

# *Clostridium difficile* infection and colonisation in children under 3 years of age: prospective comparative study

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## ABSTRACT

**Aims:** Despite an increasing trend in *Clostridium difficile* infections (CDI) and high *C. difficile* colonization rate especially among younger children, infants remain quite resistant to the disease. The goals of this study were to distinguish whether there exists a difference in CDI between children with or without diarrhoea, ascertain the prevalence of CDI, and assess CDI severity in children under 3 years with diarrhoea in our institution.

**Methods:** A prospective study was conducted from May 2015 to June 2016. Children 3 years of age or younger were enrolled and into two groups. Every faecal sample was tested using a diagnostic two-step screening algorithm including an immunochromatographic test and polymerase chain reaction.

**Results:** The study enrolled 147 children with diarrhoea and 75 control patients. The prevalence of CDI in children with diarrhoea was 2% (3/147), the prevalence of toxigenic *C. difficile* in the diarrhoeal group compared to the control group was 11.6% (17/147) vs. 10.6% (8/75) ( $p < 0.9999$ ).

**Conclusions:** No significant difference was observed between infants with diarrhoea and the control group. We recommend not examining for *C. difficile* children not exhibiting specific risk factors.

## KEYWORDS

infants – neonates – *Clostridium difficile* – diarrhoea – paediatrics – colonisation

## SOUHRN

Musil V., Homola L., Vrba M., Braunová A., Kravalová T., Malá M., Krbková L.: Infekce a kolonizace *Clostridium difficile* u dětí do 3 let věku: prospektivní srovnávací studie

**Cíl:** Navzdory narůstajícímu trendu klostridiové kolitidy a vysoké míře kolonizace *Clostridium difficile* mezi mladšími dětskými pacienty, malé děti zůstávají k nemoci poměrně rezistentní. Cílem studie bylo rozlišit, zda existuje rozdíl v klostridiové kolitidě mezi dětmi s průjmem a bez průjmu, určit počet záchytů klostridiové kolitidy ve sledované skupině, zhodnotit závažnost klostridiové kolitidy u dětí do 3 let věku s průjmem.

**Metody:** Prospektivní hodnocení bylo vedeno od května 2015 do června 2016. Děti mladší 3 let byly zapsány do dvou skupin. Každý

vzorek stolice byl testován dvoustupňovým algoritmem zahrnující imunochromatografický test a polymerázovou řetězovou reakci.

**Výsledky:** Zapsáno 147 pacientů s průjmem a 75 kontrol. Počet záchytů klostridiové kolitidy u dětí s průjmem byl 2% (3/147), počet záchytů toxigenních kmenů ve skupině s průjmem ve srovnání s kontrolní skupinou byl 11,6% (3/147) vs. 10,6% (8/75) ( $p < 0.9999$ ).

**Závěr:** Nebyl pozorován žádný významný rozdíl mezi dětmi s průjmem a bez průjmu. Nedoporučujeme cíleně vyšetřovat děti na *Clostridium difficile* při absenci rizikových faktorů.

## KLÍČOVÁ SLOVA

děti – novorozenci – *Clostridium difficile* – průjem – pediatrie – kolonizace

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## INTRODUCTION

*Clostridium difficile* is a non-invasive bacteria, the pathogenesis of which is based upon the action of enterotoxin A and cytotoxin B binding to receptors of intestinal epithelial cells. The toxins mediate intestinal disease with a broad spectrum of clinical manifestations ranging from mild diarrhoea to life-threatening disease. The Development CDI requires disruptions in the human intestinal microbiota [1–3].

Epidemiology and severity of *C. difficile* infections have been widely studied in adults. Some reports assume that CDI in the child population is on the rise, but recent study dispute that there is an upward trend [4–7]. A considerable role in *C. difficile*'s changing epidemiology

has been attributed to the discovery of an epidemic hypervirulent strain, North American Pulsed Field Type 1, ribotype 027 (NAP1), which has been responsible for outbreaks [8–10]. Other ribotypes have been revealed in subsequent years. In relation to the hypervirulent strains, a third toxin, the so-called binary toxin, has been uncovered [1, 11].

Despite the high colonization rate, infants remain quite resistant to the disease. The mechanism of that resistance, however, is obscure. Some hypotheses suppose that the cellular components essential for attachment of the toxin are absent or that infants lack appropriate toxin receptors on the bowel mucous membrane, as some studies have demonstrated on animal models [12–14].

## PŮVODNÍ PRÁCE

The aims of the present study were to determine whether or not there exist differences between children with and without diarrhoea, ascertain the prevalence of CDI, and assess the severity of CDI within our institution in children under 3 years of age with diarrhoea.

### MATERIALS AND METHODS

#### Patients and samples

We performed a prospective case-control study on hospitalized children ( $\leq 3$  years of age) with or without diarrhoea. The study was conducted from 11 May 2015 to 11 June 2016 in the Department of Paediatric Infectious Diseases, University Hospital Brno. This 60-bed hospital facility provides paediatric outpatient and inpatient care, including intensive care, for children younger than 19 years of age with a spectrum of paediatric diagnoses. Children  $\leq 3$  years of age were enrolled and separated into two groups and several subgroups. The first group was composed of patients investigated and admitted due to diarrhoeal disease regardless of the final diagnosis (diarrhoea as a sign of intestinal or extra-intestinal cause). Inpatients without diarrhoea were enrolled into the control group. Inpatients with varying final diagnoses (respiratory infection, infectious mononucleosis, tonsillitis, scarlet fever, cervical lymphadenitis, staphylococcal scalded skin syndrome, herpetic stomatitis, chickenpox, urinary infections, aseptic meningitis, influenza, toxoallergic rash) were recruited at our institution. The subgroups were divided according to age: neonates (defined as infants aged 28 days or younger), infants  $\leq 1$  year old, 1–2 years old, and 2–3 years old. Only one faecal specimen was obtained from each patient. In the first group, the faecal samples were submitted at the beginning of hospitalization (1st or 2nd day). In the second group, the samples were collected whenever possible. Two infants required therapy in the intensive care unit.

#### Inclusion and exclusion criteria

The criteria for inclusion into the diarrhoea patients group was defined as a presence of liquid stools with defecation frequency of 3 or more stools within 24 hours or more frequently than is normal for the individual on the basis of a personal history. The stool frequency was ascertained from the patient's parent. In neonates, diarrhoea was evaluated as a change in stool calibre. The control group was determined by absence of diarrhoea before admission to the hospital. CDI was defined as presence of diarrhoea and a positive stool test result for toxigenic *C. difficile* (IT toxin/PCR gene or toxin +) and with exclusion of non-*C. difficile* enteropathogenic organisms. Severe CDI was defined as a patient with CDI who fulfilled the following criteria within 30 days of symptom onset: history of admission to the intensive care unit (ICU), history of surgery for toxic megacolon, motility disorder, perforation or refractory colitis, or death caused by CDI (because of their usefulness, the criteria from Sathyendran et al. were used) [15]. None of the enrolled children required hospitalization for CDI before stool sample collection. Community-acquired CDI was determined

by the absence of hospitalization in the previous 3 months and a hospital stay not exceeding 48 h at the time of sampling. Hospital-acquired CDI was determined by a history of hospitalization (less than 3 months) and/or duration of hospitalization lasting more than 48 h at the time of sampling. Children with previous CDI history were excluded.

#### Data collection

The electronic clinical record was reviewed on the day of stool sampling. Patient data included age, gender, duration of hospitalization, risk factors (hematologic and solid malignancies, hematologic diseases, immunosuppression, solid organ transplantation, inflammatory bowel disease or other gastrointestinal disorders, repeated hospitalizations, previous antibiotic and cytotoxic drug exposure, and gastric acid suppression within 10 weeks of the day of the stool specimen), and details of treatment. Laboratory data including blood test results and additional microbiological testing were recorded. The information was obtained from the medical history record in the hospital's database.

#### Detection and identification of *C. difficile*

Faecal specimens were stored at 4 °C and transported in sterile containers. Stool analysis for *C. difficile* was conducted within 24 h after stool sampling. Diagnosis of CDI was based on a diagnostic two-step screening algorithm. Each faecal sample was tested by IT for the presence of the GDH and toxins A and B (using TECHLAB C. DIFF QUICK CHEK COMPLETE). Faecal samples that were GDH negative and toxin A/B negative were evaluated as negative. Stools which were GDH positive and toxin A/B positive were interpreted as positive. Faecal samples that were GDH positive but toxin A/B negative were confirmed by PCR (GeneXpert Clostridium difficile Cepheid) for presence of a gene for toxin B, the binary toxin, and/or *tcdC* deletion in nucleotide 117. Stools which were GDH positive and PCR toxin B gene positive were interpreted as "suspected CDI" (Toxigenic strains of *C. difficile*) [16]. Patients with this result are included in a cluster of CDI. None of the stool samples for *C. difficile* were cultured. Each faecal sample in the first group was also tested for non-*C. difficile* diarrhoeal pathogens using stool culture and immunochromatographic tests for detection of rotavirus, norovirus, and adenovirus (Immunoquick NoRotAdeno, BIOSYNEX). Recommendations for CDI were followed [16].

#### Informed consent

Ethical approval was obtained from the University Hospital Ethics Committee. Written informed consent for the University Hospital to sample stools was received from parents or legal guardians at the time of hospital admissions.

#### Statistical analysis

Children with diarrhoea were compared to control patients with respect to a number of variables. Fisher's exact test was used as appropriate for comparing categorical variables. Significance level for statistical hypothesis testing was  $\alpha = 0.05$ .

**Table 1.** Demographic, clinical and laboratory characteristics

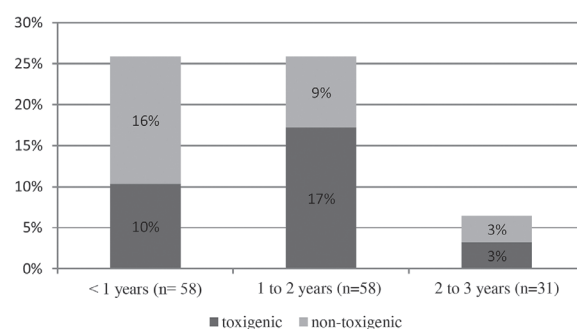
Test Results	Patients with diarrhoea (n = 147)	Control Patients (n = 75)	
Median age (range) in years	1.2 (from 3 days to 34 months)	1.3 (from 3 days to 35 months)	
Female (%)	58 (39.5)	34 (45.3)	
Male (%)	89 (60.5)	41 (54.7)	
≤ 28 days (%)	4 (2.7)	3 (4.0)	
≤ 1 years (%)	54 (36.7)	24 (32.0)	
1 to 2 years (%)	58 (39.5)	23 (30.7)	
2 to 3 years (%)	31 (21.1)	25 (33.3)	
Duration of hospitalisation - Mean (± SD) <sup>a</sup> in days	4.3 (± 2.2)	6 (± 2.5)	
Rectal body temperature > 37.5 °C	93 <sup>b</sup>	/	
Vomiting (n)	14 <sup>c</sup>	/	
Bloody diarrhoea (n)	17 <sup>d</sup>	/	
Resolution of diarrhoea - Mean (± SD) in days	1.6 (± 1.44)	/	
Risk Factors			p-Value <sup>e</sup>
Antibiotic exposure <sup>e</sup> (%)	13 (8.9)	19 (25.3)	< 0.05
Hospitalisation <sup>f</sup> (%)	15 (10.2)	12 (16)	0.28
Hirschsprung's disease (n)	1	1	/
Stoma in gastrointestinal system (n)	2	0	/
CMPA <sup>g</sup> (n)	6 <sup>h</sup>	3	/
surgery procedure (n)	1	3	/
central hypotonia (n)	1	0	/
Laboratory results of <i>C. difficile</i>			
GDH <sup>i</sup> /toxin-	118	55	
GDH+/toxin-	15	13	
GDH+/toxin A,B+	4	2	
GDH+/PCR <sup>j</sup> gene B+	11	6	
GDH+/PCR gene B, BT <sup>k</sup>	2	0	

<sup>a</sup>SD - standard deviation<sup>b</sup>93/139, missing data for 8 children<sup>c</sup>14/137, missing data for 10 children<sup>d</sup>17/136, missing data for 11 children<sup>e</sup>antibiotic exposure 12 weeks prior to stool collection<sup>f</sup>hospitalisation 60 days prior to stool collection<sup>g</sup>cow's milk protein allergy (CMPA)<sup>h</sup>1x suspected CMPI<sup>i</sup>GDH - Glutamate-dehydrogenase<sup>j</sup>PCR - Polymerase Chain Reaction<sup>k</sup>binary toxin<sup>l</sup>significance level of  $\alpha = 0.05$  for statistical hypothesis testing

## RESULTS

Patient demographic data, clinical characteristics, and laboratory results of our samples are presented in Table 1. One hundred and forty-seven patients with diarrhoea and seventy-five control patients were enrolled in the study. The median age was 1.2 years (range from 3 days to 34 months) and 1.5 years (range from 3 days to 36 months) in the first and control groups, respectively. The stool samples were tested for *C. difficile* over a 13-month period. The results according to sex and age subgroups are not major and are presented in Table 1 and Figure 1.

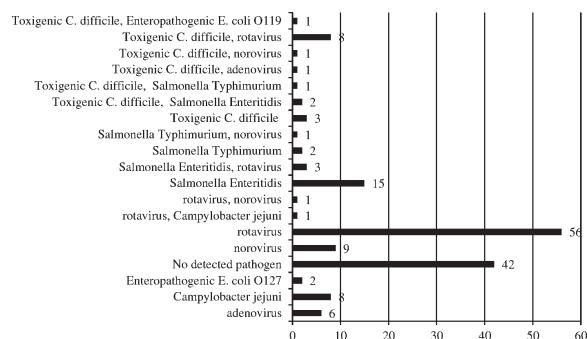
Among symptomatic children, 11.6% (n = 17) were toxigenic *C. difficile* isolates (see Table 1), including 2.7% (n = 4) of samples detected by IT and 7.5% (n = 11) con-

**Figure 1.** Toxigenic *C. difficile* rate in each age group (neonates are not included separately)

Diagnosis of CDI is based on the diagnostic two step-screening algorithm, which was described in "materials and methods" (combination of IT and PCR).

Abbreviations: CDI - *Clostridium difficile* infection, IT - immunochromatographic test, PCR - polymerase chain reaction

firmed by PCR, while in 1.4% (n = 2) of cases the binary toxin was revealed but no deletion in nucleotide 117 was presented. The CDI definition was fulfilled in 2% (n = 3) of patients. One child might be categorized under hospital-acquired *C. difficile*. By comparison, in the control group, of the total 10.7% (n = 8) toxigenic *C. difficile* isolates, 2.7% (n = 2) were toxin A/B positive while gene for toxin B was detected in 8% (n = 6) of stools. Binary toxin and deletion in nucleotide 117 were not recorded in any samples.

**Figure 2.** The aetiology of diarrhoea among hospitalized children (n = 147)

The numbers of agents, some children had multiple aetiologies.

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**Table 2.** The group of symptomatic children with positive toxigenic *C. difficile* in stool samples

Test results	Absolute number
Number of patients	17
Female	9
Male	8
≤ 28 days	0
≤ 1 years	6
1 to 2 years	10
2 to 3 years	1
Duration of hospitalisation – Mean (± SD <sup>a</sup> ) in days	3.8 (± 1.0)
Rectal body temperature > 37.5 °C	13
Vomiting	6
Bloody diarrhoea	2
Resolution of diarrhoea <sup>b</sup> – Mean (± SD) in days	1.7 (± 1.3)
<b>Risk factors<sup>c</sup></b>	
Antibiotic exposure <sup>d</sup>	3
Hospitalisation <sup>e</sup>	2
Hirschsprung's disease <sup>f</sup>	1
Stoma in gastrointestinal system	1
Central hypotonia	1
<b>Laboratory results of <i>C. difficile</i></b>	
GDH <sup>g</sup> +/toxin A,B+	3
GDH+/PCR <sup>h</sup> gene B+	14
CDI group <sup>i</sup>	
Cases <sup>j</sup>	3
Recurrences	0
Specific CDI treatment	1 <sup>k</sup>
Community-acquired	2
Hospital-acquired	1

<sup>a</sup>SD – standard deviation<sup>b</sup>self-limited resolution without specific CDI treatment<sup>c</sup>some patients had more risk factors<sup>d</sup>antibiotic exposure 12 weeks prior to stool collection<sup>e</sup>hospitalisation 60 days prior to stool collection<sup>f</sup>the combination of partial colectomy (with a colostomy) and fidaxomicin 200 mg twice daily for 10 days<sup>g</sup>GDH – Glutamate-dehydrogenase<sup>h</sup>PCR – Polymerase Chain Reaction<sup>i</sup>CDI (*Clostridium difficile* infection) definition is determined in the „Materials and Methods“<sup>j</sup>one boy (age: 2.1 years), two girls (age: 1.3 and 1.5 years, respectively)

Altogether non *C. difficile* intestinal pathogens were discovered in 106 stool samples (data are summarized in Figure 2) 72% (76/106) of which were viral agents. Of 17 positive *C. difficile* toxin specimens, a parallel bowel pathogen was detected in 14 stools (rotavirus, *n* = 8; norovirus, *n* = 1; adenovirus, *n* = 1; *Salmonella enteritidis*, *n* = 2; *Salmonella typhimurium*, *n* = 1; enteropathogenic *Escherichia coli*, *n* = 1) while no pathogen was revealed in the 3 remaining stool samples. *Salmonella enteritidis* was cultivated from 1 infant with diarrhoeal disease (GDH+/

PCR gene B+), but the same bacterium had been detected during a previous hospitalization. The infection could have been due to intestinal carriage, the patient was excluded from the CDI group.

Among clinical symptoms in children with toxigenic *C. difficile* isolates (*n* = 17) during hospitalisation (the results are presented in Table 2), the majority (*n* = 13) had fever, vomiting occurred in 6 patients, and bloody diarrhoea was observed in 2 children. Considering the age group concerned, abdominal pain was not assessed. Seven patients had common complications of dehydration – hypoglycaemia, mineral disturbances, high levels of nitrogen-containing compounds (urea, creatinine). In one child with underlying comorbidities (Down syndrome, Hirschsprung's disease), transversostomy, repeated hospitalizations, and antibiotic treatment in last month (cefotaxime, metronidazole, gentamicin, co-trimoxazole), intensive care unit therapy and specific CDI treatment (colectomy + fidaxomicin) of the child was required. A colonoscopy was conducted, whereby inflammatory changes and superficial ulcers in the intestinal lining were described. In other children, endoscopy, X-ray, and ultrasonography were not required. The patient met the CDI criteria for severe form. The remaining 2 patients (who met the criteria for likely CDI) did not need treatment because their diarrhoeal infections were self-limited. One infant had gastrostomy and VACTERL syndrome. No risk factor was revealed in the remaining case (other risk factors are listed in Table 1 and Table 2).

## DISCUSSION

In the chosen age group, the colonization rate has been shown to be high despite low morbidity [6, 7, 12, 13, 15]. Although earlier national guidelines had discouraged stool testing for *C. difficile* in younger infants, the updated recommendations have considered association with CDI in every child group. Unfortunately the global guidelines are typically based on adult patients, not children [6, 7, 10, 15, 17–19].

The human intestine is sterile at birth. By 1 year of age, child and adult intestinal microbiota become similar [20]. Studies reported an average *C. difficile* colonization rate of 37% of stools in neonates (in the first 28 days after birth) with wide variations [20,21] Between 1 and 12 months of age, colonization declines to an approximate rate of 10% of children. After the first year of life, the asymptomatic carriage appears to approximate the colonization rate in adults (3–4%) [2, 20, 22]. The high colonization rate in infancy has been attributed to lower capacity of bowel microflora to inhibit *C. difficile* growth [23]. It is also affected by infant nutrition [20, 24]. In our study, the total prevalence of toxigenic *C. difficile* strains in the diarrhoeal group compared to that of the control group was 11.6% (17/147) vs. 10.7% (8/75), which was statistically insignificant (*p* > 0.9999). We suggest there is a minimal causal relationship between the presence of toxin and diarrhoea. Our findings could be influenced is fact that the majority of those individuals enrolled were otherwise healthy infants with an absence of risk factors. In other study, Sathyendran et al. had reported 15.6% positive stool specimens among patients with median age of 1.2 years, of which 28% were toxin-positive and 72% were PCR positive [15]. The toxigenic *C. difficile* rate had been in-



fluenced by the recruited sample of children. The detection of toxigenic *C. difficile* seems to occur fairly frequently, and positive results must be interpreted while viewing each case in a comprehensive manner.

It is generally presumed that the majority of CDI cases in children are attributable to colonization. No clear definition of CDI in this age group yet exists [7]. Detection of CDI has been complicated by an absence of reliable faecal biomarkers for CDI [25]. In our study, the overall prevalence of CDI in children with diarrhoea was 2% (3/147), while 17.6% (3/17) of children (with positive toxin/gene for toxin B in faeces) met the criteria for CDI. Most studies have included epidemiology of toxigenic *C. difficile* strains without evaluating the individual CDI cases or have been conducted among children of different age groups (different research design). Sathyendran et al. published outcomes using the same molecular methods and algorithm, determining that 14% (46/320) of children (median age of 1.2 years) met criteria for CDI [15]. This distinction could be due to a higher proportion of children within risk groups. Sixty-three percent of those patients were treated as compared to 33% (1/3) in our study. The main contributing reason can be seen in the elected sample of patients [15].

Although older studies had reported a high colonization rate of toxigenic strains in neonates [20], our study found no neonates testing positive for the presence of toxigenic *C. difficile* strains. Unfortunately, the number of enrolled neonates was low ( $n = 7$ ). Similar findings had been presented by Rousseau et al. [21], with only 4.8% of positive samples, and by Sathyendran et al. [15], where the prevalence remained at 2.6% (1/39) despite a longer hospitalization period. Those authors had hypothesized prevention practices (e.g. hand hygiene) to be a contributing factor [15]. Moreover, the current diagnostic algorithm should be more sensitive. We would also point out the possibility of transient carriage [20] and emphasize the low number of published cases of pseudomembranous colitis in neonates [5].

According to age subgroups in both groups, detection of toxigenic strains was around 10% in infants ( $\leq 1$  year of age), which was comparable to the conclusions from other studies [15, 20, 21]. None of the aforementioned children with toxigenic *C. difficile* strains fulfilled the CDI definition. The American Academy of Pediatrics recommends that children younger than 1 year of age not be tested unless a child has bowel motility problems or is situated in an outbreak situation [10]. We could not agree more with this statement, because our records did not identify these factors in the enrolled children. In the older age population (1–2 years of age), high toxigenic *C. difficile* prevalence in the diarrhoeal case group compared to the control group (16.9% and 8.7%, respectively) was surprisingly revealed, but only 20% of those (2/10) in fact met the CDI criteria. In view of the fact that most of the recruited children were otherwise healthy and with an absence of risk factors, the high rate remains a mystery in comparison to other reported data [20, 21]. This conclusion could be partially influenced by the small number of children in the subgroups. The oldest age group exhibited the presumptive low toxigenic rate [20]. Statistical analysis was not beneficial for comparison of the subgroups.

A majority of reported studies highlight previous antibiotic exposure and repeated hospitalization as significant risk factors [6, 7, 15, 17]. An absence of risk factors

in the diarrhoeal group can be explained by the method of patient selection, as there is a predominance of viral infections in the selected age groups and low morbidity in the general child population. The significantly higher rate of antibiotic exposure among controls ( $p < 0.05$ ) might be due to the high level of antibiotic prescriptions among children with respiratory infections during winter and spring months (data are not included in the text). Meanwhile, cow's milk protein allergy is a controversial risk factor in relation to CDI [6].

Our study has a number of limitations. The timing of stool collection differed between the two groups – symptomatic patients ( $< 48$  hours of admission) and asymptomatic patients (at any time during admission). This difference could have impacted positive results in control group (duration of hospitalization could increase *C. difficile* colonisation). The majority of those individuals enrolled were otherwise healthy children with an absence of risk factors. Statistical analysis was beneficial only for comparison of the two main groups, while the subgroups, genders and numbers of risk factors were too small for meaningful statistical analysis. Moreover, the reliance on clinical data and medical history could be confounding. The criteria used for determining CDI had limitations. Defecation frequency before hospitalization was based on the parents' perceptions. Identification of viral pathogens was limited to three viruses – an unknown pathogen could not be excluded. Diarrheagenic parasites were not identified because of their presumed low incidence in developed countries [26]. Generally, CDI severity was not appraised due to the low number CDI cases. The research was conducted solely in our department. It could be difficult to follow the study because of its design.

## CONCLUSION

Our findings indicate that *C. difficile* is a significant but infrequent pathogen in hospitalized children with diarrhoea. Essentially healthy children usually exhibit a favourable course of infection. Cases should be evaluated individually in the absence of optimal CDI diagnostics in infants. We recommend not to examine the youngest paediatric populations if children do not exhibit the aforementioned risk factors. Stool testing for non *C. difficile* bowel pathogens should be emphasized for children  $\leq 3$  years of age in cases of *C. difficile*-positive stool samples. There is a need for more contemporary works on *C. difficile* colonization/infection in infants and young children.

## List of abbreviations

BT	– Binary toxin
<i>C. difficile</i>	– <i>Clostridium difficile</i>
CDI	– <i>Clostridium difficile</i> infection
EPEC	– Enteropathogenic <i>Escherichia coli</i>
GDH	– Glutamate-dehydrogenase
IBD	– Inflammatory bowel disease
ICU	– Intensive care unit
IT	– Immunochromatographic test
PCR	– Polymerase Chain Reaction
VACTERL (acronym)	– refers to the nonrandom co-occurrence of birth defects: Vertebral anomalies, Anal atresia, Cardiac defects, Tracheoesophageal fistula and/or Esophageal atresia, Renal & Radial anomalies and Limb defects

## PŮVODNÍ PRÁCE

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## Conflict of interest statement

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## Author contributions

All of the authors of this manuscript are responsible for the reported findings. VM collected all clinical data of practical approach. All authors participated in the searching of the published literature for this review; analysis and interpretation of the data; drafting and/or revising and/or making intellectual contributions to the content of the manuscript; All authors approved the final version of the manuscript submitted for publication.

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