

Bacterial pathogens associated with bloody diarrhea in Uruguayan children

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ABSTRACT

Diarrheal disease continues to be a serious health problem, especially in developing countries. Bloody diarrhea represents approximately 20-30% of all cases and has higher morbidity and mortality. Treatment with antibiotics is beneficial in cases of *Shigella*, *Campylobacter*, *Yersinia* and *Salmonella* infection, principally in those children with a higher risk of invasive disease. The aims of this study were to detect the bacterial agents associated with bloody diarrhea in children and to determine their antimicrobial susceptibility patterns. Between June 2001 and January 2008, 249 children with bloody diarrhea were studied. *Shigella* and Shiga toxin-producing *Escherichia coli* (STEC) were recovered from 48 (19.3%) and 3 (1.2%) of the total of cases, respectively. In 49 out of 249 children, in whom other enteropathogens were investigated, we recovered *Campylobacter jejuni* from 7 children (14.3%), *Salmonella* spp. from 2 (4.1%) and *Aeromonas* spp. from 1 (2%) in addition to *Shigella* from 7 children (14.3%). Thirty-four (70%) *Shigella* isolates showed resistance to ampicillin and 13 (27%) to trimethoprim-sulfamethoxazole. All *Shigella* isolates were susceptible to nalidixic acid, ciprofloxacin and ceftriaxone. *Salmonella* and STEC isolates were susceptible to all antibiotics assayed. Thus, the use of trimethoprim-sulfamethoxazole or ampicillin would not be appropriate for the empirical treatment of *Shigella* – associated diarrhea.

Key words: bloody diarrhea, *Shigella*, antibiotics

RESUMEN

Patógenos bacterianos asociados a diarrea con sangre en niños uruguayos. La enfermedad diarreica es un problema de salud importante, especialmente en los países en desarrollo. La diarrea sanguinolenta representa el 20-30% del total de casos y determina una mayor morbimortalidad. La antibioticoterapia es beneficiosa en casos de infección por *Shigella*, *Campylobacter*, *Yersinia* y *Salmonella*, principalmente en niños con riesgo elevado de enfermedad invasora. Los objetivos del presente trabajo fueron conocer las bacterias asociadas a diarrea sanguinolenta y determinar su patrón de sensibilidad a los antimicrobianos. Entre junio de 2001 y enero de 2008 se estudiaron 249 niños con diarrea sanguinolenta. *Shigella* y *Escherichia coli* productor de toxina Shiga (STEC) se recuperaron de 48 (19,3%) y 3 (1,2%) casos, respectivamente. En la subpoblación de 49 niños cuyo estudio coprobacteriológico fue más completo se aisló, además de *Shigella* (14,3%), *Campylobacter jejuni* (14,3%), *Salmonella* spp. (4,1%) y *Aeromonas* spp. (2%). Treinta y cuatro (71%) aislamientos de *Shigella* mostraron resistencia a ampicilina y 13 (27%) a trimetoprima-sulfametoxazol. Todos los aislamientos de *Shigella* fueron sensibles a ácido nalidixico, ciprofloxacina y ceftriaxona. Los aislamientos de *Salmonella* y STEC fueron sensibles a todos los antibióticos ensayados. En función de estos resultados, se concluye que el uso de trimetoprima-sulfametoxazol y de ampicilina no sería apropiado para el tratamiento empírico de la diarrea asociada a *Shigella*.

Palabras clave: diarrea con sangre, *Shigella*, antibióticos

Diarrheal disease continues to be a health problem worldwide, especially in developing countries. In these regions, it accounts for approximately 2.5 million deaths per year in children under 5 years of age. Furthermore, acute diarrhea considerably contributes to morbidity and increases health care costs in children from industrialized countries (6). Bloody diarrhea (BD) represents approximately 20-30% of all cases, causing important inflammatory intestinal illness and, un-

der some circumstances, producing severe complications, like sepsis, hemorrhagic colitis and hemolytic uremic syndrome (HUS) (1, 6).

The bacterial pathogens associated with BD include species of *Shigella*, *Campylobacter*, *Salmonella*, *Escherichia coli* pathotypes, especially Shiga toxin-producing *Escherichia coli* (STEC) and enteroinvasive *E. coli* (EIEC), as well as *Yersinia enterocolitica*. The prevalence

of these agents and their antimicrobial susceptibility patterns vary among different regions (4, 12, 13).

Treatment with antibiotics is indicated in cases of *Shigella* infections and in special situations of *Campylobacter*, *Yersinia* and *Salmonella* infections, principally in those children at higher risk of invasive disease (5, 15).

However, the lack of rapid diagnostic methods for the detection of all these agents supports the implementation of empirical treatment regimens based on the epidemiological knowledge of the prevalent agents and the corresponding antimicrobial susceptibility.

The aims of this study were to detect the bacterial pathogens associated with bloody diarrhea in children and to determine their antimicrobial susceptibility patterns.

Within the framework of an institutional program aimed at the regional surveillance of STEC infections, a prospective, multicenter study was done involving health care centers in Montevideo and other cities of Uruguay. The analysis period extended from June 2001 to January 2008, 249 children with bloody diarrhea who required medical attention in these centers were studied. Stool samples were consecutively obtained and only one sample per child was processed. Each stool specimen was sent to the Department of Bacteriology and Virology in a Cary-Blair transport medium, whereas another part of the sample without transport medium was sent in a sterile plastic vial. Both were submitted in insulated, refrigerated boxes and were examined less than 12 hours after extraction. All samples were examined for the presence of fecal leukocytes, STEC and *Shigella* spp. Fecal leukocytes were semiquantified by microscopic examination of smears stained with methylene blue as follows: < 1 cell (+), 1-10 cells (++) and > 10 cells (+++) by high-power field. STEC detection was performed by culture and PCR using specific primers to *stx*₁ and *stx*₂ genes as previously described (14). *Shigella* species were investigated by classic procedures (3). Furthermore, between January 2005 and June 2006, at the request of the pediatrician in charge, we also studied forty-nine (20%) of these 249 children for the presence of *Salmonella*, *Yersinia*, EIEC and *Campylobacter* species, as described by Torres *et al.* (12). STEC, *Salmonella* and *Shigella* isolates were serotyped by standard procedures using commercial antisera (Difco®) and others available in our institute's collection (3). Antimicrobial disk susceptibility tests for *Shigella*, *Salmonella* and STEC isolates were done according to the Clinical and Laboratory Standards Institute guidelines (2). The following agents were tested: ampicillin (AMP), trimethoprim-sulfamethoxazole (TMP-SMX), chloramphenicol (CMP), tetracycline (TET), ciprofloxacin (CIP), nalidixic acid (NAL), gentamicin (GEM), ceftriaxone (CRO), ceftazidime (CAZ), and cefoxitin (FOX). The Fisher's test was used to assess the association between qualitative variables.

The median age was 12 months (range, 5 days to 14 years of age), and 90% of the children were under five years old. More than 90% of the children were assisted

at public health centers. No epidemiological link could be established among the 249 children studied. In the whole period of study, *Shigella* spp. was recovered from 48 (19.3%) children and STEC from 3 (1.2%) of the 249 cases. One child showed co-infection with two different STEC serogroups (O26 and O145). In those children (49) whose stool culture also included the search for *Salmonella*, *Yersinia*, EIEC and *Campylobacter* species, *Shigella* spp. was recovered from 7 (14.3%) children, *Campylobacter jejuni* from 7 (14.3%), *Salmonella* from 2 (4.1%), and *Aeromonas* spp. from one (2%) child. Fecal leukocytes were present in all *Shigella*-positive samples but only in 2 of the 10 children from whom another enteropathogen was isolated ($p = 0.0018$). Fecal leukocytes were present in 43 (89.6%) of the 48 fecal samples from which *Shigella* spp. was recovered; 32 of them showed ≥ 10 cells per field (+++). Ninety-four per cent of the cases of bloody diarrhea associated with *Shigella* were observed in the warmest months (between November and April). Thirty-seven isolates corresponded to *Shigella flexneri* and 11 to *Shigella sonnei*. Neither *Salmonella dysenteriae* nor *Shigella boydii* were recovered during this study. Twenty-one of the 37 *S. flexneri* strains corresponded to serotype 2a, 13 to serotype 3c and 3 to serotype 1.

Thirty out of 37 (81%) *S. flexneri* isolates showed resistance to AMP (17 isolates belonged to serotype 2a; 11 to serotype 3c and 2 to serotype 1) and 13 (35%) to TMP-SMX (7 isolates belonged to serotype 2a; five corresponded to serotype 3c and one to serotype 1). However, when we analyzed the periods 2001-2003 and 2004-2008, the results were 95% and 62% resistance to AMP; 48% and 18% resistance to TMP-SMX, respectively ($p > 0.05$). *S. flexneri* isolates showed eight antimicrobial resistance phenotypes and the TET, AMP, CMP resistance pattern was the most frequent (13 strains), followed by the TET, AMP, CMP, TMP-SMX resistance pattern (Figure 1). Two *S. flexneri* strains were susceptible to all antimicrobials tested. On the other hand, *S. sonnei* showed a unique AMP-resistance profile AMP^R. Four out of 11 *S. sonnei* isolates had this profile and the other 7 isolates were susceptible to all antimicrobial agents tested. All *Shigella* isolates were susceptible to NAL, CIP, GEM, CRO, CAZ, FOX. STEC isolates belonged to serogroups O26 (2 strains), O111 (1 strain) and O145 (1 strain). All STEC strains carrying *eae* and *ehxA* genes. Isolates O26 carried *stx* genes, while strain O145 was positive for *stx*₂ and strain O111 carried both *stx*₁ and *stx*₂ genes. The *Salmonella* strains corresponded to *Salmonella enterica* subsp. *enterica* serovar Enteritidis. The *Salmonella* and STEC cultures were susceptible to all antimicrobials tested. Neither EIEC nor *Y. enterocolitica* were recovered during this study.

As reported in other regions, *Shigella* spp. was the most frequently recovered pathogen from BD cases (11, 13). Nevertheless, when more complete stool cultures were done, *C. jejuni* was an important agent of BD. Therefore, in the future it would be important to perform antimi-

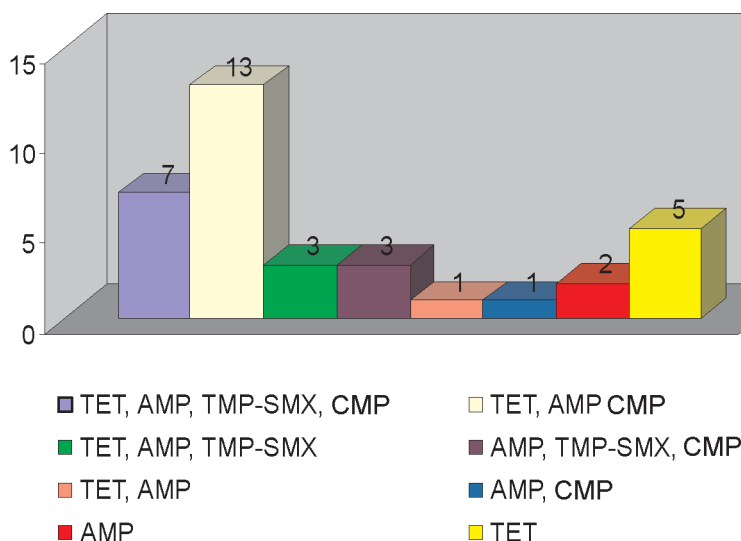


Figure 1. Antimicrobial resistance patterns in *Shigella flexneri* isolates.

crobal susceptibility tests to know their resistance pattern. The absence of *S. dysenteriae* was not surprising. In Uruguay, *S. dysenteriae* strains are rarely found and they belong to serotype 2 (12). According to our results, the BD cases that occurred during the warmest months and showed fecal polymorphonuclear leukocytes were likely caused by *Shigella* spp. ($p = 0.017$). Serotype 2a was the most frequent among *S. flexneri* strains. These findings are comparable with results from Chile (9). It would be important to define antigenic determinants that should be included in a future vaccine.

As it occurs in other regions (13), we have not recovered *Y. enterocolitica* or EIEC in a subgroup of these 249 children. However, these results are limited due to the small sample size and probably on account of the methodology used for the detection of these agents which included only phenotypic tests. Although STEC infection represented a small proportion of all acute bloody diarrhea cases studied (1.2%), the serious associated complications, such as HUS, seem to justify the importance of searching for this agent. In this sense, 2 out of 3 children infected with STEC developed HUS during the follow-up. *Salmonella* isolates were susceptible to all antibiotics tested. Results obtained with *Salmonella typhimurium* by Macedo *et al.* (7) showed a marked decrease in resistance in the last years. In our country, since December 2003, the recommended drugs for empirical treatment of BD cases in children have been: oral azithromycin (5 mg/kg/day, for 5 days) for outpatients (over 6 months of age and without toxicity signs) and parenteral ceftriaxone (100 mg/kg, *i/m*, one dose) for inpatients (under 6 months of age or with toxicity signs). When we analyzed *Shigella*

spp. strains isolated between 2001 and 2003 *versus* those isolated in 2004 and 2008, we observed that AMP resistance decreased from 95% to 62%, and TMP-SMX resistance from 48% to 18%. This trend may be explained by appropriate application of this therapeutic guideline; however, studies with larger number of samples are required to determine whether this difference is significant and sustained. In our country, like in other regions, the use of TMS or AMP continues to be inappropriate for the empirical treatment of BD cases. All *Shigella* strains recovered in this study were susceptible to NAL. This antibiotic has been the drug of choice for the last two decades in certain regions, such as Southern Africa and South Asia (15); nevertheless, physicians should anticipate increasing resistance to this drug as its use increases. In addition, *Shigella* strains resistant to NAL showed some degree of cross-resistance to CIP. Thus, the widespread use of NAL for treatment of shigellosis may reduce the efficacy of CIP (10). Fluoroquinolones are also recommended drugs for treating *Campylobacter*, *Salmonella* and *Shigella*-associated gastroenteritis (5, 15). According to the results of this study, ciprofloxacin could be used for the treatment of children with BD. However, we believe that this drug should be reserved for the treatment of severe cases for which there is no other treatment option available, taking into account the potential risk of joint damage.

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