Comparing Invasive and Non-Invasive of Isolated *Shigella flexneri* by Electron Microscopy of Cell Culture, SDS-PAGE and Congo Red Method

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Received 12 November 2005; revised 24 June 2006; accepted 9 August 2006

ABSTRACT

**Background:** The aim of this study was to compare invasive and non-invasive strains of *Shigella flexneri* isolated from Tehran by a 120 kDa protein band by SDS-PAGE, electron microscopy of cell culture and Congo red dye methods. **Methods:** *S. flexneri* strains were isolated by standard bacterial methods from fecal specimens of children attending to the 3 children’s hospitals. Phenotype analysis for screening virulent of strains of *S. flexneri* was done on a plate of tryptic soy agar contained 0.003% Congo red dye. Whole membrane protein preparations were used to examine the protein profiles of the inner and outer membrane of these Gram-negative bacteria. The protein mixture was electrophoresed through a polyacrylamide gel. The gel was stained with Coomassie brilliant blue R250 and destained with ethanol and acetic acid. HeLa cell culture was done by two-step preparations: one for light microscopy and the other for electron microscopy. **Results:** Some of *S. flexneri* (46%) were Congo red positive colonies. *S. flexneri* with negative Congo red phenotype could not enter the HeLa cell culture. A 120 kDa protein band was found in 46% of these bacteria which could enter into HeLa cell culture. Pseudopod structures which facilitate bacterial cell-to-cell spread were readily identified by electron microscopy. **Discussion:** Since the existence of 120-kDa protein band was corresponded to enter of *S. flexneri* into the HeLa cell culture and correlated with Congo red dye positive, for identification of invasive and non-invasive *S. flexneri* strains, the use of a 120-kDa protein band by SDS-PAGE or a simple, rapid and very cheap Congo red dye method is recommended. Because, there are some deaths due to *Shigella sp.* in our country, notification on the isolation of these bacteria in both children hospitals laboratories and private clinical laboratories is important. *Iran. Biomed. J*. 11 (1): 47-52, 2007

Keywords: *Shigella flexneri*, HeLa cell culture, Electron microscopy, Congo red dye

INTRODUCTION

Among pathogenic microorganisms invasive bacteria have the ability to penetrate mammalian epithelial cells both *in vivo* and *in vitro* [1]. *Shigella flexneri*, which is responsible for a dysenteric syndrome in human, belongs to the invasive group of pathogens [2-4]. It is known that the ability of *S. flexneri* to penetrate epithelial cells is encoded by a 20-kilo base portion of a 220-kilo base plasmid [1, 4, 5]. This is unexpected for a non-motile microorganism which has been related to icsA inter cellular spread, vir G, a gene encoding a 120-KDa outer membrane protein which allows interaction with microfilaments [1, 6]. The entry of *S. flexneri* into epithelial cells is achieved through internalization of the bacterium into a membrane bound vacuole derived from the host cell plasma membrane [1]. *Shigella* species remain within human intestinal epithelial cells where they cause the destruction of enterocytes and induce an inflammatory response [2]. *S. flexneri* requires both adhesive and invasive phenotype to efficiently colonize follicle-associated epithelium (FAE) [7]. Recent studies on the enter invasive pathogen, *S. flexneri*, have shown that, in addition to allowing...
intracellular growth [8], lysis of the phagocyte vacuole also allows bacteria to spread intracellular and infect adjacent cells [8, 3]. Cell cultures are commonly used to assess the ability of intracellular bacteria to invade susceptible eukaryotic cells [9].

In the years of 1980, there were a lot of studies on identification and the intracellular existence of S. flexneri. But in recent years, because of health improvement in conditions in developed countries, search for isolation of this bacteria decreased but in developing countries, such as Iran and Taiwan, many children still lose their life for infection of S. flexneri [10, 11]. So, the aim of this study was to compare the invasive and non-invasive properties of isolated S. flexneri by electron microscopy (EM) of cell culture, SDS-PAGE and Congo red.

MATERIALS AND METHODS

**Bacterial strains.** S. flexneri strains were isolated from fecal specimens of children attending to the 3 children’s hospitals (Markaz-e-Te’bii Kodakan, Aliasghar and Mofid Hospitals) from January 2001 to December 2003. In this study, 350 Shigella sp. were isolated and identified by standard methods. After that, 100 S. flexneri strains were randomly chosen and stored in peptone and glycerol at -70°C.

**Bacterial suspensions.** Bacteria were harvested in tryptic soy broth in the exponential phase, washed in PBS [NaCl (8.8 gL⁻¹); Na₂HPO₄ 2H₂O (2.250 gL⁻¹); NaH₂PO₄, H₂O (0.257 gL⁻¹); pH 7.4] and suspended at the appropriate density of 2 × 10⁶ cfu ml⁻¹ in MEM.

**Congo red binding.** Phenotype analysis for screening virulent of strains of S. flexneri was done by Congo red dye. A colony of fresh culture of isolated bacteria was inoculated on a plate contained TSA (tryptic soy agar) and Congo red solution at final concentration of 0.003% to detect red pigmentation colony [12, 13].

**Protein preparation and SDS-PAGE.** Whole membrane preparations were used to examine the protein profiles of the inner and outer membrane of Gram-negative bacteria. In this process, pellet of the cells were suspended in 30 mM Tris-HCl (pH 8.1) and resuspended in 20% sucrose/30 mM Tris-HCl (pH 8.1) plus lysozyme. Then, 3 mM EDTA (pH 7.3) was added. Terminal pellets were suspended in 1 x LUG buffer [Tris-HCl (pH 6.8), 50 ml; SDS 25 ml, 0.25 M; glycerol, 2g; beta-mercaptoethanol, 5 ml; bromophenol blue, 2 ml; of 1%, distilled water, 100 ml] [14]. Following certain preparative steps, the protein mixture was electrophoresed through a polyacrylamide 8% gel. The gel was stained with Coomassie brilliant blue R250 and continuously destained with mixture of ethanol and acetic acid. The 120-kDa band was compared with protein ladder (protein ladder, Page Ruler™ # Smo661, Fermentas, Lituani).

**HeLa cell cultures and EM.** To proceed for experiment, two-step preparations were done: one for light microscopy and the other, for EM. HeLa cell was obtained from the Public Health Medical School of Tehran University (Tehran, Iran). New cultures were prepared in tissue culture (trays) wells consisting of cover slip (22 × 22 mm) and concentration of 2 × 10⁵ cell/ml using 0.5 ml MEM (Minimal Essential Medium) with 5% FCS. Then, bacterial suspensions with MOI (multiplicity of infection) 10 bacteria/cells were added to each well. Process of bacterial suspension preparation was as explained below:

After gentle mixing, the trays were inoculated in the CO₂ incubator at 37°C for 1 h and then washed three times and fresh tissue culture medium (MEM) with 5% FCS containing gentimicin at a final concentration of 40 mg/ml was added to each well and were incubated in the CO₂ incubator at 37°C for another 2 h. The cover slips were fixed with methanol at 4°C overnight and then stained with Giemsa stain and washed with Giemsa buffer for light microscopy examination.

For EM investigation, HeLa cell monolayer was seeded in 100 ml culture flasks and after washing for three times, the bacterial suspensions were added. All flasks scraped off and centrifuged. Fixation with glutaraldehyde 2% was done on pellets of cells for 2 hours. Continuously, a centrifuge step was done and melted agar 2% was added to each cell pellet, mixed well and post fixed for 1 h with Osmium tetroxide, and then, 4 concentrations of aceton were added for dehydration and samples were embedded in resin spur (R1032, Agar Scientific, UK). Gold sections were taken and stained with saturated uranyl acetate and lead citrate [15]. A grid of cell culture without infection with S. flexneri was used as control.
RESULTS

Congo red binding. All 100 *S. flexneri* isolated were tested for binding to Congo red dye. Then, isolated bacteria were identified as positive invasive phenotype with red colonies and non-invasive phenotype identified with white colonies. In this study, 46 (46%) isolated *S. flexneri* were Congo red positive colonies on TSA contained 0.003% Congo red dye. All the Congo red positives (100%) strains produced β haemolysin on blood agar.

SDS-PAGE. From 100 isolated *S. flexneri*, a 120-kDa band was detected in 46 isolates (46%). The range of protein bands was from 30 kDa to 150 kDa. The protein profiles of some strains are shown in Table 1 and Figure 1. On the bases of protein bands, these were 12 distinct groups. Protein bands with the same size were present in several strains, for example, most of strains (20.7%) were in group I.

HeLa cell culture and EM. As it was described above, all isolates were tested for binding to Congo red dye; so, invasive and non-invasive strains were separated. Bacterial suspensions from *S. flexneri* 2a strain (Reference Strain, RS) with Congo red positive and probably IcsA positive, *S. flexneri* strains with Congo red positive and probably IcsA positive and *S. flexneri* strains with Congo red negative and probably IcsA negative isolated from patients were used to infect HeLa cell line. HeLa cell cultures which were infected with either RS or *S. flexneri* strains with Congo red positive and probably IcsA positive at MOI 10:1, showed marked loss of confluence and viable cells were heavily infected with bacteria and showed morphology with indistinct membrane (46%).

According to pictures, invasive *shigella* strains penetrated into these cells (Fig. 2). HeLa cells infected with *S. flexneri* strains with Congo red positive and probably IcsA positive, exhibited structures similar to those of non-infected cells. However, *S. flexneri* strains with Congo red positive and probably IcsA positive adhered and occasionally entered to HeLa cells. Furthermore, pseudopod structures used to facilitate bacterial cell-to-cell spread were readily identified.

Table 1. Protein profiles of *S. flexneri* isolated.

<table>
<thead>
<tr>
<th>Group</th>
<th>30</th>
<th>40</th>
<th>50</th>
<th>60</th>
<th>70</th>
<th>85</th>
<th>100</th>
<th>120</th>
<th>150</th>
<th>Bacteria (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>I.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>20.6</td>
</tr>
<tr>
<td>II.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>12.1</td>
</tr>
<tr>
<td>III.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>7.3</td>
</tr>
<tr>
<td>IV.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>6.0</td>
</tr>
<tr>
<td>V.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>2.3</td>
</tr>
<tr>
<td>VI.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>17.0</td>
</tr>
<tr>
<td>VII.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>1.2</td>
</tr>
<tr>
<td>VIII.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>2.4</td>
</tr>
<tr>
<td>IX.</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
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<td>+</td>
<td>-</td>
<td>12.3</td>
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<td>X.</td>
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<td>-</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>1.2</td>
</tr>
<tr>
<td>XI.</td>
<td>+</td>
<td>-</td>
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<td>-</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>3.0</td>
</tr>
<tr>
<td>XII.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>14.6</td>
</tr>
</tbody>
</table>

+, exist; -, not exist.
**DISCUSSION**

In this study, 350 *Shigella spp.* from fecal of patients were isolated and 142 (40.57%) were *S. flexneri*. One hundred of *S. flexneri* strains which recovered from stool specimens of children’s were chosen randomly for presumptive determination of IcsA protein (120 kDa band) and Congo red binding, then, infect HeLa cell line as noted above. Forty six (46%) of *S. flexneri* strains were Congo red positive colonies on TSA plate contained 0.003% Congo red dye. Relation between the virulence of *S. flexneri* 2a and its ability to absorb Congo red was examined in Mounier study [16]. This property is correlated to invasiveness phenotype of *S. flexneri* in cell culture [17]. Present results showed, all Congo red positive isolates had adhesion and invasion properties to HeLa cells.

These data were agreed with those of other studies [1, 9, 18-22]. Francis and Thomas showed that, infected Caco-2 or HeLa cell cultures by *L. monocytogenes* at high MOI, had extensive invasion, intra-cellular multiplication and finally cell lysis [10, 21].

As it was noted above, in 46 (46%) of isolated *S. flexneri* strains, a 120 kDa band was detected in SDS- PAGE method, that might be correlated to IcsA protein. Presence of a 120 kDa band and phenotype Congo red positive were detected in 46 (46%) of isolates in the same time. In this study, both *S. flexneri* 2a RS and *S. flexneri* strains with Congo red positive and probably IcsA positive were isolated from patients, showed similar internalization, cell existence and cell disruption. In addition, pseudopod filaments were confirmed by EM.

In contrast, most of *S. flexneri* strains probably IcsA negative could not enter HeLa cells culture. Moreover, a few of these bacteria could adhere and enter to the HeLa cells, but cell disruption was not detected. These bacteria produce necrosis later, but can not be detected after 3 h incubation. Unlike the Hly (Haemolysin) positive strains of *L. monocytogenes*, none of Hly negative bacteria had spread inter-cellular after 2 h of incubation and only a few number of them had spread after 4 h of incubation [18].

In other study, interaction of *Salmonella typhimurium*, *Listeria monocytogenes* with murin’s M cells were compared. Tissue infected with the lower dose of organisms did not show significant M

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**Fig. 2.** Penetration of invasive *S. flexneri* (vacuole formation) which had a 120-kDa band (presumptive IcsA protein) and Congo red positive.

**Electron microscopy.** *S. flexneri* 2a (RS) with Congo red positive and probably IcsA positive and patient’s isolated *S. flexneri* strains with Congo red positive and patient’s isolated *S. flexneri* strains with Congo red negative and probably IcsA negative which infected HeLa cell cultures were used for EM investigation. Internalization, existence of bacteria into cells and cell disruption of both Congo red positive and probably IcsA positive *S. flexneri* strains and *S. flexneri* 2a (RS) were detected. Existence of pseudopod filaments were confirmed by EM (Fig. 3).

**Fig. 3.** Pseudopod filaments formation in infected HeLa cell with *S. flexneri* strain which had a 120-kDa band and Congo red positive.
cell disruption at the various times examined and had the same appearance as the FAE. In contrast, at the higher dose, the interactions between S. flexneri strains and the epithelium of peyer’s patches were similar to those observed for L. monocytogenes. Destroyed regions revealed, membrane blebs and a denuded epithelial surface that closely resembled to those observed for Listeria. These data demonstrated that L. monocytogenes and S. flexneri possess the ability to induce massive destruction of FAE when inoculated into intestinal loops at inoculums of 4 × 10⁹ cfu per ml [5].

In this study, the 120 kDa protein band which detected by SDS-PAGE has been correlated with IcsA protein (1), the detection of this protein band may be a useful tool in epidemiological studies for searching prevalence sources.

Results showed, the existence of the 120 kDa protein band was corresponded to enter of S. flexneri into the HeLa cell culture, demonstrated by EM and also, this band protein was found in all S. flexneri strains with Congo red dye positive. In the other hand, our results detected by different methods (SDS-PAGE, EM and Congo red) could confirm each other. Moreover, the ability of S. flexneri to enter the host cells, spread to adjacent cells and intra cellular growth could damage and lyse the host cells, which indicated invasiveness property (8). So, for identification of invasive and non-invasive S. flexneri strains, detection of the 120 kDa protein band by SDS-PAGE or a simple, rapid and very cheap Congo red dye method is recommended.

Because, there are some deaths particularly in children due to Shigella spp in our country [12] notification on the isolation of this bacteria in both children hospitals laboratories and private clinical laboratories is important.

ACKNOWLEDGEMENTS

We gratefully acknowledge from Dr. H. Monavari in Hazrate Rasol Hospital Laboratory (Tehran, Iran) who helped us for cell culture. Moreover, we thank Miss Nafisi, Mrs. Abedini from Microbiology Laboratory of Mofid and Tebbi Kodakan Hospitals respectively.

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