# **Original Article**



# The changing epidemiology of *Clostridioides difficile* infection and the NAP1/027 strain in two Québec hospitals: a 17-year time-series study

Sandrine Couture MD<sup>1</sup>, Charles Frenette MD<sup>1,2</sup> <sup>(1)</sup>, Ian Schiller M.Sc.<sup>3</sup>, Rowin Alfaro B.Sc.<sup>3</sup>, Nandini Dendukuri PhD<sup>1,3</sup>,

Daniel Thirion Pharm D<sup>4,5</sup> (1), Yves Longtin MD<sup>1,6</sup> (1) and Vivian G. Loo MD, M.Sc.<sup>1,2</sup> (1)

<sup>1</sup>McGill University, Montréal, QC, Canada, <sup>2</sup>Division of Infectious Diseases and Medical Microbiology, McGill University Health Centre, Montréal, QC, Canada, <sup>3</sup>Research Institute of the McGill University Health Centre, Montréal, QC, Canada, <sup>4</sup>Université de Montréal, Montréal, QC, Canada, <sup>5</sup>Department of Pharmacy, McGill University Health Centre, Montréal, QC, Canada and <sup>6</sup>Jewish General Hospital, Montréal, QC, Canada

# Abstract

Objective: To describe the epidemiology of healthcare-associated *Clostridioides difficile* infection (HA-CDI) in two Québec hospitals in Canada following the 2003 epidemic and to evaluate the impact of antibiotic stewardship on the incidence of HA-CDI and the NAP1/027 strain.

Design: Time-series analysis.

Setting: Two Canadian tertiary care hospitals based in Montréal, Québec.

Patients: Patients with a positive assay for toxigenic *C. difficile* were identified through infection control surveillance. All cases of HA-CDI, defined as symptoms occurring after 72 hours of hospital admission or within 4 weeks of hospitalization, were included.

Methods: The incidence of HA-CDI and antibiotic utilization from 2003 to 2020 were analyzed with available *C. difficile* isolates. The impact of antibiotic utilization on HA-CDI incidence was estimated by a dynamic regression time-series model. Antibiotic utilization and the proportion of NAP1/027 strains were compared biannually for available isolates from 2010 to 2020.

Results: The incidence of HA-CDI decreased between 2003 and 2020 at both hospitals from 26.5 cases per 10,000 patient-days in 2003 to 4.9 cases per 10,000 patient-days in 2020 respectively. Over the study period, there were an increase in the utilization of third-generation cephalosporins and a decrease in usage of fluoroquinolones and clindamycin. A decrease in fluoroquinolone utilization was associated with a significant decrease in HA-CDI incidence as well as decrease in the NAP1/027 strain by approximately 80% in both hospitals.

Conclusions: Decreased utilization of fluoroquinolones in two Québec hospitals was associated with a decrease in the incidence of HA-CDI and a genotype shift from NAP1/027 to non-NAP1/027 strains.

(Received 15 February 2024; accepted 25 April 2024)

#### Introduction

*Clostridioides difficile* is the leading infectious cause of healthcareassociated (HA) diarrhea in developed countries.<sup>1–3</sup> It is associated with significant morbidity, with clinical manifestations ranging from asymptomatic colonization, severe diarrhea, to toxic megacolon, and death.<sup>1</sup> In 2003, many hospitals in Québec, Canada, experienced an epidemic of *C. difficile* infection (CDI) associated with increased severity of disease and recurrences.<sup>4–6</sup> A fourfold increase in *C. difficile* attributable mortality was observed in Canada between 1997 and 2005.<sup>7</sup> This epidemic was associated with the dissemination of the hypervirulent North American Pulsed Field

Corresponding author: Vivian G. Loo; Email: vivian.loo@mcgill.ca

Type 1 (NAP1)/ribotype 027 strain.<sup>8</sup> Outbreaks associated with this strain have been reported worldwide.<sup>9</sup> Recent studies from the United States demonstrate a decreasing trend for HA-CDI in the last decade with a reduced prevalence of the NAP1/027 strain.<sup>10,11</sup>

Antibiotic use is the major risk factor associated with CDI. Antibiotic classes historically associated with high risk of CDI include cephalosporins and clindamycin.<sup>12</sup> The epidemic dissemination of the NAP1/027 strain was found to be related to fluoroquinolone utilization.<sup>13,14</sup> Interventions aimed at restricting the use of these high-risk antibiotics have been associated with a marked decrease in the incidence of CDI in Québec, the United States, and Europe.<sup>15–18</sup> A UK study combining whole genome sequencing and antimicrobial prescribing data revealed that CDI regression in England was driven by the effect of fluoroquinolone restriction on fluoroquinolone-resistant strains.<sup>19</sup> In the United States, inpatient fluoroquinolone use was found to be a predictor of

© The Author(s), 2024. Published by Cambridge University Press on behalf of The Society for Healthcare Epidemiology of America. This is an Open Access article, distributed under the terms of the Creative Commons Attribution-NonCommercial-NoDerivatives licence (http://creativecommons.org/licenses/by-nc-nd/4.0/), which permits non-commercial re-use, distribution, and reproduction in any medium, provided that no alterations are made and the original article is properly cited. The written permission of Cambridge University Press must be obtained prior to any commercial use and/or adaptation of the article.

Data was presented in a preliminary abstract at IDWeek 2021.

**Cite this article:** Couture S, Frenette C, Schiller I, *et al.* The changing epidemiology of *Clostridioides difficile* infection and the NAP1/027 strain in two Québec hospitals: a 17-year time-series study. *Antimicrob Steward Healthc Epidemiol* 2024. doi: 10.1017/ash.2024.95

HA-CDI due to the NAP1/027 strain.<sup>20</sup> Furthermore, a recent large-scale 6-year ecological study performed in over 500 US hospitals demonstrated that temporal changes in antibiotic utilization correlated with changes in HA-CDI.<sup>21</sup> These studies support antibiotic stewardship as a major factor in reducing HA-CDI incidence. However, reductions in the prevalence of the NAP1/027 strain were observed in absence of nationwide reductions in fluoroquinolones use.<sup>22</sup> Other studies and experts have advocated that other unmeasured factors (eg, infection control interventions) were likely implicated in the observed decrease in HA-CDI incidence.<sup>23,24</sup>

The *C. difficile* epidemic in Québec peaked in 2004–2005 with an incidence of 12.6 cases per 10,000 patient-days.<sup>25</sup> In the past 15 years, the incidence of HA-CDI has decreased in Québec, reaching 3.14 cases per 10,000 patient-days in 2019–2020.<sup>26</sup> A series of infection control measures, including an antibiotic stewardship program,<sup>26</sup> were implemented at our institutions over this period in response to the Québec epidemic. The aim of this study was to assess the impact of antibiotic utilization on HA-CDI incidence and the NAP1/027 strain in two major academic centers in Montreal, Québec, Canada.

#### Methods

#### Population and surveillance period

We conducted an observational study with antibiotic utilization data to study its effect on both HA-CDI incidence and *C. difficile* genotypes at the Royal Victoria Hospital (RVH) and Montreal General Hospital (MGH), two adult tertiary care hospitals of the McGill University Health Centre (MUHC), in Montreal, Canada between April 1, 2003, and March 31, 2020.

#### HA-CDI incidence

Patients with a positive assay for toxigenic C. difficile were identified by the MUHC laboratory and reported to the infection prevention and control service. Laboratory detection was performed by cell cytotoxin neutralization assay (CCNA) until June 2009, following which the assay was changed to a commercial nucleic acid amplification test (NAAT). Patients were considered to have HA-CDI if they met one of the following criteria: diarrhea with at least three liquid to semi-liquid stools in 24 hours without any other apparent etiology, or toxic megacolon with a positive C. difficile toxin assay, a change in stool characteristics from baseline without apparent cause in a patient known for chronic diarrhea with a positive C. difficile toxin assay, endoscopic evidence of pseudomembranous colitis, or histopathologic evidence of C. difficile colitis.<sup>25</sup> A case was considered as HA if symptoms occurred more than 72 hours after admission or within 4 weeks of a previous hospitalization. Outpatients with CDI who were not associated with a previous hospitalization in the last 4 weeks, and patients admitted to the obstetrics and psychiatry wards were excluded from the study. HA-CDI incidence was calculated as the number of cases per 10,000 patient-days.

# Antibiotic utilization

Antibiotic utilization over the study period was extracted from MUHC pharmacy data. Antibiotic consumption was measured in defined daily doses (DDDs) per 1,000 patient-days. We focused on the following antibiotic classes: first-generation cephalosporins, second-generation cephalosporins, third-generation cephalosporins, narrow spectrum penicillins (penicillin, ampicillin, cloxacillin, amoxicillin), beta-lactam/beta-lactamase inhibitors, carbapenems, fluoroquinolones, aminoglycosides, sulfonamides (trimethoprimsulfamethoxazole), tetracyclines, macrolides, lincosamides (clindamycin), and glycopeptides (vancomycin). These classes of antibiotics were selected based on the frequency of their use and suspected risk for CDI.<sup>12</sup> We excluded oral vancomycin from total antibiotic utilization given the assumed direct correlation of CDI incidence with its usage, as well as metronidazole due to the inability to determine retrospectively what proportion of its utilization was for the treatment of CDI.

#### C. difficile isolate genotyping

All stool samples that were positive for *C. difficile* toxin gene detection by NAAT at our institution were routinely genotyped by the infection control service for outbreak surveillance from 2010 at one institution and from 2014 at the second institution. These isolates were typed by pulsed-field electrophoresis using standard methods from January 2010 to May 2012.<sup>27</sup> Ribotyping by standard methods was used from June 2012 to March 2020.<sup>28</sup>

#### Infection control interventions

Infection control interventions implemented at our institution over the study period were reviewed.

# Statistical analysis

A dynamic regression model for time series was used to estimate the impact of antibiotic usage (predictor) on CDI incidence (outcome) over time (quarterly data from 2003-2004 to 2019-2020 for both sites) while taking into account the possibility of autocorrelated error terms.<sup>29</sup> Recognizing that a change in antibiotic usage may not have an immediate effect on CDI incidence, we performed a multiplier analysis by allowing for a two-period lagged effect, with each period representing roughly 3 months. It allowed us to estimate the impact on the CDI incidence we could expect to observe if antibiotic usage was equally increased contemporaneously, 3 months and 6 months ago. We compared antibiotic usage and the proportion of NAP1/027 strains on a biannually basis from 2014-2015 to 2019-2020 for the MGH site and from 2010-2011 to 2019-2020 for the RVH site by calculating the observed change in percentage in both variables between 2014-2015 and 2019-2020 for MGH and between 2010-2011 and 2019-2020 for RVH. A Sieve-bootstrap Mann-Kendall's trend test was used to test for monotonic trend in antibiotic usage and proportion of NAP1/027 strains.<sup>30</sup> Analyses were performed using the R statistical software version 4.0.3.<sup>31</sup> Model selection was performed using the stepwise algorithm outlined in Hyndman and Khandakar<sup>32</sup> and implemented in the *forecast* R package.<sup>33</sup>

#### Ethics approval

Institutional review board approval was obtained for this study. All patient information was de-identified.

#### Results

#### Healthcare-associated CDI incidence

The incidence of HA-CDI decreased between April 2003 and March 2020 at both institutions. At the MGH, the incidence decreased from 19.1 cases per 10,000 patient-days in the first quarter of 2003 to 3.0 cases per 10,000 patient-days in the last quarter of 2020. At the RVH, the incidence decreased from 35.7 cases per 10,000 patient-days in the first quarter of 2003 to 4.8 cases

Table 1. Antibiotic utilization at the MGH and RVH between 2003 and	2020
---	------

	MGH			RVH		
Antibiotic	Use in 2003–2004 first quarter (DDDs per 1,000 patient-days)	Use in 2019–2020 last quarter (DDDs per 1,000 patient-days)	Change (%)	Use in 2003–2004 first quarter (DDDs per 1,000 patient-days)	Use in 2019–2020 last quarter (DDDs per 1,000 patient-days)	Change (%)
Penicillins	67.38	32.93	-51.13	100.93	76.15	-24.55
Beta-lactam/beta-lactamase inhibitors	80.61	147.03	82.4	68.36	192.79	182.04
Carbapenems	12.20	47.68	290.74	30.33	90.80	199.37
First-generation cephalosporins	101.58	88.19	-13.18	57.85	89.31	54.37
Second-generation cephalosporins	5.08	4.75	-6.56	9.04	3.28	-63.71
Third-generation cephalosporins	14.79	66.78	351.46	13.14	68.90	424.27
Fluoroquinolones	102.18	47.50	-53.51	122.63	78.80	-35.74
Clindamycin	10.72	8.53	-20.48	16.22	7.60	-53.11
Macrolides	12.38	23.20	87.42	45.16	46.52	3.03
Tetracyclines	5.88	41.68	609.19	4.40	43.92	898.09
Sulfonamides	29.57	35.52	20.11	33.37	57.06	70.97
Aminoglycosides	21.16	9.43	-55.44	15.55	40.28	159.07
Total	585.97	631.75	7.81	705.70	900.85	27.65

Note. MGH, Montreal General Hospital; RVH, Royal Victoria Hospital; DDDs, defined daily doses.

per 10,000 patient-days in the last quarter of 2020. At both institutions, a transient rise in the incidence of HA-CDI was observed between 2009 and 2012 after implementation of *C. difficile* laboratory detection by NAAT.

# Antibiotic utilization

Antibiotic utilization at the RVH and MGH between April 2003 and March 2020 is presented in Table 1. A decrease in the incidence of HA-CDI between April 2003 and March 2020 occurred, although total antibiotic utilization was increasing at both hospitals, from 585.97 to 631.75 DDDs per 1,000 patient-days at the MGH and from 705.70 to 900.85 DDDs per 1,000 patientdays at the RVH. At both institutions, we observed an increase in consumption of third-generation cephalosporins (14.79-66.78 DDDs per 1,000 patient-days at the MGH, 13.14-68.90 DDDs per 1,000 patient-days at the RVH), beta-lactam/beta-lactamase inhibitors (80.61-147.03 DDDs per 1,000 patient-days at the MGH, 68.36-192.79 DDDs per 1,000 patient-days at the RVH), and carbapenems (12.20-47.68 DDDs per 1,000 patient-days at the MGH, 30.33-90.80 DDDs per 1,000 patient-days at the RVH). At both hospitals, a decrease in utilization of fluoroquinolones (102.18-47.50 DDDs per 1,000 patient-days at the MGH, 122.63-78.80 DDDs per 1,000 patient-days at the RVH) and clindamycin (10.72-8.53 DDDs per 1,000 patient-days at the MGH, 16.22-7.6 DDDs per 1,000 patient-days at the RVH) was observed over the study period.

# Effect of antibiotic utilization on HA-CDI incidence

Utilization of high-risk antibiotics and HA-CDI incidence at the MGH and RVH are presented in Figures 1 and 2, respectively. At both institutions, a decrease of 10 DDDs in fluoroquinolone

utilization was associated with a decrease in HA-CDI incidence (MGH impact -1.40 cases per 10,000 patient-days, 95% CI [-2.44, -0.35]; RVH impact -0.98 cases per 10,000 patient-days, 95% CI [-1.68, -0.28]). A decrease of 10 DDDs in the utilization of firstgeneration cephalosporins was associated with a decrease in HA-CDI at the MGH (impact -1.87 cases per 10,000 patient-days, 95% CI [-3.62, -0.12]). Such an association was not observed at the RVH. A change in the utilization of second-generation cephalosporins was not associated with a significant change in HA-CDI incidence. A decrease in 10 DDDs of third-generation cephalosporin utilization was associated with an increase in HA-CDI incidence at the MGH (impact 2.30 cases per 10,000 patientdays, 95% CI [0.73, 3.86]). There was no significant change in HA-CDI following the change in third-generation cephalosporin utilization at the RVH. Changes in clindamycin utilization were not associated with a significant change in HA-CDI incidence at either institution.

# Antibiotic utilization and impact on the NAP1/027 strain

Between April 2014 and March 2020 at the MGH, there was a significant decrease in the proportion of the NAP1/027 strain (-85.7%, p = 0.046). During the same period, there was a decrease in fluoroquinolone utilization (-47.79%, p = 0.013), a significant increase in third-generation cephalosporin utilization (160.81%, p = 0.001), and no significant change in clindamycin utilization (-10.97%, p = 0.1) (Figure 3). At the RVH, between April 2010 and March 2020, there was a decrease in the proportion of the NAP1/027 strain (-75%, p = 0). During the same period, there was a trend in decreased fluoroquinolone utilization (-48.13%, p = 0.18), a significant increase in third-generation cephalosporin utilization (182.18%, p = 0), and a significant decrease in clindamycin utilization (-43.21%, p = 0.002) (Figure 4).



Figure 1. Healthcare-associated *C. difficile* infection incidence and high-risk antibiotic utilization at the MGH between 2003 and 2020. *Antibiotic:* — cephalosporin first generation, — cephalosporin second generation, — cephalosporin third generation, — clindamycin, — fluoroquinolone. *Note: C. difficile, Clostridioides difficile;* MGH, Montreal General Hospital.



Figure 2. Healthcare-associated *C. difficile* infection incidence and high-risk antibiotic utilization at the RVH between 2003 and 2020. *Antibiotic:* — cephalosporin first generation, — cephalosporin second generation, — cephalosporin third generation, — clindamycin, — fluoroquinolone. *Note: C. difficile, Clostridioides difficile;* RVH, Royal Victoria Hospital.

#### Infection control interventions

Infection control interventions that were implemented during the study period are presented in Table 2.

## Discussion

Our 17-year analysis of the epidemiology of HA-CDI in two tertiary care hospitals in Montreal, Québec, demonstrated an approximate 80% decrease in the incidence of HA-CDI at both institutions. This decline occurred while total antibiotic consumption was increasing, but use of certain high-risk antibiotics fluoroquinolones, second-generation cephalosporins, and clindamycin—was decreasing. A decrease in fluoroquinolone utilization was associated with a significant decrease in HA-CDI incidence at the 6-month interval at both hospitals. A decrease in firstgeneration cephalosporins was associated with a decrease in



Figure 3. Proportion of NAP1/027 C. difficile genotype and utilization of high-risk antibiotics at the MGH between 2014 and 2020. Antibiotic: — cephalosporin third generation, — clindamycin, — fluoroquinolone. Note: C. difficile, Clostridioides difficile; MGH, Montreal General Hospital.



Figure 4. Proportion of NAP1/027 C. difficile genotype and utilization of high-risk antibiotics at the RVH between 2010 and 2020. Antibiotic: — cephalosporin third generation, — clindamycin, — fluoroquinolone. Note: C. difficile, Clostridioides difficile; RVH, Royal Victoria Hospital.

HA-CDI incidence at one of the two centers. In the later years of our study, we observed a significant decrease over time in the *C*. *difficile* NAP1/027 strain which is inherently resistant to fluoroquinolones. There was a parallel significant decrease in fluoroquinolone utilization at one of the two hospitals.

Our findings are consistent with previous studies reporting an association between HA-CDI and the use of high-risk antibiotic classes, particularly fluoroquinolones and cephalosporins.<sup>20,21,34</sup> Interventions targeting the restriction in the utilization of these antibiotics have been successful at driving a decline in HA-CDI in

North America and Europe.<sup>16,17,19,34–36</sup> In Québec, at the height of the HA-CDI epidemic between 2003 and 2006, implementation of antibiotic stewardship led to a decrease in the utilization of high-risk antibiotics and a decrease in HA-CDI incidence.<sup>15</sup>

The effect of fluoroquinolone utilization is of particular interest as the HA-CDI epidemic in Québec was associated with dissemination of the NAP1/027 strain which is known to be resistant to this antibiotic class.<sup>13</sup> Our findings demonstrated an association between a decrease in fluoroquinolone utilization and a decrease in total *C. difficile* incidence. We were further able to

#### Table 2. Infection control interventions implemented at the MUHC between April 1, 2003, and March 31, 2020

Voor	Intervention
2004	Contact precautions for all patients with diarrhea     Single room or cohort rooms for patients with diarrhea
	Revision of antibiotic usage
	Two-step cleaning
	Routine use of hypochlorite for daily cleaning
	Increased frequency of bathroom cleaning
2007	Early outbreak identification
2008	Three-step cleaning
	Glow germ fluorescent markers to assess adequacy of surface cleaning
2009	Initiation of a formal antibiotic stewardship program
	Laboratory detection of <i>C. difficile</i> changed to NAAT test from CCNA
2010	Antibiotic stewardship audits and feedback on targeted units
2012	Hand byging program with audits and foodback
	Systematic use of audits for cleaning and disinfection
	Continued antibiotic stewardship
2013	Use of daily chlorbexidine wipes for patients on outbreak units
	Implementation of TCAB-CSI* structure on targeted units
	Clostridioides difficile precautions maintained until discharge
2014	Reinstitution of antibiotic stewardship on targeted units
	Dedicated housekeeping team for cleaning patient rooms at discharge
	Systematic audit program for housekeeping
2015	<ul> <li>Move to a single-room facility for one hospital</li> </ul>
	TCAB-CSI* interventions implemented on all units
	Ongoing systematic audits of housekeeping and hand hygiene
	Orderly training program for cleaning and disinfection
2016	
2010	Checklists and systematic quality control for disinfection of environment and equipment     Nocospray® (MedPro Defense Montreal Canada) for terminal disinfection of C difficile and VDE rooms
2017	- necospray- (neci to belense, montreat, canada) for terminal disinfection of C. difficite and the rooms
2017	Ongoing implementation of Nocospray®
2018	Increased use of hydrogen peroxide cleaning for equipment weekly and at discharge
	Assigning dedicated housekeepers to clean patient equipment

Note. MUHC, McGill University Health Centre; NAAT, nucleic acid amplification test; VRE, vancomycin-resistant Enterococcus; CDI, Clostridioides difficile infection.

\*Transforming care at the bedside: weekly huddles where infection control would meet with the nurse manager, attending physician, senior resident, and other allied health professionals to review CDI rates and address interventions that could be improved.

demonstrate that a significant decrease in the incidence of the NAP1/027 strain was concomitant with decrease in fluoroquinolone utilization at one of our two hospital sites in the final years of the study. There was a trend in decreased fluoroquinolone utilization over time with the decrease in proportion of isolates of the NAP1/027 strain at our second study hospital, but this did not reach statistical significance. Our findings are similar to the results of a study conducted in the United Kingdom between 2006 and 2013. This study demonstrated that the decline in *C. difficile* incidence was primarily driven by the elimination of fluoroquinolone-resistant strains.<sup>19</sup>

In our study, the association between the utilization of cephalosporins and HA-CDI incidence is equivocal. The change in the utilization of the different cephalosporin classes had inconsistent effects across the two hospitals. We found that an increase in third-generation cephalosporin use was associated with a decrease in HA-CDI incidence at the MGH. This contrasts with previous literature that supported the restriction of third-generation cephalosporins to control HA-CDI.<sup>21,34</sup> One hypothesis is that third-generation cephalosporins replaced fluoroquinolones

in the treatment of urinary and pulmonary pathogens at our institution. Since we did not determine the specific effect of an antibiotic independent of changes in other antibiotic classes, it is possible that the potential effect of the increase in utilization of third-generation cephalosporins on HA-CDI was offset by the concomitant effect of decreasing fluoroquinolones on the predominant NAP1/027 strain.

Our findings regarding the association between the reduction of use of high-risk antibiotics and incidence of HA-CDI contrast with a coincident study conducted in 12 Québec hospitals between 2012 and 2017. That study demonstrated that the change in high-risk antibiotics did not entirely account for the decrease of HA-CDI in the province.<sup>23</sup> These different results may be explained by the shorter study duration of 5 years in the Fortin et al publication. Our study was of longer duration and included more time points for analyses.

We reviewed infection control interventions that were implemented at our institution during the study period and identified measures other than antibiotic stewardship that were potentially at play in the changing epidemiology of HA-CDI. Both of our hospital sites experienced an increase in HA-CDI incidence between 2009 and 2012. This temporally correlates with the change to the NAAT test from CCNA for the detection of *C. difficile* in 2009. US data from the Center for Disease Control and Prevention suggested that such a change in testing method could account for a rise in HA-CDI ranging from 43% to 67%.<sup>37</sup> In Québec, 35 institutions modified their testing algorithm between 2010 and 2014 and those detecting toxigenic *C. difficile* instead of *C. difficile* toxin had higher rates of HA-CDI.<sup>38</sup> Despite the initial increase in HA-CDI incidence with the change in testing, HA-CDI incidence in our institutions continued to decline after 2012.

Another possible contributor to the decline in HA-CDI incidence was the move to a single-room facility for the RVH site in 2015. A previous time-series analysis conducted at our institution did not demonstrate an association of HA-CDI incidence and housing patients in single rooms.<sup>39</sup>

There are several limitations to this investigation. Firstly, only two hospitals were included in this analysis. More data is needed from multiple institutions to clarify whether the reduction in fluoroquinolone use has been a major driving factor to the decrease in HA-CDI incidence and the NAP1/027 strain in North America and Europe. Secondly, there were multifaceted infection control interventions that were implemented over the study period. Hence, the effect of changes in antibiotic utilization and HA-CDI incidence are subject to potential confounders that are difficult to measure. Previous epidemiological studies have also demonstrated that there are shifts in epidemic C. difficile strains in the absence of substantial changes in antibiotic use.<sup>40</sup> Furthermore, the change in laboratory diagnostic method for C. difficile detection to NAAT which is a more sensitive test than CCNA likely led to a transient increase in incidence during the study period and may have blunted certain associations between antibiotic utilization and HA-CDI. In addition, we did not evaluate for associations between antibiotic classes and the impact of the change of an antibiotic class independent of the change of other classes. Lastly, the retrospective nature of the study led to limited accessibility to C. difficile strain genotyping from the earlier years of the study. Due to this significant amount of unavailable data, we could not evaluate the effect of changes in antibiotic utilization on the NAP1/ 027 strain during the entire study period. Despite this, we were still able to demonstrate an association of the decrease in the proportion of C. difficile isolates of the NAP1/027 genotype with the decrease in fluoroquinolone utilization in one hospital and a trend in the second hospital.

In conclusion, the decline in HA-CDI following the epidemic of the early 2000s at two hospitals in Montreal, Canada, was accompanied by a reduction in the utilization of high-risk antibiotics. Specifically, a decrease in the use of fluoroquinolones was associated with a significant decrease in HA-CDI incidence. Moreover, at one of our study sites, the decrease in fluoroquinolone utilization was also associated with a parallel decrease in the proportion of *C. difficile* isolates of the NAP1/027 genotype. Our study demonstrates that antibiotic stewardship is an essential infection control strategy to decrease HA-CDI incidence and can effectively suppress hypervirulent strains at a facility level.

Acknowledgments. None.

**Author contributions.** No one other than the named authors had a role in the gathering or preparation of data or in the writing of the manuscript.

Financial support. There was no funding received for this study.

# References

- Kelly CP, Pothoulakis C, LaMont JT. Clostridium difficile colitis. New Engl J Med 1994;330:257–262. https://doi.org/10.1056/nejm199401273300406.
- Poutanen SM, Simor AE. Clostridium difficile-associated diarrhea in adults. CMAJ: Can Med Assoc J = J De L'Assoc Med Can 2004;171:51–58.
- McFarland LV. Epidemiology of infectious and iatrogenic nosocomial diarrhea in a cohort of general medicine patients. *Am J Infect Control* 1995;23:295–305.
- Valiquette L, Low DE, Pepin J, McGeer A. *Clostridium difficile* infection in hospitals: a brewing storm. *CMAJ: Can Med Assoc J = J De L'Assoc Med Can* 2004;171:27–29.
- Pepin J, Valiquette L, Alary ME, et al. Clostridium difficile-associated diarrhea in a region of Quebec from 1991 to 2003: a changing pattern of disease severity. CMAJ: Can Med Assoc J = J De L'Assoc Med Can 2004;171:466–472. https://doi.org/10.1503/cmaj.1041104.
- Pepin J, Valiquette L, Cossette B. Mortality attributable to nosocomial *Clostridium difficile*-associated disease during an epidemic caused by a hypervirulent strain in Quebec. *CMAJ: Can Med Assoc J = J De L'Assoc Med Can* 2005;173:1037–1042. https://doi.org/10.1503/cmaj.050978.
- Gravel D, Miller M, Simor A, et al. Health care-associated Clostridium difficile infection in adults admitted to acute care hospitals in Canada: a Canadian Nosocomial Infection Surveillance Program Study. Clin Infect Dis 2009;48:568–576. https://doi.org/10.1086/596703.
- Warny M, Pepin J, Fang A, et al. Toxin production by an emerging strain of Clostridium difficile associated with outbreaks of severe disease in North America and Europe. Lancet (London, England) 2005;366:1079–1084. https://doi.org/10.1016/s0140-6736(05)67420-x.
- He M, Miyajima F, Roberts P, et al. Emergence and global spread of epidemic healthcare-associated *Clostridium difficile*. Nat Genet 2013;45: 109–113. https://doi.org/10.1038/ng.2478.
- Giancola SE, Williams RJ, 2nd, Gentry CA. Prevalence of the *Clostridium difficile* BI/NAP1/027 strain across the United States Veterans Health Administration. *Clin Microbiol Infect* 2018;24:877–881. https://doi.org/10. 1016/j.cmi.2017.11.011.
- Guh AY, Mu Y, Winston LG, et al. Trends in U.S. Burden of Clostridioides difficile infection and outcomes. New Engl J Med 2020;382:1320–1330. https://doi.org/10.1056/NEJMoa1910215.
- Owens RC, Jr., Donskey CJ, Gaynes RP, Loo VG, Muto CA. Antimicrobialassociated risk factors for *Clostridium difficile* infection. *Clin Infect Dis* 2008;1:S19–S31. https://doi.org/10.1086/521859.
- Loo VG, Poirier L, Miller MA, et al. A predominantly clonal multiinstitutional outbreak of *Clostridium difficile*-associated diarrhea with high morbidity and mortality. *New Engl J Med* 2005;353:2442–2449. https://doi. org/10.1056/NEJMoa051639.
- Pepin J, Saheb N, Coulombe MA, et al. Emergence of fluoroquinolones as the predominant risk factor for *Clostridium difficile*-associated diarrhea: a cohort study during an epidemic in Quebec. *Clin Infect Dis* 2005;41: 1254–1260. https://doi.org/10.1086/496986.
- Valiquette L, Cossette B, Garant MP, Diab H, Pepin J. Impact of a reduction in the use of high-risk antibiotics on the course of an epidemic of *Clostridium difficile*-associated disease caused by the hypervirulent NAP1/027 strain. *Clin Infect Dis* 2007;2:S112–S121. https://doi.org/10.1086/519258.
- 16. Kallen AJ, Thompson A, Ristaino P, et al. Complete restriction of fluoroquinolone use to control an outbreak of *Clostridium difficile* infection at a community hospital. *Infect Control Hosp Epidemiol* 2009;30:264–272. https://doi.org/10.1086/595694.
- 17. Schönherr SG, Ranft D, Lippmann N, Lübbert C. Changes in antibiotic consumption, AMR and *Clostridioides difficile* infections in a large tertiarycare center following the implementation of institution-specific guidelines for antimicrobial therapy: a nine-year interrupted time series study. *PLoS One* 2021;16:e0258690. https://doi.org/10.1371/journal.pone.0258690.

- Dancer SJ, Kirkpatrick P, Corcoran DS, Christison F, Farmer D, Robertson C. Approaching zero: temporal effects of a restrictive antibiotic policy on hospital-acquired *Clostridium difficile*, extended-spectrum β-lactamase-producing coliforms and methicillin-resistant Staphylococcus aureus. *Int J Antimicrob Agents* 2013;41:137–142. https://doi.org/10.1016/j.ijantimicag. 2012.10.013.
- Dingle KE, Didelot X, Quan TP, et al. Effects of control interventions on Clostridium difficile infection in England: an observational study. Lancet Infect Dis 2017;17:411–421. https://doi.org/10.1016/S1473-3099(16)30514-X.
- Silva SY, Wilson BM, Redmond SN, Donskey CJ. Inpatient fluoroquinolone use in Veterans' Affairs hospitals is a predictor of *Clostridioides difficile* infection due to fluoroquinolone-resistant ribotype 027 strains. *Infect Control Hosp Epidemiol* 2021;42:57–62. https://doi.org/10.1017/ice.2020.383.
- Kazakova SV, Baggs J, McDonald LC, et al. Association between antibiotic use and hospital-onset Clostridioides difficile infection in US acute care hospitals, 2006–2012: an ecologic analysis. Clin Infect Dis 2020;70:11–18. https://doi.org/10.1093/cid/ciz169.
- Hensgens MP, Goorhuis A, Notermans DW, van Benthem BH, Kuijper EJ. Decrease of hypervirulent *Clostridium difficile* PCR ribotype 027 in the Netherlands. *Euro Surveill* 2009;14. https://doi.org/10.2807/ese.14.45.19402-en.
- Fortin E, Thirion DJG, Ouakki M, et al. Role of high-risk antibiotic use in incidence of health-care-associated *Clostridioides difficile* infection in Quebec, Canada: a population-level ecological study. *Lancet Microbe* 2021;2:e182–e90. https://doi.org/10.1016/s2666-5247(21)00005-7.
- Eyre DW, Dingle KE, Didelot X, et al. Clostridium difficile in England: can we stop washing our hands?—Authors' reply. Lancet Infect Dis 2017;17: 478–479. https://doi.org/10.1016/s1473-3099(17)30185-8.
- 25. Comité de surveillance provinciale des infections nosocomiales (SPIN). Surveillance des diarrhées associées à *Clostridium difficile* au Québec: Bilan du 22 août 2004 au 18 août 2007: Institut national de santé publique du Québec (INSPQ); 2008 [cited 2021]. https://www.inspq.qc.ca/sites/default/ files/publications/745\_cdifficile\_bilan2004-2007.pdf.
- 26. Comité de surveillance provinciale des infections nosocomiales (SPIN). Diarrhées associées au *Clostridium difficile*: Résultats de surveillance 2020– 2021. https://www.inspq.qc.ca/publications/3374.
- Corkill JE, Graham R, Hart CA, Stubbs S. Pulsed-field gel electrophoresis of degradation-sensitive DNAs from *Clostridium difficile* PCR ribotype 1 strains. J Clin Microbiol 2000;38:2791–2792.
- Fawley WN, Knetsch CW, MacCannell DR, et al. Development and validation of an internationally-standardized, high-resolution capillary gel-based electrophoresis PCR-ribotyping protocol for *Clostridium difficile*. *PLoS One* 2015;10:e0118150. https://doi.org/10.1371/journal.pone.0118150.

- 29. Hyndman RJ, Athanasopoulos G. Forecasting: principles and practice: OTexts; 2018.
- Noguchi K, Y.R. Gel, C.R. Duguay. Bootstrap-based test for trends in hydrological times series, with application to ice phenology data. *J Hydrol* 2011;410. https://doi.org/10.1016/j.jhydrol.2011.09.008.
- R Core Team. R: A language and environment for statistical computing Vienna, Austria: R Foundation for Statistical Computing; 2020 [cited 2022]. https://www.R-project.org/.
- 32. Hyndman R AG, Bergmeir C, Caceres G, *et al.* Yasmeen F forecast: Forecasting functions for time series and linear models: R package version 8.15; 2021. https://pkg.robjhyndman.com/forecast/.
- Hyndman RJ, Khandakar Y. Automatic time series forecasting: the forecast package for R. J Stat. Softw 2008;27:1–22.
- Pereira JB, Farragher TM, Tully MP, Jonathan Cooke J. Association between *Clostridium difficile* infection and antimicrobial usage in a large group of English hospitals. *Br J Clin Pharmacol* 2014;77:896–903. https://doi.org/10. 1111/bcp.12255.
- 35. Muto CA, Blank MK, Marsh JW, et al. Control of an outbreak of infection with the hypervirulent Clostridium difficile BI strain in a university hospital using a comprehensive "bundle" approach. Clin Infect Dis 2007;45: 1266–1273. https://doi.org/10.1086/522654.
- 36. Lawes T, Lopez-Lozano JM, Nebot CA, et al. Effect of a national 4C antibiotic stewardship intervention on the clinical and molecular epidemiology of Clostridium difficile infections in a region of Scotland: a non-linear time-series analysis. Lancet Infect Dis 2017;17:194–206. https:// doi.org/10.1016/s1473-3099(16)30397-8.
- 37. Gould CV, Edwards JR, Cohen J, et al. Effect of nucleic acid amplification testing on population-based incidence rates of *Clostridium difficile* infection. *Clin Infect Dis* 2013;57:1304–1307. https://doi.org/10.1093/cid/ cit492.
- Bogaty C, Lévesque S, Garenc C, et al. Trends in the use of laboratory tests for the diagnosis of *Clostridium difficile* infection and association with incidence rates in Quebec, Canada, 2010–2014. Am J Infect Control 2017;45:964–968. https://doi.org/10.1016/j.ajic.2017.04.002.
- McDonald EG, Dendukuri N, Frenette C, Lee TC. Time-series analysis of health care-associated infections in a new hospital with all private rooms. *JAMA Intern Med* 2019;179:1501–1506. https://doi.org/10.1001/ jamainternmed.2019.2798.
- Belmares J, Johnson S, Parada JP, et al. Molecular epidemiology of Clostridium difficile over the course of 10 years in a tertiary care hospital. Clin Infect Dis 2009;49:1141–1147. https://doi.org/10.1086/ 605638.