



AKADÉMIAI KIADÓ

Acta Microbiologica et
Immunologica Hungarica

67 (2020) 2, 79-86

DOI: [10.1556/030.66.2019.024](https://doi.org/10.1556/030.66.2019.024)

© 2019 Akadémiai Kiadó, Budapest

Growing consumption of antibiotics and epidemiology of *Clostridioides difficile* infections in Poland: A need to develop new solutions

ESTERA JACHOWICZ^{1,2*}, MONIKA POBIEGA²,
ANNA RÓŻAŃSKA¹ and JADWIGA WÓJKOWSKA-MACH¹

¹Department of Microbiology, Faculty of Medicine, Jagiellonian University Collegium Medicum, Kraków, Poland

²Biophage Pharma SA, Kraków, Poland

Received: May 30, 2018 • Accepted: July 19, 2019

ORIGINAL ARTICLE



ABSTRACT

Clostridioides (formerly *Clostridium*) *difficile* infections (CDIs) are becoming more common and more serious. *C. difficile* is the etiologic agent of antibiotic-associated diarrhea, pseudomembranous enterocolitis, and toxic megacolon while CDIs recur in 7.9% of patients. About 42.9 CDI cases/10,000 patient-days are diagnosed each day in Europe, whereas in Poland 5.6 CDI cases/10,000 patient-days are reported; however, the median for European countries is 2.9 CDI cases/10,000 patient-days. Epidemiology of CDIs has changed in recent years and risk of developing the disease has doubled in the past decade that is largely determined by use of antibiotics. Studies show that rate of antibiotic consumption in the non-hospital sector in Poland is much higher than the European average (27 vs. 21.8 DDD/1,000 patient-days), and this value has increased in recent years. Antibiotic consumption has also increased in the hospital sector, especially in the intensive care units – 1,520 DDD/1,000 patient-days (ranging from 620 to 3,960 DDD/1,000 patient-days) – and was significantly higher than in Germany 1,305 (ranging from 463 to 2,216 DDD/1,000 patient-days) or in Sweden 1,147 (ranging from 605 to 2,134 DDD/1,000 patient-days). The recent rise in CDI incidence has prompted a search for alternative treatments. Great hope is placed in probiotics, bacteriocins, monoclonal antibodies, bacteriophages, and developing new vaccines.

KEYWORDS

Clostridium difficile infections, bacteriophage, probiotics, epidemiology, antibiotic consumption

INTRODUCTION

Clostridioides (formerly *Clostridium*) *difficile* is a Gram-positive, anaerobic, spore-forming rod causing disease in humans and animals, present in the gut of 14% of healthy people [1]. This pathogen was first described in 1935 and called *Bacillus difficilis* – a rod that is difficult to grow [2].

At present, *C. difficile* infection (CDI) poses an enormous challenge to contemporary medicine. The number of CDIs is constantly growing and their treatment is becoming increasingly more difficult and more expensive. *C. difficile* is the main etiologic agent of antibiotic-associated diarrhea; according to literature data, as many as 16%–35% of them are caused by *C. difficile* [3]. Symptoms of CDI are watery diarrhea, abdominal pain, vomiting, nausea, and fever [4]. Severe forms of infection may lead to pseudomembranous colitis and toxic megacolon (*megacolon toxicum*) [5]. According to literature reports, 25% of patients with CDI do not respond to antibiotic treatment and approximately 7.9% relapse [6, 7]. Severe complications, such as surgeries, *megacolon toxicum*, or even patient death, are determined in 16.7% of all CDI cases [7].

* Corresponding author:

Estera Jachowicz
Department of Microbiology, Faculty of
Medicine, Jagiellonian University
Collegium Medicum, Gołębia 24, 31-
007, Kraków, Poland
Biophage Pharma SA, Kraków, Poland
Phone: +48 126 332 567;
Fax: +48 124 233 924
E-mail: estera.jachowicz@doctoral.uj.edu.pl

Hospitalized patients are the most exposed to CDI. Seventy percent of all CDIs are associated with hospital treatment [7]. The most significant risk factor for the development of CDI is taking antibiotics that interfere with the natural intestinal flora [8]. The antibiotics with the highest risk of developing CDI include, first of all, clindamycin that increases the risk of developing CDI approximately 20-fold; while the frequently used fluoroquinolones, cephalosporins, and carbapenems increase this risk fivefold (compared to people not treated with an antibiotic) [9]. Restoring the patient's microbiota to its original state, the one from before antibiotic therapy, takes up to 2 years [10]. There are attempts to look for relationships between development of CDI and the application of drugs that alter the pH of gastric juice, such as H2 blockers and proton pump inhibitors. However, we lack enough evidence to confirm this hypothesis [11]. A meta-analysis made by Furuya-Kanamori et al. demonstrates that exposure to corticosteroids increases the risk of CDI, but more accurate studies of this phenomenon are necessary [12]. Other significant risk factors for CDI are immunosuppression, e.g., in transplant patients, especially heart transplants, as well as cystic fibrosis, vitamin D deficiency, intestinal diseases, age over 65 years, low levels of anti-toxin A IgG, and previous CDI: after the first episode, the risk increases to 45%, and by the third it is 65% [6].

EPIDEMIOLOGY

Epidemiology of CDI has changed significantly in recent years. It is estimated that the past decade brought a twofold increase in the disease risk, but this change applies especially to the clinical picture as the USA, for instance, experienced a threefold rise in the group of patients requiring hospitalization due to community-acquired CDI (CDI-related hospitalizations) and additionally the disease affects people who have been considered earlier not particularly vulnerable to CDI, among others, healthy persons living in the community and peripartum women [13, 14].

In the USA, the incidence of hospital-acquired CDI (HA-CDI) amounted to 9.3/10,000 patient-days and the community-acquired 4.8/10,000 patient-days [15]. In European countries, the total CDI incidence of HA-CDI was 2.4 cases/10,000 patient-days.

Epidemiology of the disease has changed, which is mainly due to the spread of the highly virulent strain NAP1, rare before 2001 and now responsible for many hospital epidemics [12]. In Europe, the most frequently isolated ribotypes are RT027 (22.9%), RT001 (7.5%), RT014 (6.7%), RT078 (5.1%), RT002 (4.2%), and RT020 (4.2%), which also correlate with the results obtained by Pituch et al. – ribotype RT027 is the most commonly isolated ribotype in Poland [7, 16].

The CDI mortality is high, in the USA, an active population- and laboratory-based study in 2011 identified 5.4% of *C. difficile* fatal case rate, whereas in the EU it was 3.9% [7].

EPIDEMIOLOGY OF CDI IN POLAND

The Polish total HA-CDI incidence was 6.2/10,000 patient-days [7]. Other Polish data, from single-center studies, reported the CDI incidence in intensive care units (ICUs) at 13/10,000 and 10.6/10,000 patient-days [17, 18]. However, such a high number of CDI has been observed in Poland for a short time – the incidence rates were 6.1, 8.6, and 9.6 CDI per 10,000 patient-days in 2011, 2012, and 2013, respectively [19].

In Poland, the first infection caused by 027 ribotype was detected in a hospital in Warsaw in 2005–2006. This isolate had *tcdA*, *tcdB*, and binary toxin genes (*cdtA* and *cdtB*) [20]. Studies conducted in 2011–2013 in 13 Polish hospitals have shown that ribotype 027 is the most frequent (62%) type related to CDI in Poland [21]. The situation is ever worse in Silesia where in 2017 more than 80% of CDIs are connected to ribotype 027 [22]. In a Polish single-center study, the crude mortality rate of HA-CDI was 12.9% in medical wards and 27.7% in the ICU setting [23].

According to Barlam, the risk of developing CDI has, among others, a direct close relationship with the rational use of antibiotics: “Antibiotic Stewardship Program” decreases the risk of CDI in particularly susceptible patients [24]. This is consistent with observations concerning the patients in Poland, where the situation in regard to CDI, and to antibiotic consumption, is serious [25]. A single-center study in a Polish ICU, where the consumption of antibiotics amounted to 1,913 DDD (defined daily dose)/1,000 patient-days, with a simultaneous exceptionally high and disturbing percentage of high level of resistant strains, found a very high CDI incidence: 10.6/10,000 patient-days [18]. Still, the unit examined was not an exception. A multicenter study conducted in Polish ICUs indicated consumption of antibiotics at an average level of 1,520 DDD/1,000 patient-days (ranging from 620 to 3,960 DDD/1,000 patient-days) and was significantly higher than analogous rates recorded in, e.g., Germany, 1,305 (463–2,216), or in Sweden, 1,147 (605–2,134 DDD/1,000 patient-days) [26–28]. What is also worrying is the high percentage of carbapenems (17%) and quinolones (14%) that was found, while it is precisely the proportion of broad-spectrum antibiotics and fluoroquinolones in the total antibiotic consumption that constitutes a parameter evaluating the quality of prescribing antibiotics [28].

Analysis of the consumption of antibiotics in hospitals using a different methodology, considering DDD consumption per 1,000 inhabitants per day (person-days and patient-days) (ECDC program), showed consumption slightly lower than the average for all European countries (1.78 vs. 2.03 DDD/1,000 patient-days), however, with upward trend in the period 2014–2017 (data for previous years are not available for Poland) [29]. In turn, the consumption of antibiotics calculated according to the same methodology for the Polish non-hospital sector in 2017 showed a value 30% higher than the European average (27 vs. 21.8 DDD/1,000 patient-days). The upward trend in the consumption of antibiotics in the non-hospital sector was recorded according to ECDC data even in the longer term (2007–2016, for which published data



are available); additionally, there was an increasing trend of the share of fluoroquinolone consumption [25].

In Poland, *C. difficile* resistance to commonly used antibiotics is high; according to reports of Lachowicz et al., 85.5% studied isolates were resistant to erythromycin, whereas 27.7% had high-level clindamycin resistance, having minimum inhibitory concentrations (MICs) greater than 256 mg/L. All strains were ciprofloxacin-resistant, 83.1% were moxifloxacin-resistant, and 87.9% strains were imipenem-resistant. All strains were sensitive to tigecycline, metronidazole, and vancomycin. All ribotype 027 and 176 *C. difficile* isolates demonstrated high-level resistance to erythromycin (MIC \geq 256 mg/L); multidrug resistance (resistance to at least three classes of antimicrobial agents) was observed in 85.5% of toxigenic *C. difficile* strains [30].

PATHOMECHANISM OF INFECTIONS

CDI is generally contracted endogenously and by fecal–oral route, primarily through the hands of medical staff [1, 31]. *C. difficile* has many virulence factors, including ability to synthesize toxins A, B and binary toxin (CDT), presence of cilia, S-layer, Cwp66 adhesin, GroEL heat shock proteins and fibronectin-binding proteins (Fbp68), as well as the ability to form biofilm and capacity for sporulation [32–37]. Intestinal diseases are caused only by the *C. difficile* strains that produce toxins A and/or B; there is also a correlation between the amount of toxins in the intestines and the disease duration [38]. This has been proven by introducing insertions into the genes *tcdA* and *tcdB* of virulent bacteria, which led to loss of toxin production and absence of disease symptoms [39]. It was an enormous challenge for contemporary medicine when, in the year 2000, the strain *North American Pulsed Field Type 1* (NAP1) appeared; also known as B1/NAP1/027, or in other words ribotype PCR 027, it is characterized by increased virulence, resistance to fluoroquinolones, formation of a greater number of spores, production of larger amounts of toxins A and B and binary toxin [40, 41]. Binary toxin causes disease in people not previously exposed to antibiotic therapy or hospitalized; the incidence of infections with strains producing binary toxin is estimated at 11%–20.5% in children [12]. Toxins produced by *C. difficile* (A, B, and binary) enter enterocytes and then damage their structure; consequently, albumins, electrolytes, and water are lost through the cell membrane [42, 43]. In a healthy individual, the first line of defense against *C. difficile* is the intestinal environment, the mucus present therein, along with antimicrobial substances suspended in it, as well as bactericidal peptides and immunoglobulins, and the natural microbiota.

There is a high percentage of *C. difficile* carriers among children; however, it does not correlate with CDI incidence due to immaturity of mucous membranes and microbiota in young children and the resulting lack or insensitivity of the receptors to *C. difficile* toxin activity. It is only in children over 2 years of age when the microbiota begins to resemble to the one in adults [44].

DIAGNOSIS AND TREATMENT

Due to serious complications, rapid and accurate diagnosis of infection and appropriate treatment play a vital role. The basic tests are the ones that detect glutamate dehydrogenase (GDH), immunochromatic tests, the ones detecting toxins, and the nucleic acid amplification test [12]. GDH is an enzyme produced by toxinogenic and non-toxinogenic strains of *C. difficile*. It is noteworthy that antibodies against GDH can cross-react with other bacterial enzymes of the genus *Clostridium* [45].

Due to the anaerobic growth conditions of *C. difficile*, its drug sensitivity is very rarely determined in routine diagnostics. According to the European Committee on Antimicrobial Susceptibility Testing (EUCAST), for epidemiological and clinical purposes, sensitivity to eight antibiotics should be determined (moxifloxacin, vancomycin, tigecycline, daptomycin, fusidic acid, metronidazole, rifampicin, and fidaxomicin). There are big problems in terms of interpretation of diagnostic results; therefore, EUCAST does not recommend routine *C. difficile* susceptibility testing (http://www.eucast.org/clinical_breakpoints/, accessed on: December 27, 2018). In 2016, in Europe, 4.6% of *C. difficile* strains isolated were resistant to metronidazole, 69.4% to moxifloxacin, and one isolate was resistant to vancomycin [7].

Standard CDI therapy involves administration of vancomycin. Treatment with metronidazole is only used in mild cases of infection if the aforementioned antibiotics are unavailable [12]. With mild forms of CDI, the effectiveness of both antibiotics is comparable; however, with severe infections, it was shown that vancomycin proved more effective [46, 47].

IN SEARCH OF NEW SOLUTIONS

A new drug against *C. difficile* in test trials is cadazolid, which inhibits the synthesis of bacterial proteins but is not absorbed from the gastrointestinal tract. The antibiotic demonstrated good activity against *C. difficile*, also against the hypervirulent strain [48]. Another antibacterial agent studied is SMT19969 [2,2'-bis(4-pyridyl)3H,3'H 5,5-bibenzimidazole], which exhibits activity similar to vancomycin in studies [49]. A randomized research with β -lactamase ribaxamase (SYN-004) demonstrated that patients treated intravenously with ceftriaxone along with ribaxamase in lower respiratory tract infections and oral therapy reduced the incidence of *C. difficile* compared to placebo (risk reduction 2.4%). The ribaxamase may prevent CDI in patients treated with intravenous β -lactam antibiotics [50]. The literature lists alternative methods for CDI treatment, such as the use of probiotics, vaccines containing toxins A and B, fecal transplant, and bacteriophage therapy. To investigate treatment options, as well as infection prevention, CDI animal models were created (examples of model animals are Syrian hamsters, mice, pigs, and rabbits), which demonstrate a similar course of the disease as the one in humans [51–53].

Another treatment method for recurrent infections that demonstrates the best results (effectiveness of about 90%) is fecal microbiota transplantation (FMT) carried out in patients with intestinal diseases, irritable bowel syndrome, or nervous system diseases. Usually, fecal samples are collected from a healthy family member as the genetic similarity is reflected in the composition of the microbiota [54]. The doses are usually administered through a lower gastrointestinal series or by means of a colonoscope [55]. The advantages of FMT include low costs of treatment and its effectiveness, whereas the disadvantages are variability of the transplanted microbiota and patients' mental resistance. FMT can take various forms, from a suspension in sodium chloride, through lyophilized powder, to an encapsulated formulation [55].

It is also recommended to use probiotics as adjunctive therapy for CDI treatment. A good probiotic must be resistant to bactericidal substances secreted by other microorganisms, capable of survival in the gastrointestinal tract, and should modulate the host immune response [56]. The most investigated probiotic strain, applicable for *C. difficile* treatment, is the yeast *Saccharomyces boulardii* [57]. Literature data indicate its positive influence on inflammatory bowel disease, a good capacity to colonize the intestine, and the ability to disappear from the body after 5 days upon stopping the supplementation [58]. Through suppression of NF- κ B activity, it inhibits IL-8 production, which is one of the mediators of the inflammatory response arising due to CDI, and limits the binding of toxin A to receptors, which was demonstrated during studies in a CDI rat model [58, 59]. According to the studies by Plummer et al. [60], people administered with probiotics showed colonization with the pathogen but the toxin was not detected in their feces, which may point to the fact that probiotics can neutralize the toxin. Literature reports reveal higher proportions of patients with CDI in the placebo groups than in the groups taking probiotics [60, 61]. Apart from *S. boulardii*, other probiotic bacteria listed as adjunctive to CDI treatment are *Lactobacillus acidophilus*, *Lactobacillus rhamnosus*, *Lactobacillus casei*, and *Lactobacillus plantarum* [62]. *L. acidophilus* LA-5 alleviates the symptoms of CDI, reduces the concentration of toxins, and improves the histopathological picture of the intestines, whereas *Lactococcus lactis* SL3 reduces the viability of *C. difficile* strains [63, 64]. It should be noted that some strains of the genus *Lactobacillus* are naturally resistant to vancomycin – the antibiotic used for therapy of CDI [65]. Hell et al. [66] gave a multistrain preparation containing bacteria of the genus *Bifidobacterium*, *Lactobacillus*, and *Enterococcus* to 10 CDI-positive people and obtained *C. difficile* eradication in 9 of them, while 1 patient died of pneumonia (not associated with CDI). It should be emphasized that the effect of probiotic strains on *C. difficile* is strain-dependent. A vital role in *C. difficile* eradication is played by H₂O₂ produced by probiotic strains [8]. Due to a limited number of studies and small patient groups, there are no official recommendations for the use of probiotics in the treatment and adjunctive therapy for CDI. At present, there are ongoing clinical studies on the use of probiotics

(e.g., *Lactobacillus* spp. and *Bifidobacterium* spp.) in CDI (www.clinicaltrials.gov: NCT03368105 and NCT03647995). It is worth noting that the strains of the *Lactobacillus* sp. produce other antibacterial substances in microaerobic and anaerobic environment, hence the conflicting results of probiotic antagonism against *C. difficile*. In *in vivo* studies, besides a different environment, the eradication of *C. difficile* is also influenced by the immune system and the local microbiota [67]. It should also be pointed out that there were cases of mycoses and fungemia caused by *S. boulardii* [68].

Bacteriocins, produced by certain bacteria, can also be applied in the fight against *C. difficile*. Lacticin 3147 and *L. lactis* niacin demonstrate a bactericidal effect against *C. difficile* but also against other bacteria, including those non-pathogenic, in a way similar to actagardine A and LFF 571. Thuricin CD, produced by *Bacillus thuringiensis*, a bacterium colonizing the intestine, has an inhibitory effect on *C. difficile* that is comparable to metronidazole or vancomycin; however, it shows no toxic effects on other bacteria naturally present in the intestine [69].

Intensive studies are being conducted into monoclonal antibodies against toxins A and B as well as vaccines against *C. difficile* antigens FliD, FliC, and Cwp 84 [70–73]. Research carried out by Wilcox et al. on the influence of bezlotoxumab and actoxumab monoclonal antibodies confirm the safety of their use. The scientists have recorded a smaller number of relapses (by around 10%) in the group taking antibodies than in the placebo group [74]. In addition, there is work underway on vaccines against *C. difficile* and the literature data available suggest that they are safe [59]. Vaccines containing toxoids showed no serious post-vaccination effects. Complications typical of vaccines with aluminum adjuvant have been observed: 46% of people who received the vaccine and 15% of people in the placebo group had a reaction [75]. People vaccinated against *C. difficile* toxins A and B and aged 18–55 years exhibited a greater immune response against toxin A than B, contrary to the situation in the age group of over 65 years. Administering another dose of the vaccine resulted in increased seroconversion of antibodies against toxin A in elderly patients [76]. In the next phase of clinical testing of the vaccine, de Bruyn et al. [77] obtained an immune response against toxin A in 97% and against toxin B in 92% of people. The vaccines deserve special attention, the research into them is in advanced clinical stages and their routine application would allow to prevent the occurrence of infections caused by *C. difficile*.

Another alternative method for CDI elimination is phage therapy. However, to date, the bacteriophages isolated have displayed a lysogenic life cycle. There are also literature reports on the role of prophages in the expression of genes for *C. difficile* toxins A and B [78]. According to Rea et al., prophages affect the host phenotype and toxin expression, which may result in increased virulence of the pathogen. The bacteriophages obtained to date have been isolated after treating the host strain with mitomycin C [79]. Difficulties to isolate the ones displaying a lytic cycle are explained by the

fact that the prophage stage is conducive to the survival of the phage, which is associated with *C. difficile* spore production. The second argument is the fact that a large number of prophages in the bacterial genome reduce its susceptibility to new bacteriophage infections (mechanism of superinfection) [79]. In their latest studies, Nale et al. demonstrated bactericidal activity against *C. difficile* bacteria of a phage cocktail containing the family *Myoviridae*: CDHM 1, 2, 5, 6 *in vitro* inhibition of biofilm formation by *C. difficile* and *in vivo* reduction in colonization of the hamster gut [80, 81]. Phago-therapy has therapeutic potential in terms of treatment of CDI; however, to date, there are still many unknowns regarding this issue.

PREVENTION

According to European Society of Clinical Microbiology and Infectious Diseases (ESCMID) Study Group for *C. difficile* (ESGCD), the cornerstones of CDI prevention and control remain appropriate microbiological testing practices based on a two-stage test, CDI surveillance with regular feedback, standard and contact precautions with special emphasis on hand hygiene, use of personal protective equipment and environmental disinfection, antimicrobial stewardship and education of healthcare workers as well as CDI cases and hospital visitors regarding CDI prevention [82].

SUMMARY

Both Europe and Poland are currently facing the enormous problem of high CDI incidence involving the highly pathogenic ribotype 027. Treatment of CDI is lengthy, costly, and sometimes, unfortunately, ineffective. Therefore, it is so important to perform intensive actions to prevent CDI and introduce modern effective methods for their treatment simultaneously.

Conflict of Interest: The authors declare no conflict of interest regarding the publication of this paper.

REFERENCES

- Murphy, C., Veron, M., Cullen, M.: Intravenous immunoglobulin for resistant *Clostridium difficile* infection. *Age Ageing* **35**, 85–86 (2006).
- Hall, I. C., O'Toole, E.: Intestinal flora in newborn infants with a description of a new pathogenic anaerobe, *Bacillus difficilis*. *Am J Dis Child* **49**, 390–402 (1935).
- Aslam, S., Hamill, R. J., Musher, D. M.: Treatment of *Clostridium difficile*-associated disease: Old therapies and new strategies. *Lancet Infect Dis* **5**, 549–557 (2005).
- Pituch, H., Albrecht, P.: *Clostridium difficile* – narastający problem diagnostyczny i terapeutyczny [*Clostridium difficile* – A growing diagnostic and therapeutic problem]. *Gastroenterol Klin* **5**, 40–51 (2013).
- Zanella Terrier, M. C., Simonet, M. L., Bichard, P., Frossard, J. L.: Recurrent *Clostridium difficile* infections: The importance of the intestinal microbiota. *World J Gastroenterol* **20**, 7416–7423 (2014).
- Kelly, C. P.: Can we identify patients at high risk of recurrent *Clostridium difficile* infection? *Clin Microbiol Infect* **18**, 21–27 (2012).
- European Centre for Disease Prevention and Control. *Clostridium difficile* Infections Annual Epidemiological Report for 2016. ECDC, Stockholm, 2018. Available at https://ecdc.europa.eu/sites/portal/files/documents/AER_for_2016-C-difficile_0.pdf
- Naaber, P., Smidt, I., Stsepetova, J., Brilene, T., Annuk, H., Mikelsaar, M.: Inhibition of *Clostridium difficile* strains by in intestinal *Lactobacillus* species. *J Med Microbiol* **53**, 551–554 (2004).
- Deshpande, A., Pasupuleti, V., Thota, P., Pant, C., Rolston, D. D., Sferra, T. J., Hernandez, A. V., Donskey, C. J.: Community-associated *Clostridium difficile* infection and antibiotics: A meta-analysis. *J Antimicrob Chemother* **68**, 1951–1961 (2013).
- Jenberg, C., Löfmark, S., Edlund, C., Jansson, J. K.: Long-term ecological impacts of antibiotic administration on the human intestinal microbiota. *ISME J* **1**, 56–66 (2007).
- Furuya-Kanamori, L., Stone, J. C., Clark, J., McKenzie, S. J., Yakob, L., Paterson, D. L., Riley, T. V., Doi, S. A., Clements, A. C.: Comorbidities, exposure to medications, and the risk of community-acquired *Clostridium difficile* infection: A systematic review and meta-analysis. *Infect Control Hosp Epidemiol* **36**, 132–41 (2015).
- McDonald, L. C., Gerding, D. N., Johnson, S., Bakken, J. S., Carroll, K. C., Coffin, S. E., Dubberke, E. R., Garey, K. W., Gould, C. V., Kelly, C., Loo, V., Sammons, J. S., Sandora, T. J., Wilcox, M. H.: Clinical practice guidelines for *Clostridium difficile* infections in adults and children: 2017 update by the Infectious Diseases Society of America (IDSA) and Society for Healthcare Epidemiology of America (SHEA). *Clin Infect Dis* **66**, e1–e48 (2018).
- Lucado, J., Gould, C., Elixhauser, A.: *Clostridium difficile* Infections (CDI) in Hospital Stays, 2009: Statistical Brief #124 Healthcare Cost and Utilization Project (HCUP) Statistical Briefs. Rockville, MD: Agency for Health Care Policy and Research (US), 2012. Available at <https://www.ncbi.nlm.nih.gov/books/NBK92613/>
- Centers for Disease Control and Prevention (CDC): Severe *Clostridium difficile*-associated disease in populations previously at low risk: Four states, 2005. *Morb Mortal Wkly Rep* **54**, 1201–1205 (2005).
- Lessa, C. F., Mu, Y., Bamberg, W. M., Beldavs, Z. G., Dumyati, G. K., Dunn, J. R., Farley, M. M., Holzbauer, S. M., Meek, I. J., Phipps, C. E., Wilson, E. L., Winston, G. L., Cohen, A. J., Limbago, M. B., Fridkin, K. S., Gerding, N. D., Clifford, L., McDonald, C. L.: Burden of *Clostridium difficile* infection in the United States. *N Engl J Med* **372**, 825–834 (2015).
- Pituch, H., Obuch-Woszczyński, P., Lachowicz, D., Kuthan, R., Dzierżanowska-Fangrat, K., Mikucka, A., Jermakow, K., Pituch-Zdanowska, A., Davies, K.: Prevalence of *Clostridium difficile* infection in hospitalized patients with diarrhoea:

- Results of a Polish multicenter, prospective, biannual point-prevalence study. *Adv Med Sci* **63**, 290–295 (2018).
17. Kołpa, M., Wałaszek, M., Różańska, A., Wolak, Z., Wójkowska-Mach, J.: Hospital-wide surveillance of health-care-associated infections as a source of information about specific hospital needs. A 5-year observation in a multiprofile provincial hospital in the south of Poland. *Int J Environ Res Public Health* **15**, E1956 (2018).
 18. Ziółkowski, G., Pawłowska, I., Krawczyk, L., Wójkowska-Mach, J.: Antibiotic consumption versus the prevalence of multidrug-resistant *Acinetobacter baumannii* and *Clostridium difficile* infections at an ICU from 2014–2015. *J Infect Public Health* **11**, 626–630 (2018).
 19. Pituch, H., Obuch-Woszczatyński, P., Lachowicz, D., Wultańska, D., Karpiński, P., Młynarczyk, G., van Dorp, S. M., Kuijper, E. J.: Polish *Clostridium difficile* study group: hospital-based *Clostridium difficile* infection surveillance reveals high proportions of PCR ribotypes 027 and 176 in different areas of Poland, 2011 to 2013. *Euro Surveill* **20**, 38 (2015).
 20. Pituch, H., Bakker, D., Kuijper, E., Obuch-Woszczatyński, P., Wultańska, D., Nurzyńska, G., Bielec, A., Bar-Andziak, E., Łuczak, M.: First isolation of *Clostridium difficile* PCR-ribotype 027/toxinotype III in Poland. *Pol J Microbiol* **57**, 267–268 (2008).
 21. Pituch, H., Obuch-Woszczatyński, P., Lachowicz, D., Wultańska, D., Karpiński, P., Młynarczyk, G., van Dorp, S. M., Kuijper, E. J.: Hospital-based *Clostridium difficile* infection surveillance reveals high proportions of PCR ribotypes 027 and 176 in different areas of Poland, 2011 to 2013. *Euro Surveill* **20**, 38 (2015).
 22. Aptekorz, M., Szczegielniak, A., Wiechuła, B., Harmanus, C., Kuijper, E., Martirosian, G.: Occurrence of *Clostridium difficile* ribotype 027 in hospitals of Silesia, Poland. *Anaerobe* **45**, 106–113 (2017).
 23. Czepiel, J., Kędzierska, J., Biesiada, G., Birczyńska, M., Perucki, W., Nowak, P., Garlicki, A.: Epidemiology of *Clostridium difficile* infection: Results of a hospital-based study in Krakow, Poland. *Epidemiol Infect* **143**, 3235–3243 (2015).
 24. Barlam, T. F., Cosgrove, S. E., Abbo, L. M., MacDougall, C., Schuetz, A. N., Septimus, E. J., Srinivasan, A., Dellit, T. H., Falck-Ytter, Y. T., Fishman, N. O., Hamilton, C. W., Jenkins, T. C., Lipsett, P. A., Malani, P. N., May, L. S., Moran, G. J., Neuhauser, M. M., Newland, J. G., Ohl, C. A., Samore, M. H., Seo, S. K., Trivedi, K. K.: Implementing an antibiotic stewardship program: Guidelines by the Infectious Diseases Society of America and the Society for Healthcare Epidemiology of America. *Clin Infect Dis* **62**, 51–77 (2016).
 25. Wójkowska-Mach, J., Godman, B., Glassman, A., Kurdi, A., Pilc, A., Rozanska, A., Skoczyński, S., Wałaszek, M., Bochenek, T.: Antibiotic consumption and antimicrobial resistance in Poland; findings and implications. *Antimicrob Resist Infect Control* **7**, 136 (2018).
 26. Hanberger, H., Erlandsson, M., Burman, L. G., Cars, O., Gill, H., Lindgren, S., Nilsson, L. E., Olsson-Liljequist, B., Walther, S.: High antibiotic susceptibility among bacterial pathogens in Swedish ICUs. Report from a nationwide surveillance program using TA90 as a novel index of susceptibility. *Scand J Infect Dis* **36**, 24–30 (2004).
 27. Meyer, E., Gastmeier, P., Deja, M., Schwab, F.: Antibiotic consumption and resistance: data from Europe and Germany. *Int J Med Microbiol* **303**, 388–395 (2013).
 28. Trejnowska, E., Deptuła, A., Tarczyńska-Słomian, M., Knapik, P., Jankowski, M., Misiewska-Kaczur, A., Tamowicz, B., Śmiechowicz, J., Antończyk, R., Armatowicz, P., Sułkowski, W., Durek, G.: Surveillance of antibiotic prescribing in intensive care units in Poland. *Can J Infect Dis Med Microbiol* **2018**, 5670238 (2018).
 29. European Centre for Disease Prevention and Control. Antimicrobial Consumption: Annual Epidemiological Report for 2017. ECDC, Stockholm, 2018. Available at https://www.ecdc.europa.eu/sites/default/files/documents/AER_for_2017-antimicrobial-consumption.pdf
 30. Lachowicz, D., Pituch, H., Obuch-Woszczatyński, P.: Antimicrobial susceptibility patterns of *Clostridium difficile* strains belonging to different polymerase chain reaction ribotypes isolated in Poland in 2012. *Anaerobe* **31**, 37–41 (2015).
 31. Barker, A. K., Ngam, C., Musuza, J. S., Vaughn, V. M., Safdar, N.: Reducing *Clostridium difficile* in the inpatient settings: a systematic review of the adherence to and effectiveness of *C. difficile* prevention bundles. *Infect Control Hosp Epidemiol* **38**, 639–650 (2017).
 32. Tasteyre, A., Karjalainen, T., Avesani, V., Delmée, M., Collignon, A., Bourlioux, P., Barc, M. C.: Phenotypic and genotypic diversity of the flagellin gene (*fliC*) among *Clostridium difficile* isolates from different serogroups. *J Clin Microbiol* **38**, 3179–3186 (2000).
 33. Hennequin, C., Porcheray, F., Waligora-Dupriet, A., Collignon, A., Barc, M., Bourlioux, P., Karjalainen, T.: GroEL (Hsp60) of *Clostridium difficile* is involved in cell adherence. *Microbiology* **147**, 87–96 (2001).
 34. Waligora, A. J., Hennequin, C., Mullany, P., Bourlioux, P., Collignon, A., Karjalainen, T.: Characterization of a cell surface protein of *Clostridium difficile* with adhesive properties. *Infect Immun* **69**, 2144–2153 (2001).
 35. Hennequin, C., Janoir, C., Barc, M. C., Collignon, A., Karjalainen, T.: Identification and characterization of a fibronectin-binding protein from *Clostridium difficile*. *Microbiology* **149**, 2779–2787 (2003).
 36. Kirby, J. M., Ahern, H., Roberts, A. K., Kumar, V., Freeman, Z., Acharya, K. R., Shone, C. C.: Cwp84, a surface-associated cysteine protease, plays a role in the maturation of the surface layer of *Clostridium difficile*. *J Biol Chem* **284**, 34666–34673 (2009).
 37. Semenyuk, E. G., Laning, M. L., Foley, J.: Spore formation and toxin production in *Clostridium difficile* biofilms. *PLoS One* **9**, 87757 (2014).
 38. Burdon, D. W., George, R. H., Mogg, G. A., Arabi, Y., Thompson, H., Johnson, M., Alexander-Williams, J., Keighley, M. R.: Faecal toxin and severity of antibiotic-associated pseudomembranous colitis. *J Clin Pathol* **34**, 548–551 (1981).
 39. Kuehne, S. A., Cartman, S. T., Heap, J. T., Kelly, M. L., Cockayne, A., Minton, N. P.: The role of toxin A and toxin B in *Clostridium difficile* infection. *Nature* **467**, 711–712 (2010).
 40. Pituch, H., Kreft, D., Obuch-Woszczatyński, P., Wultańska, D., Meisel-Mikołajczyk, F., Łuczak, M., van Belkum, A.: Clonal spread of a *Clostridium difficile* strain with a complete set of a



- toxin A, toxin B, and a binary toxin genes among Polish patients with *Clostridium difficile* associated diarrhea. J Clin Microbiol **43**, 772–775 (2005).
41. Freeman, J., Fawley, W. N., Baines, S., Wilcox, M.: Measurement of toxin production by *Clostridium difficile*. Lancet **367**, 982–983 (2006).
 42. Castagliuolo, I., Kelly, C. P., Qiu, B. S., Nikulasson, S. T., LaMont, J. T., Pothoulakis, C.: IL-11 inhibits *Clostridium difficile* toxin A enterotoxicity in rat ileum. Am J Physiol **273**, 333–341 (1997).
 43. Pothoulakis, C., LaMont, J. T.: *Clostridium difficile* colitis and diarrhea. Gastroenterol Clin North Am **22**, 623–637 (1993).
 44. Khanna, S., Baddour, L. M., Huskins, W. C., Kammer, P. P., Faubion, W. A., Zinsmeister, A. R., Harmsen, W. S., Pardi, D. S.: The epidemiology of *Clostridium difficile* infection in children: A population-based study. Clin Infect Dis **56**, 1401–1406 (2013).
 45. Surawicz, C. M., Brandt, L. J., Binion, D. G., Ananthakrishnan, A. N., Curry, S. R., Gilligan, P. H., McFarland, L. V., Mellow, M., Zuckerbraun, B. S.: Guidelines for diagnosis, treatment, and prevention of *Clostridium difficile* infections. Am J Gastroenterol **108**, 478–98 (2013).
 46. Zar, F. A., Bakkanagari, S. R., Moorthi, K. M., Davis, M. B.: A comparison of vancomycin and metronidazole for treatment of *Clostridium difficile*-associated diarrhea, stratified by disease severity. Clin Infect Dis **45**, 302–307 (2007).
 47. Di, X., Bai, N., Zhang, X., Liu, B., Ni, W., Wang, J., Wang, K., Liang, B., Liu, Y., Wang, R.: A meta-analysis of metronidazole and vancomycin for the treatment *Clostridium difficile* infection, stratified by disease severity. Braz J Infect Dis **19**, 339–349 (2015).
 48. Gerding, D. N., Hecht, D. W., Louie, T., Nord, C. E., Talbot, G. H., Cornely, O. A., Buitrago, M., Best, E., Sambol, S., Osmolski, J. R., Kracker, H., Locher, H. H., Charef, P., Wilcox, M.: Susceptibility of *Clostridium difficile* isolates from a phase 2 clinical trial of cadazolid and vancomycin in *C. difficile* infection. J Antimicrob Chemother **71**, 213–219 (2015).
 49. Freeman, J., Vernon, J., Vickers, R., Wilcox, M. H.: Susceptibility of *Clostridium difficile* isolates of varying antimicrobial resistance phenotypes to SMT19969 and 11 comparators. Antimicrob Agents Chemother **60**, 689–692 (2015).
 50. Kokai-Kun, J. F., Roberts, T., Coughlin, O., Le, C., Whalen, H., Stevenson, R., Wachter, V. J., Sliman, J.: Use of ribaxamase (SYN-004), a β -lactamase, to prevent *Clostridium difficile* infection in β -lactam-treated patients: A double-blind, phase 2b, randomised placebo-controlled trial. Lancet Infect Dis **19**, 487–496 (2019).
 51. Taylor, N. S., Thorne, G. M., Bartlett, J. G.: Comparison of two toxins produced by *Clostridium difficile*. Infect Immun **34**, 1036–1043 (1981).
 52. Dabard, J., Dubos, F., Martinet, L., Ducluzeau, R.: Experimental reproduction of neonatal diarrhea in young gnotobiotic hares simultaneously associated with *Clostridium difficile* and other *Clostridium* strains. Infect Immun **24**, 7–11 (1979).
 53. Triadafilopoulos, G., Pothoulakis, C., O'Brien, M. J., LaMont, J. T.: Differential effects of *Clostridium difficile* toxins A and B on rabbit ileum. Gastroenterology **93**, 273–279 (1987).
 54. van Nood, E., Vriese, A., Nieuwdorp, M., Fuentes, S., Zoetendal, E. G., de Vos, W. M., Visser, C. E., Kuijper, E. J., Bartelsman, J. F., Tijssen, J. G., Speelman, P., Dijkgraaf, M. G., Keller, J. J.: Duodenal infusion of donor feces for recurrent *Clostridium difficile*. N Engl J Med **368**, 407–415 (2013).
 55. Borody, T. J., Connelly, N., Mitchell, S. W.: Fecal microbiota transplantation in gastrointestinal diseases. Pol Arch Med Wewn **125**, 852–858 (2015).
 56. FAO: Report of a Joint FAO/WHO Working Group on Drafting Guidelines for the Evaluation of Probiotics in Food. FAO, London Ontario, Canada, 2002. Available at http://www.who.int/foodsafety/fs_management/en/probiotic_guidelines.pdf
 57. Johnson, S., Maziade, P. J., McFarland, L. V., Trick, W., Donskey, C., Currie, B., Low, D. E., Goldstein, E. J.: Is primary prevention of *Clostridium difficile* infection possible with specific probiotics? Int J Infect Dis **16**, 786–792 (2012).
 58. Castagliuolo, I., LaMont, J. T., Nikulasson, S. T., Pothoulakis, C.: *Saccharomyces boulardii* protease inhibits *Clostridium difficile* toxin A effects in the rat ileum. Infect Immun **64**, 5225–5232 (1996).
 59. Czepiel, J., Biesiada, G., Drózd, M., Gdula-Argasińska, J., Żurańska, J., Marchewka, J., Perucki, W., Wołkow, P., Garlicki, A.: The presence of IL-8 +781 T/C polymorphism is associated with the parameters of severe *Clostridium difficile* infection. Microb Pathog **114**, 281–285 (2018).
 60. Plummer, S., Weaver, M. A., Harris, J. C., Dee, P., Hunter, J.: *Clostridium difficile* pilot study: Effects of probiotic supplementation on the incidence of *C. difficile* diarrhea. Int Microbiol **7**, 59–62 (2004).
 61. Kotowska, M., Albrecht, P., Szajewska, H.: *Saccharomyces boulardii* in the prevention of antibiotic-associated diarrhoea in children: A randomized double-blind placebo-controlled trial. Aliment Pharmacol Ther **21**, 583–590 (2005).
 62. Boonma, P., Spinler, J. K., Venable, S. F., Versalovic, J., Tumwasorn, S.: *Lactobacillus rhamnosus* L34 and *Lactobacillus casei* L39 suppress *Clostridium difficile*-induced IL-8 production by colonic epithelial cells. BMC Microbiol **14**, 177 (2014).
 63. Kaur, S., Vaishnavi, C., Prasad, K. K., Ray, P., Kochhar, R.: Effect of *Lactobacillus acidophilus* & epidermal growth factor on experimentally induced *Clostridium difficile* infection. Indian J Med Res **133**, 434–441 (2011).
 64. Lee, J. S., Chung, M. J., Seo, J. G.: *In vitro* evaluation of antimicrobial activity of lactic acid bacteria against *Clostridium difficile*. Toxicol. Res. **29**, 99–106 (2013).
 65. Leclercq, R., Cantón, R., Brown, D. F., Giske, C. G., Heisig, P., MacGowan, A. P., Mouton, J. W., Nordmann, P., Rodloff, A. C., Rossolini, G. M., Soussy, C. J., Steinbakk, M., Winstanley, T. G., Kahlmeter, G.: EUCAST expert rules in antimicrobial susceptibility testing. Clin Microbiol Infect **19**, 141–160 (2013).
 66. Hell, M., Bernhofer, C., Stalzer, P., Kern, J. M., Claassen, E.: Probiotics in *Clostridium difficile* infection: Reviewing the need for a multistrain probiotic. Benef Microb **4**, 39–51 (2013).
 67. Hütt, P., Shchepetova, J., Löivukene, K., Kullisaar, T., Mikelsaar, M.: Antagonistic activity of probiotic lactobacilli and bifidobacteria against entero- and uropathogens. J Appl Microbiol **100**, 1324–1332 (2005).
 68. Cassone, M., Serra, P., Mondello, F., Girolamo, A., Scafetti, S., Pistella, E., Venditti, M.: Outbreak of *Saccharomyces cerevisiae*

- subtype *boulardii fungemia* in patients neighboring those treatment with a probiotic preparation of the organism. *J Clin Microbiol* **41**, 5340–5343 (2003).
69. Rea, M. C., Sit, C. S., Clayton, E., O'Connor, P. M., Whittal, R. M., Zheng, J., Vederas, J. C., Ross, R. P., Hill, C.: Thuricin CD, a posttranslationally modified bacteriocin with a narrow spectrum of activity against *Clostridium difficile*. *Proc Natl Acad Sci U S A* **107**, 9352–9357 (2010).
70. Pechine, S., Gleizes, A., Janoir, C., Gorges-Kergot, R., Barc, M. C., Delmée, M., Collignon, A.: Immunological properties of surface proteins of *Clostridium difficile*. *J Med Microbiol* **54**, 193–196 (2005).
71. Pechine, S., Janoir, C., Collignon, A.: Variability of *Clostridium difficile* surface proteins and specific serum antibody response in patients with *Clostridium difficile*-associated disease. *J Clin Microbiol* **43**, 5018–5025 (2005).
72. Pechine, S., Deneve, C., Le Monnier, A., Hoys, S., Janoir, C., Collignon, A.: Immunization of hamsters against *Clostridium difficile* infection using the Cwp84 protease as an antigen. *FEMS Immunol Med Microbiol* **63**, 73–81 (2011).
73. Wright, A., Drudy, D., Kyne, L., Brown, K., Fairweather, N. F.: Immunoreactive cell wall proteins of *Clostridium difficile* identified by human sera. *J Med Microbiol* **57**, 750–756 (2008).
74. Wilcox, M. H., Gerding, D. N., Poxton, I. R., Kelly, C., Nathan, R., Birch, T., Cornely, A. O., Rahav, G., Bouza, E., Lee, C., Jenkin, G., Jensen, W., Kim, Y., Yoshida, J., Gabryelski, L., Pedley, A., Eves, K., Tipping, R., Guris, D., Kartsonis, N., Dorr, M.: Bezlotoxumab for prevention of recurrent *Clostridium difficile* infection. *N Engl J Med* **376**, 305–317 (2017).
75. Foglia, G., Shah, S., Luxemburger, C., Pietrobon, P. J.: *Clostridium difficile*: Development of a novel candidate vaccine. *Vaccine* **30**, 4307–4309 (2012).
76. Greenberg, R. N., Marburg, T. C., Foglia, G., Warny, M.: Phase I dose finding studies of an adjuvanted *Clostridium difficile* toxoid vaccine. *Vaccine* **30**, 2245–2249 (2012).
77. de Bruyn, G., Saleh, J., Workman, D., Pollak, R., Elinoff, V., Fraser, N. J., Lefebvre, G., Martens, M., Mills, R. E., Nathan, R., Trevino, M., van Cleeff, M., Foglia, G., Ozol-Godfrey, A., Patel, D. M., Pietrobon, P. J., Gesser, R.: Defining the optimal formulation and schedule of a candidate toxoid vaccine against *Clostridium difficile* infection: A randomized Phase 2 clinical trial. *Vaccine* **34**, 2170–2178 (2016).
78. Sekulovic, O., Meessen-Pinard, M., Fortier, L. C.: Prophage-stimulated toxin production in *Clostridium difficile* NAP1/027 lysogens. *J Bacteriol* **193**, 2726–2734 (2011).
79. Rea, M. C., Alemayehu, D., Rosss, R. P., Hill, C.: Gut solutions to a gut problem: Bacteriocins, probiotics and bacteriophage for control of *Clostridium difficile* infection. *J Med Microbiol* **62**, 1369–1378 (2013).
80. Nale, J. Y., Spencer, J., Hargreaves, K. R., Buckley, A. M., Trzepiński, P., Douce, R. G., Clokie, M. R. J.: Bacteriophage combinations significantly reduce *Clostridium difficile* growth *in vitro* and proliferation *in vivo*. *Antimicrob Agents Chemother* **60**, 968–981 (2016).
81. Nale, J. Y., Redgwell, T. A., Millard, A., Clokie, M. R. J.: Efficacy of optimised bacteriophage cocktail to clear *Clostridium difficile* in a batch fermentation model. *Antibiotics (Basel)* **7**, 13 (2018).
82. Tschudin-Sutter, S., Kuijper, E. J., Durovic, A., Vehreschild, M. J. G. T., Barbut, F., Eckert, C., Fitzpatrick, F., Hell, M., Norèn, T., O'Driscoll, J., Coia, J., Gastmeier, P., von Müller, L., Wilcox, M. H., Widmer, A. F.: Guidance document for prevention of *Clostridium difficile* infection in acute healthcare settings. *Clin Microbiol Infect* **24**, 1051–1054 (2018).