. J Med Microbiol 57: 938-946.
- CARTER JA, BLONDEL CJ, ZALDÍVAR M, ÁLVAREZ SA, MAROLDA CL, VALVANO MA, CONTRERAS I (2007) O-antigen modal chain length in *Shigella flexneri* 2a is growth-regulated through RfaH-mediated transcriptional control of the *wzy* gene. Microbiology 153: 3499-3507.
- CHEREPANOV PP, WACKERNAGEL W (1995) Gene disruption in Escherichia coli: Tc^R and Km^R cassettes with the option of Flp-catalyzed excision of the antibiotic-resistance determinant. Gene 158: 9-14.
- DARVEAU RP, HANCOCK RE (1983) Procedure for isolation of bacterial lipopolysaccharides from both smooth and rough *Pseudomonas aeruginosa* and *Salmonella typhimurium* strains. J Bacteriol 155: 831-838.
- DATSENKO KA, WANNER BL (2000) One-step inactivation of chromosomal genes in *Escherichia coli* K-12 using PCR products. Proc Natl Acad Sci USA 97: 6640-6645.
- DORMAN CJ, PORTER ME (1998) The *Shigella* virulence gene regulatory cascade: a paradigm of bacterial gene control mechanisms. Mol Microbiol 29: 677-684.
- GOLDBERG MB, THERIOT JA (1995) Shigella flexneri surface protein IcsA is sufficient to direct actin-based motility. Proc Natl Acad Sci USA 92:
- HITCHCOCK PJ, BROWN TM (1983) Morphological heterogeneity among *Salmonella* lipopolysaccharide chemotypes in silver-stained polyacrylamide gels. J Bacteriol 154: 269-277.
- HOARE A, BITTNER M, CARTER J, ÁLVAREZ S, ZALDÍVAR M, BRAVO D, VALVANO MA, CONTRERAS I (2006) The outer core lipopolysaccharide of *Salmonella enterica* serovar Typhi is required for bacterial entry into epithelial cells. Infect Immun 74: 1555-1564.
- HONG M, PAYNE SM (1997) Effect of mutations in *Shigella flexneri* chromosomal and plasmid-encoded lipopolysaccharide genes on invasion and serum resistance. Mol Microbiol 24: 779-791.

- KANIPES MI, HOLDER LC, CORCORAN AT, MORAN AP, GUERRY P (2004) A deep-rough mutant of *Campylobacter jejuni* 81-176 is noninvasive for intestinal epithelial cells. Infect Immun 72: 2452-2455.
- KLENA J, ZHANG P, SCHWARTZ O, HULL S, CHEN T (2005) The core lipopolysaccharide of *Escherichia coli* is a ligand for the dendritic-cellspecific intercellular adhesion molecule nonintegrin CD209 receptor. J Bacteriol 187: 1710-1715.
- KÖHLER H, RODRIGUES SP, MCCORMICK BA (2002) Shigella flexneri interactions with the basolateral membrane domain of polarized model intestinal epithelium: Role of lipopolysaccharide in cell invasion and in activation of the mitogen-activated protein kinase ERK. Infect Immun 70: 1150-1158.
- LESSE AJ, CAMPAGNARI AA, BITTNER WE, APICELLA MA (1990) Increased resolution of lipopolysaccharides and lipooligosaccharides utilizing tricine-sodium dodecyl sulfate-polyacrylamide gel electrophoresis. J Immunol Methods 126: 109-117.
- MALONEY K E, VALVANO M A (2006) The *mgtC* gene of *Burkholderia* cenocepacia is required for growth under magnesium limitation conditions and intracellular survival in macrophages. Infect Immun 74: 5477-5486.
- NIKAIDO H, VAARA M (1985). Molecular basis of bacterial outer membrane permeability. Microbiol Rev 49: 1-32.
- NIKAIDO H (2003) Molecular basis of bacterial outer membrane permeability revisited. Microbiol Mol Biol Rev 67: 593-656.
- OKAMURA N, NAKAYA R (1977) Rough mutant of *Shigella flexneri* 2a that penetrates tissue culture cells but does not evoke keratoconjunctivitis in guinea pigs. Infect Immun 17: 4-8.
- OKAMURA N, NAGAI T, NAKAYA R, KONDO S, MURAKAMI M, HISATSUNE K (1983) HeLa cell invasiveness and O antigen of *Shigella flexneri* as separate and prerequisite attributes of virulence to evoke keratoconjunctivitis in guinea pigs. Infect Immun 39: 505-513.
- PARSOT C, MÉNARD R, GOUNON P, SANSONNETTI PJ (1995) Enhanced secretion through the *Shigella flexneri* Mxi-Spa translocon leads to assembly of extracellular proteins into macromolecular structures. Mol Microbiol 16: 291-300.
- PARSOT C (2009) *Shigella* type III secretion effectors: how, where, when, for what purposes? Curr Opin Microbiol 12: 110-116.
- RAETZ CR, WHITFIELD C (2002) Lipopolysaccharide endotoxins. Annu Rev Biochem 71: 635-700.
- RAJAKUMAR K, JOST BH, SASAKAWA C, OKADA N, YOSHIKAWA M, ADLER B (1994) Nucleotide sequence of the rhamnose biosynthetic operon of *Shigella flexneri* 2a and role of lipopolysaccharide in virulence. J Bacteriol 176: 2362-2373.
- SANDLIN RC, LAMPEL KA, KEASLER SP, GOLDBERG MB, STOLZER AL, MAURELLI AT (1995) Avirulence of rough mutants of *Shigella flexneri*: requirement of O antigen for correct unipolar localization of IcsA in the bacterial outer membrane. Infect Immun 63: 229-237.
- SCHNAITMAN CA, KLENA JD (1993) Genetics of lipopolysaccharide biosynthesis in enteric bacteria. Microbiol Rev 57: 655-682.
- SCHROEDER GN, HILBI H (2008) Molecular pathogenesis of *Shigella* spp.: controlling host cell signaling, invasion, and death by type III secretion. Clin Microbiol Rev 21: 134-156.
- SCHUCH R, MAURELLI AT (1997) Virulence plasmid instability in *Shigella flexneri* 2a is induced by virulence gene expression. Infect Immun 65: 3686-3692.
- TSAI CM, FRASCH CE (1982) A sensitive silver stain for detecting lipopolysaccharides in polyacrylamide gels. Anal Biochem 119: 115-119.
- WEST NP, SANSONETTI P, MOUNIER J, EXLEY RM, PARSOT C, GUADAGNINI S, PRÉVOST MC, PROCHNICKA-CHALUFOUR A, DELEPIERRE M, TANGUY M, TANG CM (2005) Optimization of virulence functions through glucosylation of *Shigella* LPS. Science 307: 1313-1317.
- ZAIDI TS, FLEISZIG SM, PRESTON MJ, GOLDBERG JB, PIER GB (1996) Lipopolysaccharide outer core is a ligand for corneal cell binding and ingestion of *Pseudomonas aeruginosa*. Invest Ophthalmol Vis Sci 37: 976-986.

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