



CODEN [USA]: IAJPB

ISSN : 2349-7750

**INDO AMERICAN JOURNAL OF
PHARMACEUTICAL SCIENCES**

SJIF Impact Factor: 7.187

<http://doi.org/10.5281/zenodo.4313378>Available online at: <http://www.iajps.com>

Research Article

**DETERMINE PARTICULAR EAECs VIRULENCE GENES
RELATED TO THE DURATION AND TYPE OF DIARRHEA IN
PAKISTANI CHILDREN**¹Dr Usama Shahbaz, ²Dr Muhammad Yasin Saeed, ³Dr Uzair Nadeem¹Faisalabad Medical University, Faisalabad.**Article Received:** October 2020**Accepted:** November 2020**Published:** December 2020**Abstract:**

Aggregative Escherichia coli is often found in diarrheal stools around the world. It has been associated with fluid bowel movements, weight reduction and failure to thrive in children living in non-industrialized countries. Various important destructive qualities of EAEC are recognized; yet their parts in intense and persevering runs have not been examined recently. The purpose of this review was to recognize the explicit destructive qualities of EAEC related to the scope and type of racing of Pakistani youth. It was our intention to improve current EAEC diagnostics and to allow the focus to be placed on strains with a normal course of extreme disease. Surveys completed by the tutors provided data on the duration of races and the presence of blood or body fluids. In addition, the transmission of EAEC qualities in strains from unhealthy, mucoid and watery bowel cases was studied. Our current research was conducted at Sir Ganga Ram Hospital, Lahore from March 2019 to February 2020. Disposition and Relapse Tree Examination (DRE) was applied to explore the relationship between the harmful qualities of EAEC and the length and type of diarrhea. Consistent loose bowel was related to strains without the peak quality ($p = 0.003$) and to the mixture of peak and sat qualities, furthermore, the non-appearance of the aggA quality ($p = 0.06$). Delayed runs were related to the mixing of the aatA and astA qualities ($p=0.05$). Non-mucosal runs were related to strains without the aatA grade ($p = 0.006$). Intense bowel relaxation was related to the aggR, aap and aggA grades by singular odds ratios. Gentamicin and ciprofloxacin obstruction was observed in 9.7 and 4% of the strains individually. Multi-drug obstruction was observed in 38% of the strains. Hereditary host factors were linked to an increased risk of EAEC-related disease. Hence, we studied a table of hazard factors in two groups of children - EAEC positive and EAEC negative - in order to distinguish additional factors favoring disease. The duration of breastfeeding was strongly related to the likelihood of having a place in the EAEC negative group.

Keywords: EAECs virulence genes, type of diarrhea in Pakistani children.**Corresponding author:****Dr. Usama Shahbaz,**

Faisalabad Medical University, Faisalabad.

QR code



Please cite this article in press Usama Shahbaz et al, **Determine Particular Eaecs Virulence Genes Related To The Duration And Type Of Diarrhea In Pakistani Children.**, Indo Am. J. P. Sci, 2020; 07(12).

INTRODUCTION:

Children who tested positive for EAEC were found to have increased levels of fecal lactoferrin and II-1b, despite the presence of gastrointestinal side effects in a Brazilian survey [1]. This demonstrates a capacity for significant worsening of EAEC and extreme illness, which may also be present in children in industrialized countries [2]. EAEC has been associated with the free bowels of young people in Germany, Great Britain and America. In addition, a few episodes of EAEC have been reported in youth in Serbia, Japan, and Korea, and children contaminated with EAEC have been found to have fluid bowel movements, which can cause significant electrolyte loss and impaired micronutrient retention [3]. Some host inheritance factors have been associated with a lack of increased defense against EAEC disease, including single nucleotide polymorphisms in the interleukin-8 region of the advertisers and in the quality of CD. In any event, virtually no general hazard factors related to EAEC disease have been studied. Defense testing of EAEC strains has revealed impressive opposition to anti-infective agents, including opposition to ciprofloxacin, opposition to several drugs and a wide range of beta-lactamases, which is of concern [4]. The pathogenic capacity of EAEC in children in industrialized countries warrants further exploration and the role of the destructive factors of EAEC in the intense and determined loosening of the intestines needs to be explained. The highest quality level for recognizable evidence of EAEC is the measurement of the HEp-2 cell. This test is performed in reference laboratories only; it requires cell culture offices and is tedious. It is based on the recognition of the supposed appearance of "stacked blocks" by unprepared professors and has been found to fluctuate from

observer to observer. In addition, this phenotypic test does not recognize pathogenic and non-pathogenic strains [5].

METHODOLOGY:

Between January 2011 and October 2013, we conducted an amulet study to distinguish children with loose bowels and identified fecal tests. The analytical units participating in the survey were the Department of Bacteria, Parasites, Fungi, the Statin Sera Institute and the Departments of Clinical Microbiology at Slag Else Hospital and Hvidovre Hospital at Copenhagen College in Pakistan. At Copenhagen College Hvidovre Hospital, only patients suffering from traveler's intestines were investigated for CEEA. Our current research was conducted at Sir Ganga Ram Hospital, Lahore from March 2019 to February 2020. Fecal tests were performed on runaway youths who had been advised by their general expert or who had been hospitalized. Youth with positive fecal tests for EAEC and no co-morbidities were qualified to participate in the study. Between January 2011 and October 2013, 299 youth tested positive for EAEC in the three branches of clinical microbiology of interest (Figure 1). Of the 290 children, 88 (33%) were avoided due to co-morbidities with any one or a few additional enteric microorganisms. Among the co-infections, the most consistently distinguished were appendicular and destructive *E. coli* (AEEC; n = 28), rotavirus (n = 15), ETEC (n = 14) and *C. difficile* (n = 10). Of the 209 youth who tested positive for ECAA, guardians of 50 children did not have a registered telephone number, could not be reached, did not speak Pakistani or did not live in Pakistan, and were shunned. We included 156 EAEC-positive youth in the EAEC HIV-positive meeting alone.

Figure 1:

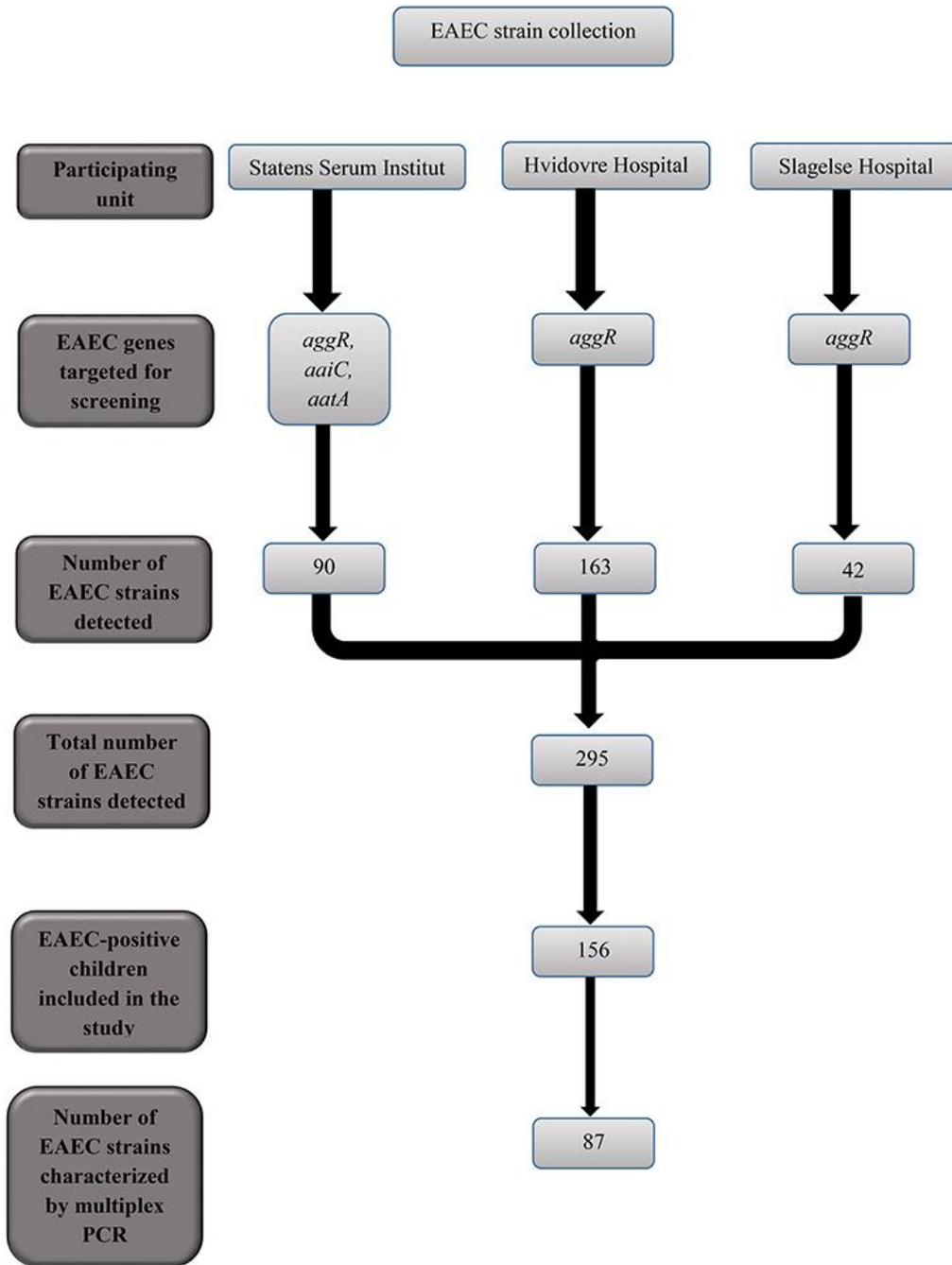


Table 1:

	Number of genes detected (%)	Description	Function
<i>aap</i>	67 (77)	Dispersin	Anti-aggregation
<i>aggR</i>	65 (75)	Transcription activator	Major regulator
<i>aggA</i>	14 (16)	Fimbrial subunit for AAF/I	Adhesion
<i>aafA</i>	6 (7)	Fimbrial subunit for AAF/II	Adhesion
<i>agg3A</i>	6 (7)	Fimbrial subunit for AAF/III	Adhesion
<i>agg4A</i>	10 (12)	Fimbrial subunit for AAF/IV	Adhesion
<i>agg5A</i>	17 (20)	Fimbrial subunit for AAF/V	Adhesion
<i>agg3/4C</i>	30 (35)	Usher	Adhesion
<i>aatA</i>	81 (93)	ABC transporter	Transporter of dispersin
<i>astA</i>	27 (31)	Aggregative heat-stable toxin	Toxin
<i>aaIC</i>	58 (67)	Type VI secretion system	Secreted protein
ORF3	65 (75)	Cryptic protein	Unknown
<i>pet</i>	9 (10)	Plasmid encoded toxin	Toxin
<i>pic</i>	45 (52)	Serine protease precursor	Toxin
<i>sat</i>	25 (29)	Secreted auto-transporter protein	Toxin
<i>sepA</i>	21 (24)	Serine protease auto-transporter toxin	Toxin
<i>sigA</i>	3 (3)	Protease-like homolog	Toxin

RESULTS:

In order to study the predominance of EAEC in the intestinal slackening of youngsters in Pakistan, we inspected the annual ubiquity of all common pathogens in excreta tests submitted to routine microbiological investigation at the Department of Bacteria, Parasites and Fungi at the Serum Institute in Statins . Between June 2011 and June 2012, 1,360 young people aged ≤ 11 were inspected (Table 4). The rates of youngsters testing positive for bacterial microbes were as follows AEEC (14%; n= 149), EAEC (8.8%; n= 94), EPEC (5.8%; n= 37), *Campylobacter* spp. (2.7%; n = 32), *Salmonella* spp. (1.3%; n = 18), ETEC (0.7%; n = 9), VTEC (0.9%; n = 9), *Yersinia* spp. (0.4%; n = 2), ETEC (0.1%; n = 1) and *Shigella* spp. (0,1 % ; n = 1). During this one-year period, EAEC was the second most recognized enteric microbe among young Danes with diarrhea. Of the 159 EAEC-positive children, 25 had an obscure length of free bowel, and in the remains of the children's collection, 89 strains of EAEC were available for multiplex PCR investigation. The circulation of destructive qualities of EAEC was sat (29%), sepA (26%), peak (54%), sigA (6%), pet (13%), astA (34%), aggR (77%), aatA (96%), aaiC (68%), aap (78%),

ORF3 (78%), agg3/4C (37%), agg3A (8%), aafA (9%), aggA (18%), agg4A (15%), and agg5A (20%). The circulation of EAEC grades in youth with intense and assiduous bowel relaxation is introduced in Table 5. To investigate the relationship between EAEC grades and intense and assiduous bowel relaxation, we examined the CART tree. This examination groups the qualities progressively according to the classes examined. For each branch, a measurable critique between the non-appearance and presence of an additional quality divides the tree into new branches and gives a separation between intense and diligent bowel relaxation, separately. Five exceptions with reports of runs of 200 days or more were eliminated from the survey, leaving 85 perceptions. Of the 83 perceptions, 29 EAEC strains were collected from children with intense and loose bowel movements, 49 strains from youth with tireless and loose bowel movements. Eleven youth had soft stools for 9 to 17 days. To begin, we present the results from the truck tree, where we treat the duration of the runs as a complete development by using the classes intensely and consistently. The EAEC strains without the photo quality necessarily caused diligent bowel relaxation (≥ 17 days), $p = 0.003$ (Figure 2).

Table 2:

	EPEC-positive group n = (156)	EPEC-negative group n = (155)	P-value with confidence interval
AGE IN YEARS			
0-2	102 (65%)	78 (50%)	$P = 0.5404^a$
3-5	38 (24%)	73 (47%)	[-0.6; 0.3]
6+	15 (10%)	4 (3%)	
Median age	2	2	
GENDER			
Boys	92 (59%)	87 (56%)	$P = 0.4326^b$ [-0.1; 0.2]
Mean birth weight	3516 g	3414 g	$P = 0.1692$ [-43.4; 246.3] ^a
HISTORY OF INFANTILE COLIC			
Yes	26 (17%)	18 (12%)	$P = 0.2008^b$
No	130 (83%)	137 (88%)	[-0.1; 0.1]
BREASTFEED CURRENTLY			
Yes	9 (6%)	3 (2%)	$P = 0.0792^b$
No	138 (94%)	146 (94%)	[-0.0; 0.1]
PREVIOUSLY BREASTFED			
Yes	131 (84%)	142 (92%)	
No	16 (10%)	7 (4%)	
Not answered	9 (6%)	6 (4%)	
BREASTFEEDING DURATION			
>0-5 months	35 (22%)	23 (15%)	$P = 0.0017^a$
6-9 months	49 (31%)	43 (28%)	[-2.7; 0.6]
10-12+ months	42 (27%)	69 (45%)	
Not answered	30 (19%)	20 (13%)	
DOMESTIC ANIMAL			
Yes	37 (24%)	33 (21%)	$P = 0.9368^b$
No	119 (76%)	118 (75%)	[-0.1; 0.1]
Unanswered	0	7 (4%)	
DIARRHEA^c			
Yes	156 (100%)	34 (22%)	
No	0 (0%)	115 (74%)	
Not answered	0 (0%)	6 (4%)	
HOSPITALIZED			
Yes	19 (12%)	0	
CONTACT WITH SICK ANIMALS^c			
Yes	7 (5%)	9 (6%)	$P = 0.5351^b$
No	118 (76%)	115 (74%)	[-0.2; 0.1]
Unsure	31 (20%)	28 (18%)	
Unanswered	0 (0%)	3 (2%)	
FOREIGN TRAVEL^c			
Yes	79 (51%)	26 (17%)	$P = 0.0000^b$
No	76 (49%)	128 (83%)	[0.2; 0.4]
Not answered	0 (0%)	1 (1%)	
USE OF ANTIBIOTICS^c			
Yes	26 (17%)	28 (18%)	$P = 0.4798^b$

(Continued)

Table 3:

	EAEC-positive group n = (156)	EAEC-negative group n = (155)	P-value with confidence interval
AGE IN YEARS			
0-2	102 (65%)	78 (50%)	$P = 0.5404^a$
3-5	38 (24%)	73 (47%)	[-0.6; 0.3]
6+	15 (10%)	4 (3%)	
Median age	2	2	
GENDER			
Boys	92 (59%)	87 (56%)	$P = 0.4326^b$
			[-0.1; 0.2]
Mean birth weight	3516 g	3414 g	$P = 0.1692$
			[-43.4; 246.3] ^a
HISTORY OF INFANTILE COLIC			
Yes	26 (17%)	18 (12%)	$P = 0.2008^b$
No	130 (83%)	137 (88%)	[-0.1; 0.1]
BREASTFEED CURRENTLY			
Yes	9 (6%)	3 (2%)	$P = 0.0792^b$
No	138 (94%)	146 (94%)	[-0.0; 0.1]
PREVIOUSLY BREASTFED			
Yes	131 (84%)	142 (92%)	
No	16 (10%)	7 (4%)	
Not answered	9 (6%)	6 (4%)	
BREASTFEEDING DURATION			
>0-5 months	35 (22%)	23 (15%)	$P = 0.0017^a$
6-9 months	49 (31%)	43 (28%)	[-2.7; 0.6]
10-12+ months	42 (27%)	69 (45%)	
Not answered	30 (19%)	20 (13%)	
DOMESTIC ANIMAL			
Yes	37 (24%)	33 (21%)	$P = 0.9368^b$
No	119 (76%)	118 (75%)	[-0.1; 0.1]
Unanswered	0	7 (4%)	
DIARRHEA^c			
Yes	156 (100%)	34 (22%)	
No	0 (0%)	115 (74%)	
Not answered	0 (0%)	6 (4%)	
HOSPITALIZED			
Yes	19 (12%)	0	
CONTACT WITH SICK ANIMALS^c			
Yes	7 (5%)	9 (6%)	$P = 0.5351^b$
No	118 (76%)	115 (74%)	[-0.2; 0.1]
Unsure	31 (20%)	28 (18%)	
Unanswered	0 (0%)	3 (2%)	
FOREIGN TRAVEL^c			
Yes	79 (51%)	26 (17%)	$P = 0.0000^b$
No	76 (49%)	128 (83%)	[0.2; 0.4]
Not answered	0 (0%)	1 (1%)	
USE OF ANTIBIOTICS^c			
Yes	26 (17%)	28 (18%)	$P = 0.4798^b$
Penicillin	14 (9%)	24 (16%)	[-0.1; 0.1]
Macrolides	2 (1%)	0 (0%)	
Other	1 (1%)	0 (0%)	
Unknown	4 (3%)	3 (2%)	
No	128 (82%)	126 (81%)	
Not answered	1 (1%)	1 (1%)	

^aIndependent sample t-test comparing the duration of breastfeeding between the study groups.

^bDifference in proportions test.

^cWithin a period of 2 months prior to sampling.

N/A, not applicable.

DISCUSSION:

EAEC is a recognized regular diarrheal microbe, but the distinctive evidence of a single pathogenic and causative EAEC quality remains uncertain. In addition, a considerable number of destructive factors, mixed together, have been associated with clinical disorders in epidemiological investigations [6]. It is conceivable that host resistance and host exposure play an important role in the pathogenicity of EAEC, therefore the mixture of harmfulness qualities in EAEC gives the important assortment to neighborhood diseases [7]. The mosaic nature of the EAEC genomes described today seems to improve our view that the pathogenicity of EAEC is subject to the host and its current status [8]. In this review, we represented different strains of EAEC collected from juveniles with intense and diligent intestinal slackening [9]. The EAEC strains were described taking into account the traditional harmfulness qualities of EAEC and antimicrobial opposition profiles. We found that lack of photo quality was related to relentless intestinal slackening, $p = 0.02$. Similarly, the mixture of peak and sat qualities and the absence of aggA quality was related to relentless defecation, $p = 0.06$ (Figure 2). The peak quality has a mucolytic movement, causing agglutination of the hematoma and opposition of the serum [10].

CONCLUSION:

The continuous passages were related to EAEC strains without the peak quality, and to strains with the mixture of peak and sat qualities and the non-participation of the aggA quality. The mixing of aatA and astA qualities was related to delayed bowel relaxation. Intensely relaxed intestines were related to the aggR, aap and aagA qualities by singular odds ratios. Strains that were found to be insufficient for aatA quality were related to non-mucosal diarrhea. Breastfeeding was considered a defense against EAEC disease after the weaning period.

RESFERENCES:

1. Sukkua K, Patungkaro W, Sukhumungoon P. Detection and molecular characterization of enteroaggregative Escherichia coli from diarrheal patients in tertiary hospitals, southern Thailand. Southeast Asian J Trop Med Public Health. 2015;46:901-910. [PubMed] [Google Scholar]
2. Haghi F, Zeighami H, Hajiahmadi F, Khoshvagh H, Bayat M. Frequency and antimicrobial resistance of diarrhoeagenic Escherichia coli from young children in Iran. J Med

- Microbiol. 2014;63:427–432. [[PubMed](#)] [[Google Scholar](#)]
3. Kubomura A, Misaki T, Homma S, Matsuo C, Okabe N. Phenotypic and molecular characterization of enteroaggregative *Escherichia coli* isolated in Kawasaki, Japan. *Jpn J Infect Dis.* 2017;70:507–512. [[PubMed](#)] [[Google Scholar](#)]
 4. Harrington SM, Dudley EG, Nataro JP. Pathogenesis of enteroaggregative *Escherichia coli* infection. *FEMS Microbiol Lett.* 2006;254:12–18. [[PubMed](#)] [[Google Scholar](#)]
 5. Nataro JP, Steiner T, Guerrant RL. Enteroaggregative *Escherichia coli*. *Emerg Infect Dis.* 1998;4:251–261. [[PMC free article](#)] [[PubMed](#)] [[Google Scholar](#)]
 6. Franca FL, Wells TJ, Browning DF, Nogueira RT, Sarges FS, Pereira AC, et al. Genotypic and phenotypic characterisation of enteroaggregative *Escherichia coli* from children in Rio de Janeiro, Brazil. *PLoS One.* 2013;8:e69971. [[PMC free article](#)] [[PubMed](#)] [[Google Scholar](#)]
 7. Brzuszkiewicz E, Thurmer A, Schuldes J, Leimbach A, Liesegang H, Meyer FD, et al. Genome sequence analyses of two isolates from the recent *Escherichia coli* outbreak in Germany reveal the emergence of a new pathotype: Entero-Aggregative-Haemorrhagic *Escherichia coli* (EAHEC) *Arch Microbiol.* 2011;193:883–891. [[PMC free article](#)] [[PubMed](#)] [[Google Scholar](#)]
 8. Cerna JF, Nataro JP, Estrada-Garcia T. Multiplex PCR for detection of three plasmid-borne genes of enteroaggregative *Escherichia coli* strains. *J Clin Microbiol.* 2003;41:2138–2140. [[PMC free article](#)] [[PubMed](#)] [[Google Scholar](#)]
 9. Hebbelstrup Jensen B, Poulsen A, Hebbelstrup Rye Rasmussen S, Struve C, Engberg JH, Friis-Møller A, et al. Genetic virulence profile of enteroaggregative *Escherichia coli* strains isolated from Pakistani children with either acute or persistent diarrhea. *Front Cell Infect Microbiol.* 2017;7:230. [[PMC free article](#)] [[PubMed](#)] [[Google Scholar](#)]
 10. Estrada-Garcia T, Perez-Martinez I, Bernal-Reynaga R, Zaidi MB. Enteroaggregative *Escherichia coli*: a pathogen bridging the north and south. *Curr Trop Med Rep.* 2014;1:88–96. [[PMC free article](#)] [[PubMed](#)] [[Google Scholar](#)]