Hospital-based Surveillance for Rotavirus Gastroenteritis Among Young Children in Bangladesh

Defining the Potential Impact of a Rotavirus Vaccine Program

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Background: In anticipation of introduction of a rotavirus vaccine into the national immunization program of Bangladesh, active hospital-based surveillance was initiated to provide prevaccine baseline data on rotavirus disease.

Methods: Children 5 years of age and younger admitted with acute gastroenteritis (AGE) (≥2 watery or looser-than-normal stools or ≥1 episode of forceful vomiting) at 7 hospitals throughout Bangladesh were identified. Clinical information and stool specimens were collected from every 4th patient. Specimens were tested for rotavirus antigen by enzyme immunoassays; ≥25% of detected rotaviruses were genotyped.

Results: From July 2012 to June 2015, rotavirus was detected in 2432 (64%) of 3783 children hospitalized for AGE. Eight enrolled children died, including 4 (50%) who were rotavirus positive. Rotavirus was detected year-round in Bangladesh with peak detection rates of >80% during November–February. Most (86%) rotavirus AGE cases were 6–23 months of age. Sixty-nine percent of children with rotavirus had severe disease (Vesikari score, ≥11). Among 543 strains genotyped, G1P[8] (31%) and G12P[8] (29%) were the most common.

Conclusions: Rotavirus is a major cause of morbidity in Bangladeshi children, accounting for nearly two-thirds of AGE hospitalizations. These data highlight the potential value of rotavirus vaccination in Bangladesh, and will be the key for future measurement of vaccine impact.

Key Words: Bangladesh, acute gastroenteritis, rotavirus, hospital-based surveillance, rotavirus vaccination

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Rotavirus, the most common cause of severe diarrhea in infants and young children, remains a significant cause of morbidity and mortality worldwide.1 In 2010, ~200,000 global diarrhea deaths among children 5 years of age and younger were attributed to rotavirus, with ~25% occurring in Southeast Asia.2 In Bangladesh, single-site studies have demonstrated a considerable burden of rotavirus on healthcare systems; rotavirus was the etiology of gastroenteritis in 20% of children seeking care in a diarrhea treatment facility in urban Dhaka,3 and in 33% of the hospitalizations because of gastroenteritis in rural Matlab.4

Two live attenuated oral rotavirus vaccines, Rotarix (RV1; Merck, Whitehouse Station, NJ), a pentavalent (G1, G2, G3, G4, P[8]) bovine-human reassortant vaccine, and Rotarix (RV1; GSK Biologicals, Rixensart, Belgium), a monovalent (G1P[8]) human strain vaccine, are currently recommended by the World Health Organization for introduction into national immunization programs worldwide.4 A key factor in the decision by countries to introduce rotavirus vaccines has been demonstrating high rotavirus disease burden through establishment of surveillance for rotavirus diarrhea. In addition, prevaccine data on disease burden have been crucial for monitoring vaccine impact after vaccine introduction. Assessing vaccine impact in resource-limited settings is particularly important given the lower efficacy of RV1 and RV5 shown in clinical trials in Africa and Asia, compared with that seen in Europe and America.5,10

In anticipation of introduction of a rotavirus vaccine into the routine immunization schedule in Bangladesh, this study aims to provide baseline data on the burden of rotavirus gastroenteritis and circulating strains at sentinel hospitals nationwide. Because the most recent verbal autopsy data from the Bangladesh Demographic and Health Survey noted an 85% reduction in diarrhea-specific mortality between 2004 and 2011 [from 7 to 1 per 1000 live births],11 we specifically aimed at measuring the contribution of rotavirus to healthcare utilization.

MATERIALS AND METHODS

Starting in July 2012, an active hospital-based rotavirus surveillance system, consisting of 7 tertiary hospitals located in all 7 divisions of Bangladesh (Fig. 1), was initiated in 3 phases. These hospitals were chosen because they have a high number of pediatric gastroenteritis admissions each year. Rotavirus surveillance was started at the Dhaka, Rajshahi and Sylhet sites in July 2012, extended to Chittagong and Rangpur in February 2013, and further extended to Khulna and Barisal in August 2013. Surveillance followed the World Health Organization protocol for rotavirus surveillance in hospital settings.12 At each hospital, from 8:30 am to 4:00 pm each day (except for weekends and holidays), field assistants identified children 5 years of age and younger admitted to pediatric wards with diarrhea by reviewing admission logbooks and screened them for acute gastroenteritis (AGE) symptoms. AGE was defined as the occurrence of ≥3 watery or looser-than-normal stools or ≥1 episode
of forceful vomiting within a 24-hour period, with symptoms lasting ≤7 days. Surveillance physicians enrolled every 4th child listed who met the surveillance case definition. Compared with other sites, diarrhea-associated hospitalizations in Rangpur were low in number so, only at this site, the protocol was modified after 3 months to enroll every child admitted with AGE. Surveillance physicians collected demographic and clinical information of enrolled children from the parents and hospital records using a standard questionnaire and assessed the extent of dehydration following clinical criteria in World Health Organization diarrhea treatment guidelines.13 In addition, discharge and death logbooks were reviewed to ascertain the outcome of children admitted with diarrhoea but not enrolled in the surveillance.

Field assistants collected bulk stool specimens (4 mL) from each child on the day of enrollment and immediately stored specimens in a −70°C liquid nitrogen dry shipper after collection. The study team shipped stool samples in these containers to the International Centre for Diarrheal Disease Research (icddr,b) virology laboratory in Dhaka every 2 weeks, where a commercially available enzyme-linked immunoassay (EIA) (Prospect; Oxoid Diagnostics Ltd, Basingstoke Hants, United Kingdom) was used to test for rotavirus antigen. At icddr,b, stool samples were stored at −70°C. Every 3 months, G and P genotyping of ~25% of rotavirus-positive specimens were done using methods described previously.14 EIA, genotyping and sequencing have been done at icddr,b virology laboratory in Dhaka.

We obtained written informed consent from the enrolled children’s parents or guardians. The study protocol was reviewed and ethics approval obtained from the ethical review committee of icddr,b.

We calculated the overall and site-specific proportions of rotavirus-associated AGE hospitalizations by dividing the number of rotavirus-positive stools by the total number of samples collected and tested for rotavirus. To estimate the number of rotavirus-associated hospitalizations, the overall proportion of rotavirus-associated AGE hospitalizations was applied to the total number of AGE admissions during the surveillance period. We compared clinical and demographic characteristics and outcomes between children testing positive versus negative for rotavirus. This included an assessment for differences in clinical severity of AGE between the 2 groups, using the 20-point Vesikari scale: illnesses with scores <7 were classified as mild, illnesses with scores ≥7 and ≤10 as moderate and illnesses with scores >11 as severe.15 Proportions were compared using χ² and Cochran–Armitage trend tests, and continuous variables were compared using the Wilcoxon rank sum test.

RESULTS

During 212 hospital months of surveillance, 129,156 children 5 years of age and younger were admitted to the pediatric wards of participating hospitals, of whom 14,814 (12%) were hospitalized with AGE. Surveillance staff collected clinical data and stool specimens from 26% (3783/14,814) of children admitted with AGE (Table 1). EIA testing identified rotavirus antigen in the stools of 64% (2432/3783) of enrolled children; detection rates varied by site from 59% to 69% (Table 1). Extrapolation of rotavirus-testing results to the untested AGE cases yielded an estimated total of 9728 rotavirus AGE admissions. Thus, about 8% (9728/129,156) of all pediatric admissions at participating hospitals were attributable to rotavirus-associated AGE during the study period.

Nearly all (96%) of the rotavirus hospitalizations were among children <2 years (Fig. 2), with 57% occurring during the first year of life. Rotavirus hospitalizations occurred year-round, with rotavirus accounting for >10% of AGE admissions during every month of the year. Overall, rotavirus was detected in >80% of AGE cases during the rotavirus peak season (November–February) (Fig. 3). There were no differences in the seasonal patterns among surveillance sites (data not shown).

The median Vesikari score among children with rotavirus-confirmed AGE was slightly higher for those with compared to those without laboratory-confirmed rotavirus infection (12 vs. 11;
P < 0.001). The duration of illness before hospitalization was longer for children with rotavirus infection compared with children who did not have evidence of rotavirus infection (3 vs. 2 days; \( P < 0.001 \)), but that there were no differences in the length of hospital stay (2 days) (Table 2). Among the children enrolled over the study period, 8 (0.25%) died in the hospital. Rotavirus was detected in the stool of 50% (4/8) of those who died from AGE, and 3 of the 4 rotavirus-associated deaths occurred in children 6–23 months of age (the remaining death was in a 3-month old infant). According to discharge and death logbooks, 104 children were admitted with diarrhea and died in hospital during the study period; 69% were not be screened or enrolled because they were admitted after surveillance hours, or because death occurred soon after arrival at the hospital (see Table, Supplemental Digital Content 1, http://links.lww.com/INF/C564)

Genotyping was performed on 22% (543/2432) of positive rotavirus stool samples (Table 3). Four G genotypes (G1, G2, G9 and G12) and 3 P genotypes (P4, P6 and P8) were identified. G12 (40%) and G1 (32%) were the most prevalent G types, and P[8] (76%) and P[6] (12%) were the most prevalent P types. G1P[8] (31%) and G12P[8] (29%) were the most commonly identified strains. Mixed rotavirus strains were seen in 13% of the children; 5 (1%) of specimens were untypable. All detected G and P genotypes were observed in sites across the 7 divisions of Bangladesh.

**DISCUSSION**

This study confirms the considerable burden of rotavirus AGE among children 5 years of age and younger in Bangladesh.
We found that rotavirus accounted for ~64% of all childhood hospitalizations for AGE or ~8% of all pediatric hospitalizations at participating hospitals during the study period. Rotavirus was detected year-round among children with AGE, and during the peak winter months from November to February, the detection rates of rotavirus among children hospitalized with AGE exceeded 80%. Children with rotavirus AGE had illness that was somewhat more severe than those with AGE from other causes. Children 6–23 months of age accounted for ~85% of all rotavirus AGE hospitalizations, indicating that rotavirus vaccines, which are administered in the first few months of life, have the potential to reduce the burden of rotavirus-related deaths, as well as case-fatality ratios. Second, our surveillance systematically excluded children who had a documented temperature, resulting in an increase in the proportion of diarrhea attributable to rotavirus. This is supported by data from a longitudinal evaluation of hospitalizations data from icddr,b showing that the proportion of diarrhea cases attributable to rotavirus nearly doubled from 2002–2004 compared with 1993–1995 (42% vs. 22%).

This study demonstrated substantial rotavirus strain diversity within Bangladesh. As opposed to previous years (1991–2012), when G1, G2 and G9 were the predominant strains detected in Bangladesh,22,23 G12 was also commonly detected genotype in the study period.24 We found that 31% of detected strains were homotypic to strains in both vaccines (ie, matched in both the G and P types), 44% were partially heterotypic (matched only the G or P type) and 12% were fully heterotypic (ie, did not match the G or P type). In addition, ~14% of the strains were mixed or untypable using current primers. Although both natural and vaccine-induced protection against a range of genotypes has been shown in developed settings,25,26 the considerable variability in circulating rotavirus strains in Bangladesh warrants an assessment of the effect prevalent strains in Bangladesh.
and settings. Third, because controls were not included in this evaluation, we may have overestimated the proportion of cases attributable to rotavirus. However, any overestimation is expected to be small, as EIA rarely detects asymptomatic infections, and the attributable fraction of rotavirus detection with EIA has been shown to be >90%.27–29

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