Epidemiology of community-acquired and recurrent *Clostridioides difficile* infection

Yichun Fu^(D), Yuying Luo and Ari M Grinspan^(D)

Abstract: Clostridioides difficile infection is a leading cause of healthcare-associated infections with significant morbidity and mortality. For the past decade, the bulk of infection prevention and epidemiologic surveillance efforts have been directed toward mitigating hospital-acquired *C. difficile*. However, the incidence of community-associated infection is on the rise. Patients with community-associated *C. difficile* tend to be younger and have lower mortality rate. Rates of recurrent *C. difficile* infection overall have decreased in the United States, but future research and public health endeavors are needed to standardize and improve disease detection, stratify risk factors in large-scale population studies, and to identify regional and local variations in strain types, reservoirs and transmission routes to help characterize and combat the changing epidemiology of *C. difficile*.

Keywords: Clostridioides difficile infection, community-acquired, epidemiology, fecal microbiota transplant, recurrent

Received: 31 January 2021; revised manuscript accepted: 19 April 2021.

Introduction

Clostridioides difficile infection (CDI) is a leading cause of healthcare-associated infections with significant morbidity and mortality. CDI is associated with 15,000–30,000 deaths annually in the United States and acute inpatient costs exceeding \$4.8 billion.¹ *C. difficile* is a spore-forming, anaerobic, gram-positive rod which causes disease through release of enterotoxin A and cytotoxin B. Certain strains, most notably NAP1/027, produce a markedly increased level of these toxins, which may contribute to its hypervirulence and association with severe disease manifestations.²

In recent years, tremendous efforts have been directed at curbing healthcare-associated CDI worldwide. The Centers for Disease Control (CDC) in the United States launched its *C. difficile* surveillance program in 2009 and highlighted six key components of prevention in its 2012 *Vital Signs* report.³ Globally, 28 countries/territories in four World Health Organization regions have also established guidelines for infection control, detection and surveillance.⁴ As a result, there is promising evidence that CDI rates have declined in Europe since 2010 and the United States since

2015.^{1,5} However, CDI is increasingly detected in individuals with no prior hospitalizations or healthcare exposures, suggesting a rise in community-acquired CDI.⁶ In addition, as many as 35% of patients will experience recurrent CDI (rCDI), with an increased risk of recurrence following each subsequent episode. This significantly impacts not only individual patients but also the healthcare system, as rCDI is associated with increased medical costs compared with primary episodes. Our objective is to review the burden and characteristics of community-acquired and rCDI, and to delineate the challenges posed by the changing epidemiology.

Community-acquired CDI

Rising incidence over the past decade

According to recent guidelines from the Society for Healthcare Epidemiology of America released in 2018, CDI cases are categorized as community associated (CA-CDI) if patients experienced diarrhea onset in the community or within 48h after hospitalization and had not been discharged from a healthcare facility in the prior 12 weeks.¹ Ther Adv Gastroenterol

Special Collection

2021, Vol. 14: 1–11 DOI: 10.1177/ 17562848211016248

© The Author(s), 2021. Article reuse guidelines: sagepub.com/journalspermissions

Correspondence to: Ari M Grinspan

Division of Gastroenterology, Icahn School of Medicine at Mount Sinai, 1 Gustave L Levy Place, Box 1190, New York, NY 10029-6574, USA ari.grinspan@mountsinai. org

Yichun Fu

Department of Medicine, Icahn School of Medicine at Mount Sinai, New York, NY, USA

Yuying Luo

Division of Gastroenterology, Icahn School of Medicine at Mount Sinai, New York, NY, USA



Creative Commons Non Commercial CC BY-NC: This article is distributed under the terms of the Creative Commons Attribution-NonCommercial 4.0 License (https://creativecommons.org/licenses/by-nc/4.0/) which permits non-commercial use, reproduction and distribution of the work without further permission provided the original work is attributed as specified on the SAGE and Open Access pages (https://us.sagepub.com/en-us/nam/open-access-at-sage).



Figure 1. Estimated national burden of total *Clostridium difficile* infection (CDI), community-associated CDI, and recurrent CDI, 2011–2017, adapted from Guh *et al.*⁶

National estimates reflect actual estimate of disease burden based on rates of nucleic acid amplification test in that year, adjusted for age, sex and race of the United States population. Estimated frequency of recurrent disease is calculated with a logistic-regression model using cases with complete data, adjusting for age, sex, race, and diagnostic method.

The population-based CA-CDI incidence was reported to be 20-40 per 100,000 people between 2005 and 2006, before more sensitive nucleic acid amplification tests for C. difficile diagnosis became available.7 Overall, the incidence of CA-CDI has risen over the past decade. In the most recent survey from the CDC Prevention Emerging Infections Program, the estimated incidence of CA-CDI in the United States increased from 52.88 per 100,000 people in 2012 to 65.93 per 100,000 people in 2017.8 Interestingly, while the absolute number of CA-CDI cases has increased from an estimated 170,000 in 2011 to 226,400 cases in 2017, there has been a slow downtrend in the total number total CDI cases nationally, from 476,400 in 2011 to 462,100 in 2017 (Figure 1).6

The uptick of CA-CDI is consistent with previous studies in different regions in North America (Table 1).^{7,9} In a retrospective multicenter cohort study from 43 hospitals in Georgia, Florida, North Carolina, South Carolina, Virginia, and West Virginia, CA-CDI increased from 48.5% to 61.0% of all CDI cases.^{10,11} CA-CDI has also led to increasing hospitalizations in the United States, from 17.1 per 100,000 persons in 2011 to 21.7 per 100,000 persons in 2017. In contrast, healthcare-associated CDI (HA-CDI) hospitalizations have been declining (60.7 per 100,000 persons to 47.9 per 100,000 persons in 2017).⁶ In Canada, CA-CDI has also increased in Quebec (from 0.51 to 0.69 per 100,000 population between 2008 and 2015) and Ontario (from 6.09 to 9.56 cases per 100,000 person-years between 2005 and 2015) during the overlapping time period while total CDI cases in the nation decreased.^{12–14}

Global CA-CDI trends vary considerably when compared with the United States and Canada. During 2006 and 2017, the reported proportion of CA-CDI varied between 10.9% and 43.0% of all CDI cases in Japan, Italy, and Australia.^{5,12,15–17} The global discrepancy likely lies in surveillance methods. Cases of CA-CDI tend to be lower using hospital-based surveillance while reaching nearly 50% with population-based surveillance, given that nearly 75% of patients with CA-CDI are treated as outpatients.^{7,17} The worldwide burden of CDI in the community warrants further investigation.

Risk factors and mortality for CA-CDI

Patients diagnosed with CA-CDI tend to be younger with a significant proportion of patients (36%) reporting no antibiotic exposure during

| Nation/region | Total CDI | CA-CDI | Reference |
|---------------|---|---|---|
| United States | decreased from 476,400 cases in 2011 to 462,100 cases in 2017 | increased from 170,000 cases (52.88 per 100,000 people) in 2011–2012 to 462,100 cases (65.93 per 100,000 people) in 2017 | Guh <i>et al.</i> ,6 Lessa <i>et al</i> .9 |
| Canada | decreased from 5.9 per 1000 patient/year in 2009 to 4.3 in 2015 | increased in Quebec from 2008 to 2015 (0.51–0.69 per 100,000 population) and increased in Ontario from 2005 to 2015 (6.09–9.56 cases per 100 000 person-years) | Katz <i>et al.</i> , ¹² Pereira <i>et al.</i> , ¹³ Zanichelli <i>et al</i> . ¹⁴ |

Table 1. Comparison of total C. difficile infection (CDI) versus community-associated CDI (CA-CDI) in United States and Canada.

Table 2. A comparison of risk factors and other characteristics in healthcare-associated *C. difficile* infection (CDI) versus communityacquired CDI versus recurrent CDI.

| | HA-CDI | CA-CDI | Recurrent CDI | Reference |
|--------------------------|---|---|---|---|
| Age | median age 72 | median age 50–51 | median age 56-75.3 | Lessa <i>et al.</i> , ⁹ Khanna <i>et al.</i> , ¹¹ Chitnis <i>et al.</i> , ¹⁷ Ma <i>et al.</i> , ²⁰ Richardson <i>et al.</i> ²¹ |
| Risk factors | prior antibiotic use, proton pump inhibitor, inflammatory bowel disease | prior antibiotic use, cardiac disease, chronic kidney disease, inflammatory bowel disease | older age, female gender, chronic kidney disease, inflammatory bowel disease, immunosuppression, prior use of corticosteroids | Khanna <i>et al.</i> , ¹¹ Chitnis <i>et al.</i> , ¹⁷ Brown <i>et al.</i> , ²² Fawley <i>et al.</i> , ²³ Ma <i>et al.</i> , ²⁰ Richardson <i>et al.</i> , ²¹ van Beurden <i>et al.</i> ²⁴ |
| 30-day mortality rate | 10.60% | 3–17% | 7.8-9.3% | Guh <i>et al.</i> , ⁶ Reveles <i>et al.</i> ²⁵ |
| Strain | 078, 106 | ribotype 002, 020, 014, 015, 027, 078, 106 | ribotype 027 | Guh <i>et al.</i> ⁶ , Lessa <i>et al.</i> ⁹ , Knetsch <i>et al.</i> ²⁶ , Romano <i>et al.</i> ²⁷ , Richardson <i>et al.</i> ²¹ , van Beurden <i>et al.</i> ²⁴ |

the 12 weeks prior to diagnosis (Table 2).^{1,9} In a population study by Khanna et al.,11 patients with CA-CDI were younger (median age of 50 versus 72 years) compared with patients diagnosed with hospital-acquired CDI (HA-CDI). In a 2015 surveillance study of national CDI burden in the United States conducted by the CDC, there was a larger proportion of younger patients (age < 65 years) with CA-CDI than HA-CDI (56.2% versus 30.5%).9 Patients with CA-CDI tend to have lower estimated mortality rates compared with HA-CDI, ranging from 0.09% in 2011 to 0.07% in 2017.6,18 In a single-center retrospective study in France, CA-CDI was weakly associated (p=0.047) with more severe disease based on a composite score compared with hospitalassociated infections; however, there was no difference in mortality.¹⁹ More prospective studies are needed to elucidate if the outcomes of patients with CA-CDI differ significantly from their counterparts with HA-CDI.

Antibiotic use has long been associated with increased risks of CDI [odds ratio (OR) 3.55, 2.56-4.94].²² As mentioned earlier, a larger portion of patients had no prior exposure to antibiotics in CA-CDI compared with HA-CDI. In a meta-analysis in the United States and a retrospective study in Britain, the proportions of CA-CDI patients without antibiotic exposure are approximately 21%-38%, compared with 6%-20.3% in HA-CDI.^{11,17,23} In terms of the classes of antibiotics, Brown et al.22 found that clindamycin [OR=16.80, 95% confidence interval (CI) 7.48-37.76], fluoroquinolones (OR=5.50, CI 4.26-7.11) and carbapenems (OR=5.68, CI 2.12-15.23) are the most likely to be associated with CDI, while macrolides (2.65, CI 1.92-3.64) and penicillins (OR=2.71, CI 1.75-4.21) are still problematic but less so.22

Among the medications that are suspected culprits for increasing risk for CDI, proton pump inhibitors (PPIs) have drawn extensive scrutiny. However, the association of PPIs in CA-CDI specifically has not been well studied. In a large prospective study from 2009 to 2011 of 984 patients with CA-CDI in eight US states, 31% of patients without prior antibiotic exposure reported PPI use.¹⁷ In a case-control study of CA-CDI cases in a registry in United Kingdom between 1994 and 2004, exposure to PPI during the 90 days before index date was correlated with increased risk of CA-CDI with an OR of 3.5.²⁸ However, while these studies suggest an association with PPI, there is a need for further, large-scale studies to better understand the relationship between PPI and CA-CDI.

Multiple chronic medical conditions have been associated with increased risk of CA-CDI, including cardiac disease (OR 4.87), chronic kidney disease (OR 12.12) and inflammatory bowel disease (IBD) (OR 5.13).²⁹ Among these, interest has grown in particular on exploring the relationship between IBD and development of CDI.¹ A population-based study of an IBD registry in Manitoba between 2005 and 2014 found the incidence of CDI to be 512 per 100,000 person-years for ulcerative colitis, 377 for Crohn's Disease, and 99 for non-IBD patients.³⁰ Importantly, it is unclear if some of these patients may be colonized with C. difficile in the setting of an IBD flare instead of having an actual infection.³¹ In a retrospective study of 1006 patients with IBD diagnosed with CDI, 56% were diagnosed with CA-CDI. Interestingly, among this sub-group, only 16% had steroid use within 1 month prior to CDI diagnosis, and only 30% of patients were exposed to an antibiotic.32 A prospective study of IBD patients in clinical remission with no recent hospitalization or antibiotic use were found to have higher rates of asymptomatic carriage of C. difficile compared with healthy controls (8.2% versus 1.0%). One potential explanation for the higher incidence of CDI in the IBD population without higher rates of antibiotic exposure is via disruption of the normal microbiome leading to decreased microbiota diversity and richness. Prior studies demonstrated that patients with IBD have alterations in their microbiome as a result of chronic inflammation even before treatment compared with healthy controls, including increased levels of Enterobacteriaceae, Pasteurellacaea, Veillonellaceae and Fusobacteriaceae, and decreased abundance in Ervsipelotrichales, Bacteroidales and Clostridiales.³³

The altered microbiome produces less butvrate which maintains gut mucosal integrity and more sialic acid, which is essential for C. difficile catabolism, resulting in a state of dysbiosis which increases susceptibility to CDI.34-36 This dysbiotic state has been well described in CDI patients without IBD.37 Medications used to treat IBD may also play a role in increased CDI rates for patients with IBD.32 A meta-analysis on risk factors of CDI in patients with IBD showed that only treatment with biologics and antibiotic use within the previous 30 days of diagnosis confers increased risk of CDI.38 However, while more than half of the studies in this meta-analysis diagnosed CDI by immunoassay or polymerase chain reaction (PCR), a large proportion used ICD10 codes; therefore, there may be discrepancies in diagnostic accuracy in this analysis. Future studies are needed to differentiate and stratify risk factors for CA-CDI in patients with IBD.

Community reservoirs, strain types and colonization

As the majority of patients with CA-CDI have no prior hospital or antibiotic exposure, multiple studies have attempted to elucidate possible routes of transmission. It has been proposed that many C. difficile reservoirs exist in the community including humans,³⁹ animals (e.g. ribotype 078 in pigs and cattle),²⁶ food sources,⁴⁰ and the environment (e.g. ribotype 014, 015 and 078 in wastewater plants).27 A recent analysis of more than 224,000 CDI cases from a large national insurance enrollee database found exposure through family members as the greatest factor for the increased incidence of CA-CDI.⁴¹ A meta-analysis from North America on the relationship of toxigenic C. difficile colonization and subsequent infection found 14.8% of long-term care residents were asymptomatic carriers of toxigenic strains.42 Another meta-analysis explored the colonization rates of patients upon admission to hospitals and found the global rate to be 8.1%.42,43 Healthy infants are commonly colonized with both toxigenic and non-toxigenic C difficile strains and contact with children <2 years old may be a risk factor for CA-CDI.44 Interestingly, colonization of nontoxigenic strains may be protective against development of CDI by alteration of host microbiome, as seen in a study of patients with cystic fibrosis.⁴⁵ However, colonization of toxigenic strains is still associated with higher risks of infection.^{43,46}

In addition to community reservoirs, several studies have employed sequencing to determine if particular C. difficile strains are responsible for CA-CDI transmission. The most common strain types associated with CA-CDI in United States are ribotypes 027, 106, 014, 002 and 020.6 The incidence of ribotype 027, previously the most prevalent strain in both HA-CDI and CA-CDI in North America, has been declining since 2011, while ribotype 106 has become the predominant strain since 2014.6,9 A study in Europe found instead five distinct ribotype clusters with no strain dominating the continent; for example, ribotypes 356 and 018 were detected predominantly in Italy, while ribotype 027 was prevalent in Hungary, Italy, Germany, Romania and Poland.⁴⁷ Given the different clustering in various regions, it is hypothesized that specific strains circulate in the community locally via communityspecific routes. A small prospective study in Canada suggested that the strains associated with CA-CDI are indistinguishable from hospitalacquired CDI.48 Overall, it remains unclear at this time if specific strain types have contributed more to the rising incidence of CA-CDI. The identification of regional clustering will potentially allow scientists to establish local routes of infection and resistance patterns, which would potentially impact infection control efforts and treatment selection.

Real-world CDI transmission rates outside of healthcare facilities remains an understudied epidemiologic question. In a study using a Markov model built with data extracted from national databases, the transmission rate from a person with CDI to an uncolonized individual in the community was estimated to be 0.1% compared with 0.05% from an asymptomatic colonized person to an uncolonized one. An uncolonized person was estimated to have a probability of 0.12% per day of acquiring CDI in the community, compared with 2.3% in the hospital and 0.37% in a long-term care facility.49 Despite these findings, the proportion of CA-CDI is rising, and the growing pool of colonized individuals is likely contributing. Future epidemiologic studies will be crucial to understand the mechanisms of community transmission to mitigate its spread.

Recurrent C. difficile infections

CDI recurrence is defined as symptom onset and positive assay result following interim resolution

of symptoms with standard of care treatment of a primary episode in the previous 2-8 weeks.¹ However, given that several testing methods are used for clinical detection, reported incidences may not accurately reflect true infections (Supplemental Table 1). PCR tests do not differentiate C. difficile colonization from C. difficile infection.50 A positive test for CDI must be assessed in clinical context. Also, different testing methods may have varying sensitivities based on ribotype. Guh et al.51 found that toxin enzyme immunoassay is more sensitive for detecting certain strains, while nucleotide acid amplification testing is not affected by strain types.⁵¹ This has implications in terms of our ability to accurately assess disease burden and epidemiology, and impacts development of tailored approaches to combating C. difficile.

Ma et al.20 reported that the annual incidence rate of CDI recurrence increased from 0.0107 to 0.0309 per 1000 person-years between 2001 and 2012.²⁰ More recently, recurrent cases started to decrease during the past decade. In the United States, the estimated number of first recurrent episodes decreased from 93,400 to 69,800 between 2011 and 2017, with incidence dropping from 19.7 per 100,000 to 12.0 per 100,000 over the same time period.⁶ The decrease in incidence rate is likely multifactorial, including vigilant antibiotic stewardship, improved testing algorithms and treatment recommendations. Nonetheless, it is estimated that 10-30% of patients in the United States with CDI will still experience at least one recurrent episode, comparable to findings from other regions.^{5,15,18,52} In the 2020 surveillance by Guh et al.,6 the number of CA-CDI recurrences has been consistently lower than recurrences in HA-CDI, by approximately 21% in 2016 and 2017. This difference is probably due to the younger age of patients and less exposure to healthcare facilities in the CA-CDI population. It is important to note that not all epidemiologic studies routinely distinguish HA-CDI versus CA-CDI when calculating rCDI. Further investigation is needed to understand the burden of recurrent infections in each environment.

Relapse and reinfection

Recurrent CDI cases can be subdivided into relapse and reinfection. A relapse originates from the same strain as the initial infection, while reinfection is caused by a new strain (Table 2). Accurate classification between the two allows for rigorous evaluation of risk factors and treatment effectiveness. Several studies have sought to characterize the epidemiology of relapse versus reinfection for rCDI. Using whole-genome sequencing, Sim et al.53 determined that 15.8% of the rCDI at a single center were due to reinfection, and one in three such cases would have been misclassified as relapse.53 Given that wholegenome sequencing may not be widely available or practical, there has been interest in determining an appropriate timeline of recurrence that would help distinguish between relapse and reinfection. In a single-center cohort study conducted in Switzerland, Durovic et al.54 found that a time cutoff of 20 weeks resulted in acceptable discrimination between relapse and reinfection [receiver operating characteristic (ROC) area under the curve (AUC) 0.741]. Another study conducted in West Australia concluded that a cutoff of 12 weeks vielded a comparable distinction threshold (ROC AUC 0.61).¹⁸ Current clinical guidelines define rCDI as within 8 weeks of successful treatment, regardless of relapse or reinfection.¹ Sequencing C. difficile strains on a patient level to distinguish relapse or reinfection can potentially aid clinical decision-making but faces numerous barriers for widespread adoption.

Early versus delayed recurrence

There are limited data comparing the rates of early *versus* delayed CDI recurrence in the general population. A single-center study of 120 patients in Korea found that 19.2% of patients developed recurrence within 8weeks, 19.5% experienced delayed recurrence 8 weeks after initial treatment and 31% experienced a recurrence after 60 months. Risk factors for delayed recurrence include age \geq 70 years, whereas nasogastric tube insertion, PPI and H₂-blocker treatment has been associated with both early and delayed recurrence.⁵⁵

Risk factors for CDI recurrence

Several risk factors have been identified for rCDI including host factors such as demographics, comorbidities, previous episodes of CDI, as well as medications and environmental factors. In general, older age and female gender have been associated with increased risk for recurrent infections.^{20,21,54} A meta-analysis showed that age >65 years is an independent risk factor for rCDI (relative risk 1.63).⁵⁶ Multiple retrospective

studies and cohort studies in North America have shown that women are at increased risk for rCDI compared with men.^{20,53} Medical conditions such as chronic renal insufficiency (OR 1.59), immunosuppression (OR 9.64) and IBD (HR 1.63) increase the risk of recurrence as well.^{15,20,57,58} The potential mechanisms of how each of these conditions and others increase risk of rCDI is beyond the scope of this review. A prior history of CDI also increases risk of rCDI episodes, with studies showing that 20% of patients recur after a single episode, 40% after two episodes and 65% after three episodes.^{52,56,59}

Certain medications are associated with increased risk of rCDI. Conflicting data exist on the role of PPI in rCDI. A strong association between PPIs and rCDI was observed in a retrospective study of 45,341 patients with CDI in the United States (OR 1.14).²⁰ A similar correlation was found in a prospective study (OR 3.75).⁶⁰ However, a 6-year retrospective study in Italy did not find a significant correlation between PPI use and increased risk for recurrence (OR 0.94, CI 0.23–3.85).¹⁵ Corticosteroids (OR 1.15) and antibiotics other than those used for CDI treatment (OR 1.79) have also been associated with recurrent infections.²⁰

Apart from host factors, many studies have examined if specific strains are associated with increased recurrence. Ribotype 027 has been associated with recurrent infections in cohort studies in the United States and the Netherlands.^{21,54} Infection with other ribotypes, such as F014-020, were more commonly seen in patients with a single episode of CDI.⁶¹ Several other ribotypes are associated with recurrent diseases in other regions (such as ribotype 001 in Sweden and ribotype 014/020 in West Australia).^{18,62} The presence of multiple strains during the first CDI episode may also be a risk factor for recurrence.⁶¹

Mortality in CDI recurrence

There are conflicting reports regarding mortality associated with rCDI. In a study of inpatient hospitalizations between 2003 and 2009 at one academic tertiary care center, rCDI was associated with a 33% increased risk of mortality at 180 days compared with patients who do not experience a recurrence.⁶³ However, more recent studies have shown a lower mortality rate in rCDI compared with the initial infection. In a retrospective study of the US veteran population from 2003 to 2014, 30-, 60- and 90-day mortality rates were significantly lower in first CDI recurrence as well as in second recurrence compared with the primary episode (p < 0.0001).²⁵ Another large study from 128 US Veteran Affairs facilities between 2010 and 2014 confirmed these findings, observing a higher 30-day mortality rate for initial CDI (10.6%) *versus* first recurrence (8.3%, p < 0.001) and second recurrence (4.2%, p = 0.004).⁶⁴ Risk factors for increased mortality include advanced age, PPI use within 7 days before CDI treatment, and any non-CDI related antibiotic use prior to, concurrent with or after *C. difficile* episode.⁶⁴

Treatment for rCDI

Treatment regimens for rCDI vary depending on the number of recurrences and severity of the current episode. For nonfulminant recurrent episodes, vancomycin pulsed taper or fidaxomicin are the recommended antibiotic therapies. Crook *et al.*⁶⁵ found fidaxomicin to be superior to vancomycin in preventing recurrence in a meta-analysis of two large randomized trials [Relative Risk (RR) 0.54, 95% CI 0.42–0.71]. Fecal microbiota transplant (FMT) can be considered after two recurrences.¹

FMT has garnered strong interest for its potential in prevention of CDI recurrence. In a recent small randomized study comparing FMT, fidaxomicin and vancomycin for the treatment of rCDI, significantly more patients who underwent FMT remained symptom-free at 8 weeks post-treatment (p=0.009 for FMT versus fidaxomicin; p=0.001 for FMT versus vancomycin).⁶⁶ In another prospective study of 19 patients, the efficacy rate of FMT was 85%, and the post-treatment stool analysis showed significantly increased bacterial diversity compared with the pre-FMT specimens.⁶⁷ In a meta-analysis of randomized controlled trials (RCTs) for prevention of rCDI, FMT was found to be superior to oral vancomycin in the prevention of rCDI.68 Recent costeffectiveness studies have shown FMT to dominate all other strategies for the treatment and prevention of rCDI.⁶⁹ However, the reported range of efficacy rates of FMT have been wide, and the more recent RCTs have shown lower efficacy rates than the open-labeled studies.⁷⁰ The quality of data is limited by variations in methodologies, including lack of control arms, difference in antibiotic suppressive therapies, and selections

of CDI diagnostic assays, all of which likely contribute to the discrepancy.⁶⁸

Bezlotoxumab, a monoclonal antibody against *C. difficile* toxin B, has also been studied as an option to reduce risk of recurrence, with the MODIFY I and II trials demonstrating significant efficacy in preventing CDI recurrence over placebo (RR 0.62, 95% CI, 0.51–0.75).⁷¹ Scaling of these newer strategies to treat rCDI will likely impact the epidemiology and incidence in the next decade. Further studies will be helpful to guide positioning of various therapies in our armamentarium for treating rCDI, balancing efficacy, safety and cost-effectiveness.

COVID-19 and CDI

Lastly, there have been ongoing efforts to monitor the trend of CDI during the COVID-19 pandemic. The published data so far showed that there has not been an increase in the rate of CDI despite rising high-risk antibiotic exposures because of COVID-19.72,73 In some centers, there has been a reduction in HA-CDI cases.74,75 Multiple factors potentially contributed to the decrease. Among them, one important aspect has been improved infection prevention measures such as higher compliance in hand hygiene and isolation precautions.74,76 Only one small study specifically commented on CA-CDI versus HA-CDI, finding that six out of 38 CDI cases were community-onset, and two out of 38 were rCDI.76 In that study, COVID-19 patients with CDI had worse outcome than those without CDI, including longer hospital stay (35 days versus 19.4 days, p < 0.01) and lower rates of full recoverv without complications (50% versus 64.9%, p=0.01).⁷⁶ However, most of these cases were hospital-onset, and the trend in CA-CDI has not been clearly delineated. Further studies are needed to explore the interplay between COVID-19 and CDI, especially given the exposure to steroids and antibiotics in patients who received treatment for COVID-19.

Conclusion

Despite an overall decrease in CDI rates worldwide in the last decade, community-associated cases are increasing while the rates of recurrence of CDI are steady. CA-CDI accounts for almost 48% of annual cases in the United States and ranges from 10% to 43% of total cases across global regions. Continued efforts directed at improving disease detection and tracking are needed to help identify community reservoirs and understand patterns of transmission. While rCDI cases have decreased in the United States, rCDI remains a leading cause of significant healthcare morbidity and global economic burden. Importantly, the risk factors and disease characteristics of CA-CDI and rCDI differ regionally, calling for rigorous future investigations of various host factors, pathogen ribotypes, local reservoirs and transmission vectors specific to each region. Standardization of laboratory testing for the detection of CDI is critical to accurately characterize true infections versus colonization. With the availability of highly sensitive tests such as whole-genome sequencing, our understanding of CDI pathogenesis, transmission and colonization will continue to evolve over the next decade.

Conflict of interest statements

The authors declare that there is no conflict of interest.

Funding

The authors received no financial support for the research, authorship, and/or publication of this article.

ORCID iDs

Yichun Fu 🕩 https://orcid.org/0000-0001-5620-3200

Ari M Grinspan ២ https://orcid.org/0000-0002-0719-1538

Supplemental material

Supplemental material for this article is available online.

References

- 1. McDonald LC, Gerding DN, Johnson S, *et al.* Clinical practice guidelines for *Clostridium difficile* infection 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* 2018; 66: e1–e48.
- Gerding DN, Young VB and Donskey CJ. *Clostridioides difficile* (formerly *Clostridium difficile*) infection. In: Bennett JE, Dolin R and Blaser MJ

(eds) Mandell, Douglas, and Bennett's principles and practice of infectious diseases. Philadelphia, PA: Elsevier, 2020.

- Centers for Disease Control and Prevention (CDC). Vital signs: preventing *Clostridium difficile* infections. *MMWR Morb Mortal Wkly Rep* 2012; 61: 157–162.
- Balsells E, Filipescu T, Kyaw MH, et al. Infection prevention and control of *Clostridium* difficile: a global review of guidelines, strategies, and recommendations. J Glob Health 2016; 6: 020410.
- Levy AR, Szabo SM, Lozano-Ortega G, et al. Incidence and costs of *Clostridium difficile* infections in Canada. *Open Forum Infect Dis* 2015; 2: ofv076.
- Guh AY, Mu Y, Winston LG, et al. Trends in U.S. burden of *Clostridioides difficile* infection and outcomes. N Engl J Med 2020; 382: 1320–1330.
- Gerding DN and Lessa FC. The epidemiology of *Clostridium difficile* infection inside and outside health care institutions. *Infect Dis Clin North Am* 2015; 29: 37–50.
- 8. Centers for Disease Control and Prevention. 2018 Annual report for the emerging infections program for Clostridioides difficile infection. Atlanta, GA: CDC, 2020.
- Lessa FC, Mu Y, Bamberg WM, et al. Burden of Clostridium difficile infection in the United States. N Engl J Med 2015; 372: 825–834.
- Turner NA, Grambow SC, Woods CW, et al. Epidemiologic trends in *Clostridioides difficile* infections in a regional community hospital network. *JAMA Netw Open* 2019; 2: e1914149.
- Khanna S, Pardi DS, Aronson SL, et al. The epidemiology of community-acquired *Clostridium* difficile infection: a population-based study. Am J Gastroenterol 2012; 107: 89–95.
- Katz KC, Golding GR, Choi KB, et al. The evolving epidemiology of *Clostridium difficile* infection in Canadian hospitals during a postepidemic period (2009–2015). *CMAJ* 2018; 190: E758–E765.
- Pereira JA, McGeer A, Tomovici A, et al. Incidence and economic burden of *Clostridioides* difficile infection in Ontario: a retrospective population-based study. *CMAJ Open* 2020; 8: E16–E25.
- Zanichelli V, Garenc C, Villeneuve J, et al. Increased community-associated *Clostridioides* difficile infections in Quebec, Canada, 2008– 2015. Emerg Infect Dis 2020; 26: 1291–1294.

- Roncarati G, Dallolio L, Leoni E, et al. Surveillance of *Clostridium difficile* infections: results from a six-year retrospective study in nine hospitals of a North Italian local health authority. *Int J Environ Res Public Health* 2017; 14: 61.
- Collins DA, Sohn KM, Wu Y, et al. Clostridioides difficile infection in the Asia-Pacific region. Emerg Microbes Infect 2020; 9: 42–52.
- Chitnis AS, Holzbauer SM, Belflower RM, et al. Epidemiology of community-associated Clostridium difficile infection, 2009 through 2011. JAMA Intern Med 2013; 173: 1359–1367.
- Alfayyadh M, Collins DA, Tempone S, et al. Recurrence of *Clostridium difficile* infection in the Western Australian population. *Epidemiol Infect* 2019; 147: e153.
- Ogielska M, Lanotte P, Le Brun C, et al. Emergence of community-acquired Clostridium difficile infection: the experience of a French hospital and review of the literature. Int J Infect Dis 2015; 37: 36–41.
- 20. Ma GK, Brensinger CM, Wu Q, *et al.* Increasing incidence of multiply recurrent *Clostridium difficile* infection in the United States: a cohort study. *Ann Intern Med* 2017; 167: 152–158.
- 21. Richardson C, Kim P, Lee C, *et al.* Comparison of *Clostridium difficile* isolates from individuals with recurrent and single episode of infection. *Anaerobe* 2015; 33: 105–108.
- Brown KA, Khanafer N, Daneman N, et al. Meta-analysis of antibiotics and the risk of community-associated *Clostridium difficile* infection. *Antimicrob Agents Chemother* 2013; 57: 2326–2332.
- Fawley WN, Davies KA, Morris T, et al. Enhanced surveillance of *Clostridium difficile* infection occurring outside hospital, England, 2011 to 2013. *Euro Surveill* 2016; 21: 30295.
- 24. van Beurden YH, Nezami S, Mulder CJJ, et al. Host factors are more important in predicting recurrent *Clostridium difficile* infection than ribotype and use of antibiotics. *Clin Microbiol Infect* 2018; 24: 85.e81–85.e84.
- 25. Reveles KR, Lawson KA, Mortensen EM, *et al.* National epidemiology of initial and recurrent *Clostridium difficile* infection in the Veterans Health Administration from 2003 to 2014. *PLoS One* 2017; 12: e0189227.
- 26. Knetsch CW, Connor TR, Mutreja A, et al. Whole genome sequencing reveals potential spread of *Clostridium difficile* between humans and farm animals in the Netherlands, 2002 to 2011. *Euro Surveill* 2014; 19: 20954.

- 27. Romano V, Pasquale V, Krovacek K, *et al.* Toxigenic *Clostridium difficile* PCR ribotypes from wastewater treatment plants in Southern Switzerland. *Appl Environ Microbiol* 2012; 78: 6643–6646.
- Dial S, Delaney JA, Schneider V, et al. Proton pump inhibitor use and risk of communityacquired *Clostridium difficile*-associated disease defined by prescription for oral vancomycin therapy. *CMAJ* 2006; 175: 745–748.
- 29. Guh AY, Adkins SH, Li Q, *et al.* Risk factors for community-associated *Clostridium difficile* infection in adults: a case-control study. *Open Forum Infect Dis* 2017; 4: ofx171.
- Singh H, Nugent Z, Yu BN, et al. Higher incidence of *Clostridium difficile* infection among individuals with inflammatory bowel disease. *Gastroenterology* 2017; 153: 430–438.e432.
- Beniwal-Patel P, Stein DJ and Munoz-Price LS. The juncture between *Clostridioides difficile* infection and inflammatory bowel diseases. *Clin Infect Dis* 2019; 69: 366–372.
- Gillespie W, Marya N, Fahed J, et al. Clostridium difficile in inflammatory bowel disease: a retrospective study. Gastroenterol Res Pract 2017; 2017: 4803262.
- Gevers D, Kugathasan S, Denson LA, et al. The treatment-naive microbiome in new-onset Crohn's disease. Cell Host Microbe 2014; 15: 382–392.
- Machiels K, Joossens M, Sabino J, et al. A decrease of the butyrate-producing species *Roseburia hominis* and *Faecalibacterium prausnitzii* defines dysbiosis in patients with ulcerative colitis. *Gut* 2014; 63: 1275–1283.
- Ng KM, Ferreyra JA, Higginbottom SK, et al. Microbiota-liberated host sugars facilitate postantibiotic expansion of enteric pathogens. *Nature* 2013; 502: 96–99.
- Rodriguez C, Romero E, Garrido-Sanchez L, et al. Microbiota insights in *Clostridium difficile* infection and inflammatory bowel disease. *Gut Microbes* 2020; 12: 1725220.
- Smits WK, Lyras D, Lacy DB, et al. Clostridium difficile infection. Nat Rev Dis Primers 2016; 2: 16020.
- Balram B, Battat R, Al-Khoury A, et al. Risk factors associated with *Clostridium difficile* infection in inflammatory bowel disease: a systematic review and meta-analysis. J Crohns Colitis 2019; 13: 27–38.
- 39. Loo VG, Brassard P and Miller MA. Household transmission of *Clostridium difficile* to family

members and domestic pets. *Infect Control Hosp Epidemiol* 2016; 37: 1342–1348.

- Rodriguez-Palacios A, Ilic S and LeJeune JT. *Clostridium difficile* with Moxifloxacin/ Clindamycin resistance in vegetables in Ohio, USA, and prevalence meta-analysis. *J Pathog* 2014; 2014: 158601.
- 41. Miller AC, Segre AM, Pemmaraju SV, et al. Association of household exposure to primary *Clostridioides difficile* infection with secondary infection in family members. JAMA Netw Open 2020; 3: e208925.
- 42. Ziakas PD, Zacharioudakis IM, Zervou FN, *et al.* Asymptomatic carriers of toxigenic *C. difficile* in long-term care facilities: a meta-analysis of prevalence and risk factors. *PLoS One* 2015; 10: e0117195.
- Zacharioudakis IM, Zervou FN, Pliakos EE, et al. Colonization with toxinogenic C. difficile upon hospital admission, and risk of infection: a systematic review and meta-analysis. Am J Gastroenterol 2015; 110: 381–390; quiz 391.
- Martin JS, Monaghan TM and Wilcox MH. *Clostridium difficile* infection: epidemiology, diagnosis and understanding transmission. *Nat Rev Gastroenterol Hepatol* 2016; 13: 206–216.
- 45. Bauer MP, Farid A, Bakker M, et al. Patients with cystic fibrosis have a high carriage rate of non-toxigenic *Clostridium difficile*. *Clin Microbiol Infect* 2014; 20: O446–O449.
- Vincent C, Miller MA, Edens TJ, et al. Bloom and bust: intestinal microbiota dynamics in response to hospital exposures and *Clostridium difficile* colonization or infection. *Microbiome* 2016; 4: 12.
- Eyre DW, Davies KA, Davis G, et al. Two distinct patterns of *Clostridium difficile* diversity across Europe indicating contrasting routes of spread. *Clin Infect Dis* 2018; 67: 1035–1044.
- Thornton CS, Rubin JE, Greninger AL, et al. Epidemiological and genomic characterization of community-acquired *Clostridium difficile* infections. *BMC Infect Dis* 2018; 18: 443.
- Durham DP, Olsen MA, Dubberke ER, et al. Quantifying transmission of *Clostridium difficile* within and outside healthcare settings. *Emerg Infect Dis* 2016; 22: 608–616.
- Polage CR, Gyorke CE, Kennedy MA, et al. Overdiagnosis of *Clostridium difficile* infection in the molecular test era. *JAMA Intern Med* 2015; 175: 1792–1801.
- 51. Guh AY, Hatfield KM, Winston LG, et al. Toxin enzyme immunoassays detect *Clostridioides*

difficile infection with greater severity and higher recurrence rates. *Clin Infect Dis* 2019; 69: 1667–1674.

- McFarland LV, Surawicz CM, Rubin M, et al. Recurrent Clostridium difficile disease: epidemiology and clinical characteristics. Infect Control Hosp Epidemiol 1999; 20: 43–50.
- 53. Sim JH, Truong C, Minot SS, et al. Determining the cause of recurrent *Clostridium difficile* infection using whole genome sequencing. *Diagn Microbiol Infect Dis* 2017; 87: 11–16.
- Durovic A, Widmer AF, Frei R, et al. Distinguishing Clostridium difficile recurrence from reinfection: independent validation of current recommendations. Infect Control Hosp Epidemiol 2017; 38: 891–896.
- 55. Doh YS, Kim YS, Jung HJ, *et al.* Long-term clinical outcome of *Clostridium difficile* infection in hospitalized patients: a single center study. *Intest Res* 2014; 12: 299–305.
- Deshpande A, Pasupuleti V, Thota P, et al. Risk factors for recurrent *Clostridium difficile* infection: a systematic review and meta-analysis. *Infect Control Hosp Epidemiol* 2015; 36: 452–460.
- 57. Razik R, Rumman A, Bahreini Z, et al. Recurrence of *Clostridium difficile* infection in patients with inflammatory bowel disease: the RECIDIVISM study. Am J Gastroenterol 2016; 111: 1141–1146.
- Scappaticci GB, Perissinotti AJ, Nagel JL, et al. Risk factors and impact of *Clostridium difficile* recurrence on haematology patients. *J Antimicrob Chemother* 2017; 72: 1488–1495.
- 59. Aslam S, Hamill RJ and Musher DM. Treatment of *Clostridium difficile*-associated disease: old therapies and new strategies. *Lancet Infect Dis* 2005; 5: 549–557.
- 60. Khanna S, Montassier E, Schmidt B, *et al.* Gut microbiome predictors of treatment response and recurrence in primary *Clostridium difficile* infection. *Aliment Pharmacol Ther* 2016; 44: 715–727.
- 61. Seekatz AM, Wolfrum E, DeWald CM, *et al.* Presence of multiple *Clostridium difficile* strains at primary infection is associated with development of recurrent disease. *Anaerobe* 2018; 53: 74–81.
- 62. Chen Y, Rashid MU, Huang H, et al. Molecular characteristics of *Clostridium difficile* strains from patients with a first recurrence more than 8 weeks after the primary infection. *J Microbiol Immunol Infect* 2017; 50: 532–536.
- 63. Olsen MA, Yan Y, Reske KA, et al. Recurrent *Clostridium difficile* infection is associated with

increased mortality. *Clin Microbiol Infect* 2015; 21: 164–170.

- 64. Appaneal HJ, Caffrey AR, Beganovic M, *et al.* Predictors of mortality among a national cohort of veterans with recurrent *Clostridium difficile* infection. *Open Forum Infect Dis* 2018; 5: ofy175.
- Crook DW, Walker AS, Kean Y, et al. Fidaxomicin versus vancomycin for *Clostridium* difficile infection: meta-analysis of pivotal randomized controlled trials. *Clin Infect Dis* 2012; 55(Suppl. 2): S93–S103.
- 66. Hvas CL, Dahl Jørgensen SM, Jørgensen SP, et al. Fecal microbiota transplantation is superior to fidaxomicin for treatment of recurrent *Clostridium difficile* infection. *Gastroenterology* 2019; 156: 1324–1332.e3.
- Barberio B, Facchin S, Mele E, et al. Faecal microbiota transplantation in *Clostridioides difficile* infection: real-life experience from an academic Italian hospital. *Therap Adv Gastroenterol* 2020; 13: 1756284820934315.
- 68. Madoff SE, Urquiaga M, Alonso CD, *et al.* Prevention of recurrent *Clostridioides difficile* infection: a systematic review of randomized controlled trials. *Anaerobe* 2020; 61: 102098.
- Luo Y, Lucas AL and Grinspan AM. Fecal transplants by colonoscopy and capsules are cost-effective strategies for treating recurrent *Clostridioides difficile* infection. *Dig Dis Sci* 2020; 65: 1125–1133.
- 70. Tariq R, Pardi DS, Bartlett MG, *et al.* Low cure rates in controlled trials of fecal microbiota transplantation for recurrent *Clostridium difficile*

infection: a systematic review and meta-analysis. *Clin Infect Dis* 2019; 68: 1351–1358.

- Wilcox MH, Gerding DN, Poxton IR, et al. Bezlotoxumab for prevention of recurrent *Clostridium difficile* infection. N Engl J Med 2017; 376: 305–317.
- 72. Luo Y, Grinspan LT, Fu Y, et al. Hospitalonset *Clostridioides difficile* infections during the COVID-19 pandemic. *Infect Control Hosp Epidemiol*. Epub ahead of print 23 September 2020. DOI: 10.1017/ice.2020.1223.
- Laszkowska M, Kim J, Faye AS, et al. Prevalence of *Clostridioides difficile* and other gastrointestinal pathogens in patients with COVID-19. *Dig Dis Sci*. Epub ahead of print 22 January 2021. DOI: 10.1007/s10620-020-06760-y.
- 74. Hazel K, Skally M, Glynn E, et al. The other 'C': hospital-acquired *Clostridioides difficile* infection during the coronavirus disease 2019 (COVID-19) pandemic. *Infect Control Hosp Epidemiol*. Epub ahead of print 13 January 2021. DOI: 10.1017/ ice.2021.3.
- Ponce-Alonso M, de la Fuente JS, Rincon-Carlavilla A, et al. Impact of the coronavirus disease 2019 (COVID-19) pandemic on nosocomial Clostridioides difficile infection -ERRATUM. Infect Control Hosp Epidemiol 2021; 42: 250.
- 76. Granata G, Bartoloni A, Codeluppi M, et al. The burden of *Clostridioides difficile* infection during the COVID-19 pandemic: a retrospective casecontrol study in Italian hospitals (CloVid). *J Clin Med* 2020; 9: 3855.

Visit SAGE journals online journals.sagepub.com/ home/tag

SAGE journals