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- 1 Title: Circulation of highly drug-resistant *Clostridium difficile* ribotypes 027 and 001 in
- 2 two tertiary-care hospitals in Mexico.
- 3 **Running title:** *Clostridium difficile* drug resistance in Mexico.
- 4
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34 Abstract

Objective: To assess drug susceptibility and characterize *C. difficile* ribotypes in
isolates from two tertiary-care hospitals in Mexico.

37 Methods: Isolates were evaluated for genotyping, antimicrobial susceptibility
38 testing and detection of mutations associated with drug resistance. PCR ribotyping was
39 performed using a combination of gel-based and capillary electrophoresis-based
40 approaches.

**Results:** MIC<sub>50</sub> and MIC<sub>90</sub> were  $\geq$ 128 mg/L for ciprofloxacin, erythromycin, 41 clindamycin, and rifampicin. There was no reduced susceptibility to metronidazole or 42 tetracycline; however, reduced susceptibility to vancomycin (>4 mg/L) and fidaxomicin 43 (>2 mg/L) was detected in 50 (40.3%) and 4 (3.2%) isolates respectively. Furthermore, 44 the rpoB Arg505Lys mutation was more frequently detected in isolates with high MIC to 45 rifampicin (>32 mg/L) (OR = 52.5: 95% CI = 5.17-532.6: p < 0.000). 46 Of the 124 C. difficile isolates recovered; 84 (66.7%) were of ribotype 027, 18 47 (14.5%) of ribotype 001, and the remainder were other ribotypes (353, 255, 220, 208, 176, 48

49 106, 076, 020, 019, 017, 014, 012, 003, and 002).

50 **Conclusion:** Ribotypes 027 and 001 were the most frequent *C. difficile* isolates 51 recovered in this study, and demonstrated higher MICs. Furthermore, we found four 52 isolates with reduced susceptibility to fidaxomicin, raising a concern since this drug is 53 currently unavailable in Mexican Hospitals.

54

Keywords: Drug resistance; Ribotypes; Fidaxomicin; Ribotype 001; *Clostridium difficile*.

#### 57 Introduction

*C. difficile* infection (CDI) symptoms may range from mild diarrhea to lifethreatening complications. Apart from NAP1/BI/027, other C. *difficile* ribotypes have been
associated with severe disease, *e.g.* ribotype 078 affects younger patients and is a frequent
causative agent of community-associated disease <sup>1</sup>; ribotype 001 is the dominant strain in
eastern Europe and has higher antimicrobial resistance than other ribotypes <sup>2</sup>.

First-line treatment for mild to moderate CDI is based on oral administration of 63 metronidazole or vancomycin, with a therapeutic efficacy >70%<sup>3</sup>. In some patients, 64 however, diarrheal symptoms may reappear within days or weeks after having stopped the 65 treatment. Fidaxomicin is a relatively new narrow-spectrum macrocyclic antibiotic drug 66 that is non-inferior to vancomycin in the management of CDI and associated with lower 67 recurrence rates than vancomycin<sup>4</sup>. However, fidaxomicin is currently unavailable in 68 Mexico, thus vancomvcin and metronidazole are still the standard treatments for CDI as 69 70 recommended in the Clinical Practice Guidelines for C. difficile Infection in Adults of the Society for Healthcare Epidemiology of America (SHEA) and the Infectious Diseases 71 Society of America (IDSA). Other therapeutic options that have been proposed in recent 72 73 years are rifamycins (good *in vitro* activity against *C. difficile*) including rifaximin (for relapsing CDI)<sup>5</sup>, and linezolid (protective rather than curative activity)<sup>6</sup>. 74

Resistance to erythromycin may be due to any of more than 20 classes of erythromycin ribosomal methylase (*erm*) genes, including *ermB*<sup>7</sup>, which is also related to clindamycin resistance. In *C. difficile*, resistance to fluoroquinolones is usually due to altered DNA gyrase because of nucleotide substitutions in *gyrA* or *gyrB* genes <sup>5</sup>. Resistance to rifamycins or fidaxomicin is mediated by mutations that lead to reduced binding to the  $\beta$ 

80	subunit of RNA polymerase (RpoB) <sup>7,8</sup> . Finally, resistance to linezolid has been related to		
81	the presence of the phenicol and lincosamide resistance gene (cfr), as described for		
82	staphylococci <sup>9</sup> .		
83	Although C. difficile is an important nosocomial pathogen, little is known about the		
84	epidemiology of this microorganism in Mexico. The aims of the present study were to		
85	determine the drug susceptibility of Mexican C. difficile isolates, particularly to the recently		
86	licensed CDI treatment, fidaxomicin, and to study circulating ribotypes in two tertiary-care		
87	hospitals in Mexico.		
88			
89	Methods		
90			
91	Settings and study population		
92	We designed an observational study of circulating C. difficile ribotypes, drug		
93	susceptibility, and drug resistance genes from two hospitals in Mexico: The Hospital Civil		
94	of Guadalajara "Fray Antonio Alcalde", is a 1000-bed tertiary-care teaching hospital, in		
95	Guadalajara; and the Hospital Universitario "Dr. José Eleuterio González", is a 450-bed		
96	tertiary-care teaching hospital in Monterrey.		
97	All patients with confirmed CDI from February 2011 through January 2016 were		
98	included in the study. Recurrences were defined as patients with a reappearance of		
99	symptoms after resolution of the previous diarrheal episode within 8 weeks or less. As		
100	patient information was anonymized and only microbiological data were analyzed,		

- informed consent was not required. The study was reviewed and approved by the Local
  Ethics Committee (Approval: 047/16).
- 103

# 104 Diagnosis of *Clostridium difficile* infection

- 105 Clinical diagnosis of CDI was suspected when patients were hospitalized for more
- than 48 h and had >3 loose stools in the previous 24 h. In the Hospital Civil of Guadalajara,
- 107 CDI was confirmed by real-time PCR using the Xpert<sup>®</sup> C. difficile/Epi assay (Cepheid,
- 108 Sunny Vale, CA, USA) and in the Hospital Universitario, the diagnosis was confirmed by
- 109 the use of the Meridian ImmunoCard ® Toxins A&B (Meridian Bioscience, Inc., Memphis,
- 110 TN, USA). Some patients were additionally diagnosed by the Xpert<sup>®</sup> C. *difficile/Epi* assay

111 in Hospital Universitario only at physicians' request.

112

#### 113 Culture

Fecal samples from all confirmed CDI cases were treated with absolute ethanol and cultured on *C. difficile* agar (Neogen Corporation, MI, USA) with cefoxitin (16 mg/L) in anaerobic conditions for up to 72 h. Plates were cultured in either an anaerobic jar or anaerobic chamber. Identification was performed by PCR as described <sup>10</sup>. Only one isolate per patient was included in the study.

119

# 120 Antimicrobial susceptibility testing

We tested antimicrobial agents used for CDI treatment such as vancomycin,
metronidazole (range from 0.03 mg/L to 128 mg/L), fidaxomicin (range from 0.002 to 8

123	mg/L), and rifampicin (range from 0.0001 mg/L to 128 mg/L); and also antimicrobials with
124	potential therapeutic use such as linezolid (range from 0.03 mg/L to 128 mg/L).
125	Furthermore, we tested antimicrobials that may be associated with induction of CDI,
126	including ciprofloxacin, moxifloxacin, erythromycin, clindamycin (range from $0.03 \text{ mg/L}$
127	to 128 mg/L), and tetracycline (range from 0.008 mg/L to 128 mg/L).
128	Susceptibility testing was performed by the agar dilution method using Wilkins-
129	Chalgren agar (Oxoid Limited, Basingstoke, Hampshire, England) and Schaedler's
130	anaerobe broth (Oxoid Limited) <sup>11</sup> . Briefly, overnight cultures in 5 ml of pre-reduced
131	Schaedler's broth were spotted onto plates of Wilkins-Chalgren agar with different
132	concentrations of antibiotics, using a multipoint inoculator (10 <sup>4</sup> colony-forming units/spot).
133	An agar plate without an antimicrobial agent was included as a growth control in both
134	aerobic and anaerobic atmosphere and the plates were read after 48 h of incubation at 37°C
135	in an anaerobic environment. C. difficile ATCC 700057 was used as quality control.
136	A stock solution of fidaxomicin (800 mg/L), was prepared in DMSO, then 1 ml of
137	stock was diluted in 5 ml of DMSO and 4 ml of 10% DMSO (final concentrations 80
138	mg/L); further dilutions were made in 10% DMSO. Stock solutions of remaining antibiotics
139	(2560 mg/L) and dilutions were dissolved accordingly to recommendations of the Clinical
140	and Laboratory Standards Institute (CLSI) document M100-S27.
141	

141

# 142 Mutations associated with drug resistance

To detect mutations associated with resistance to rifampicin or fidaxomicin, two
regions of the *rpoB* gene were amplified; for rifampicin, we used previously reported

- 145 primers <sup>9</sup>, and for fidaxomicin, we designed the primers CdrpoB-FD-F (5'-
- 146 TCATGGAAAATGGAACACCA-3') and CdrpoB-FD-R (5'-
- 147 CCAAACCTCCATCTCTCCAA-3'). We designed the primers CdrpoC-VAN-F (5'-
- 148 GAATGGGTGCTGAAGCTGTA-3) and CdrpoC-VAN-R (5'-
- 149 GACGGAAACGACCTTGCTTA -3 ') to amplify a region in the *rpoC* gene that has been
- linked to vancomycin resistance<sup>12</sup>. Furthermore, the presence of the *cfr* gene was
- investigated by PCR in selected strains as previously described  $^{13}$ .
- 152 Sequencing of PCR-purified products was performed by Macrogen Inc. (Seoul,
- 153 Korea). The sequences were analyzed using the NCBI Basic Local Alignment Search Tool
- 154 (BLAST).

155

#### 156 **Typing of isolates**

All isolates were typed for *tcdA*, *tcdB*, *cdtA*, *cdtB*, and for deletions in *tcdC* by PCR
as previously described <sup>14,15</sup>. For ribotyping, amplification of the 16S-23S rRNA intergenic
spacer region was conducted by PCR as described <sup>16</sup>. The ATCC strain BAA-1805
(ribotype 027) was used as a control. Selected isolates were ribotyped by capillary
electrophoresis at the *C. difficile* Ribotyping Network Reference Laboratory (CDRN) at
Leeds Teaching Hospitals Trust, Leeds, UK.

164 **Results** 

## 166 Culture

167	Samples were cultured in anaerobic jar (n=196, 57.1%) or in anaerobic chamber		
168	(n=147, 42.9%). In total, we cultured 343 samples, of which 124 (36.1%) yielded a positive		
169	C. difficile culture (one isolate per patient). Most of the cases were from the Hospital Civil		
170	of Guadalajara (n = 76, $61.3\%$ ); the Hospital Universitario accounted for 48 of the cases		
171	(38.7%).		
172			
173	Antimicrobial susceptibility profiles		
174	Four isolates (3.2%) had reduced susceptibility to fidaxomic (MIC = $2 \text{ mg/L}$ ),		
175	whereas no isolate was resistant to either tetracycline or metronidazole. $MIC_{50}$ and $MIC_{90}$		
176	were $\geq$ 128 mg/L to ciprofloxacin, erythromycin, clindamycin, and rifampicin (Table 1).		
177	MIC distributions of CDI treatment drugs (vancomycin, metronidazole and fidaxomicin)		
178	are shown in Figure 1.		
179			
180	Molecular analysis of drug resistance		

181 When analyzing a region of *rpo*B in fidaxomicin-susceptible isolates (n= 18) and 182 isolates with reduced susceptibility (n=3), we found 7 mutations; one of these caused an 183 amino acid change that was not associated to reduced susceptibility strains (Table 2). 184 Furthermore, the *rpoB* gene was amplified and partially sequenced in 14 rifampicin-185 susceptible and 22 rifampicin-resistant isolates. We detected 3 mutations that generated 186 amino acid changes in both susceptible and resistant isolates; the Arg505Lys mutation was

187	more frequently detected in resistant isolates (21/22, 95.4%) than in susceptible isolates
188	(4/14, 28.5%) (OR = 52.5; 95% CI = 5.17- 532.6; p<0.000) (Table 2).

189	We also analyzed the $rpoC$ gene in 14 vancomycin-susceptible isolates and 12
190	isolates with reduced susceptibility to vancomycin and detected 22 mutations; two of them
191	were associated with an amino acid change (Table 2). The presence of drug resistance
192	genes was analysed in selected isolates (i.e. isolates with the highest and lowest MIC
193	values). The <i>cfr</i> gene was not amplified in any of the linezolid-susceptible (n=17) or
194	linezolid-resistant (n=9) C. difficile isolates evaluated.

195

#### 196 **Ribotypes**

Toxin A and toxin B genes were detected in all isolates. The binary toxin gene was
detected in 89 isolates (71.8%), of which 87 (97.8%) contained the *tcdC* 18-bp deletion
(Table 3).

Eighty-four isolates (67.7%) demonstrated the same ribotype banding patterns to the control strain BAA-1805 (ribotype 027) (Table 2) and we randomly selected twelve strains that were all confirmed to be ribotype 027 by the CDRN (Leeds, UK). Similarly, of 18 isolates (14.5%) that demonstrated similar banding pattern to ribotype 001 (90% of similarity), four randomly selected isolates were confirmed to be ribotype 001 by the CDRN (Leeds, UK). The ribotypes and presence of toxin genes of the other isolates are summarized in table 3.

# **Discussion**

209	In our study, ribotype 027 was the predominant strain, accounting for 67.7% of the
210	cases; this ribotype is considered epidemic and has been reported worldwide <sup>17</sup> . In previous
211	publications we have found 027 strain as the predominant ribotype in our settings <sup>18,19</sup> , the
212	latter is in contrast to diverse studies were there is a high diversity of ribotypes and 027
213	stains account for less than $30\%$ <sup>20-22</sup> . The second most frequent ribotype was 001,
214	accounting for 14.5% of the cases; this ribotype is the main ribotype circulating in Korea <sup>23</sup> ,
215	Czech Republic <sup>2</sup> , Croatia <sup>24</sup> , and Slovakia <sup>25</sup> , however, this is the first report on ribotype
216	001 circulation in Mexico. This strain has been associated with high drug resistance,
217	including resistance to ciprofloxacin, erythromycin, and clindamycin <sup>24</sup> .
218	Fidaxomicin is an FDA-approved antibiotic for the treatment of CDI <sup>8</sup> . A previous
219	study that included 1,323 isolates showed a MIC <sub>90</sub> of 0.5 mg/L against <i>C. difficile</i> <sup>26</sup> ;
220	similarly, Snydman et al reported 925 isolates that were inhibited at a fidaxomicin
221	concentration $\leq 1$ mg/L, and the MIC <sub>90</sub> was 0.5 mg/L $^{20}$ . In the present study, we observed a
222	MIC <sub>90</sub> of 0.06 mg/L; however, we detected 4 ribotype 027 isolates with a fidaxomicin MIC
223	of 2 mg/L; these isolates were recovered from patients with recurrent CDI. This finding is
224	of interest since fidaxomicin is unavailable in Mexican hospitals and none of the patients
225	included in the study were exposed to this drug. However, we performed susceptibility
226	testing only once; nevertheless, the MIC of control strain were reproducible and the four
227	strains with MIC = $2 \text{ mg/L}$ were detected in a batch of 36 isolates being tested at the same
228	time. MICs of the remaining 32 isolates were $\leq 0.125$ mg/L.

229	Reduced susceptibility to fidaxomicin may be due to point mutations in the <i>rpoB</i>
230	gene (RNA polymerase subunit $\beta$ ) <sup>12</sup> . We detected one point mutation (Glu1036Gln) in
231	RpoB, but it was not associated with drug resistance ( $P = 0.489$ ). Leeds <i>et al.</i> identified two
232	mutations in <i>rpoB</i> that were associated with reduced susceptibility to fidaxomicin; one of
233	them coded a Gln1073Arg substitution and the second was a frameshift after amino acid
234	117 of a homolog of the MarR family of transcriptional regulators <sup>12</sup> ; however, we did not
235	detect these mutations in our strains. Other mutations associated with fidaxomicin reduced
236	susceptibility have been reported, but all of them have been obtained through the serial
237	passage of strains into media containing fidaxomicin <sup>27</sup> . Goldstein <i>et al.</i> isolated a strain
238	with a MIC of 16 mg/L to fidaxomicin from a patient with an episode of recurrence during
239	a clinical trial <sup>28</sup> ; this isolate harbored a Val1143Gly substitution in <i>rpoB</i> . However, the
240	authors did not consider that resistance had developed during the clinical trial and did not
241	explain the clinical relevance of this finding. To our knowledge, there is no report of
242	clinical resistance to fidaxomicin. It is widely known that fidaxomicin reaches high levels
243	in the gut (1,000 $\mu$ g/g of faeces) <sup>28</sup> ; thus, the actual implication of the high MIC in the
244	strains is unknown, considering the lack of reports on clinical resistance and particularly in
245	Mexico, where this drug is not available.

In our study, we detected reduced susceptibility to vancomycin of 40.3% with 48.8% of reduced susceptibility in 027 strains and 33.3% in 001 strains; proportions as high as 87.7% have been reported in ribotype 027 strains <sup>29</sup>. Similarly, to fidaxomicin, it is unlikely that reduced susceptibility impacts on clinical response, due to the high levels of vancomycin reached in the gut (>2000 mg/L); however, among patients from whom we recovered isolates with reduced susceptibility (50 patients), nine died because of CDI and

five of these patients received vancomycin. Nevertheless, other factors may have acted inthese patients' response to treatment.

254 Although this bacterial species has a *vanG* homolog inducible by vancomycin, it does not promote vancomycin resistance <sup>30</sup>. Leeds *et al.* identified an Asp244Tyr 255 substitution in rpoC that was associated with reduced susceptibility to vancomycin<sup>12</sup>. 256 257 Despite our efforts to detect this mutation, it was not found. Leeds et al. reported additional mutations: a Pro108Leu substitution in a transferase encoded by murG/CD2725, a stop 258 codon after amino acid 326 in an exonuclease encoded by CD3659, and a single amino acid 259 deletion in an L-serine dehydrogenase (*sdaB*). Therefore, it seems that diverse mechanisms 260 261 are responsible for reduced susceptibility to vancomycin, particularly those involved in cell 262 wall biosynthesis.

On the other hand, we observed no resistance to metronidazole; similar findings have been reported in other studies  $^{21,31}$ . Clinical failures with metronidazole treatments have been attributed to the development of heteroresistance and deficiencies in the pharmacokinetics of the drug resulting in low luminal concentrations following oral administration. Resistance to metronidazole is known to be unstable, with the loss of levels of resistance due to laboratory manipulation <sup>5</sup>.

High MICs to linezolid has occasionally been described in C. difficile  $^{9,32}$ .

270 Interestingly, the isolates of ribotype 001 showed higher MICs (8-32 mg/L) than 027

isolates (8 mg/L). Although linezolid is not used for the treatment of CDI, linezolid is

widely used in the Hospital Civil of Guadalajara for the treatment of nosocomial

273 pneumonia, surgical wound infections, and bloodstream infections not associated with a

274	catheter. Marín <i>et al.</i> found nine isolates resistant to linezolid and were <i>cfr</i> -positive <i>C</i> .
275	difficile isolates that belonged to the same clonal cluster, suggesting possible horizontal
276	transmission of these strains among patients in their hospital setting <sup>9</sup> . However, we did not
277	detect the cfr gene in any of the selected C. difficile isolates we studied.
278	We also found a high proportion of isolates with elevated MICs to rifampicin in 027
279	strains (95.2%) and 001 strains (83.3%). In contrast, Tenover et al found lower proportions
280	(27.5%) of isolates with high MIC to rifampicin <sup>21</sup> . For this antimicrobial agent, a bimodal
281	distribution of MICs has been reported; Norén et al. found 80% of strains to with low MIC
282	(>0.016 mg/L) or high MIC (>256 mg/L) $^{33}$ . In our isolates, we detected three previously
283	reported amino acid substitutions in RpoB associated with rifampicin resistance:
284	Arg505Lys, His502Asn, and Ile548Met. Curry et al <sup>13</sup> reported all three changes, including
285	Arg505Lys, which was present in isolates with MICs $>32$ mg/L. We also confirmed the
286	importance of this mutation in rifampicin-resistant isolates (OR = 52.5, CI 5.17- 532.6, $p$ =
287	0.000). Similarly, the authors reported an Ile548Met change in isolates with MICs $>32$
288	mg/L, however, in the C. difficile strains evaluated in the present study, this change was not
289	associated with rifampicin resistance ( $p = 0.074$ ).

Our study has some limitations. First, we were unable to recover all isolates from all samples, in fact, the recovery rate was low. The low recuperation can be attributed to the medium used, which does not incorporate sodium taurocholate as spore germinant; and the use of ethanol to eliminate any vegetative organisms that survived freezing. Consequently, the isolates are not homogeneously distributed throughout the study period, making it difficult to study distribution over time, and perhaps generating bias on ribotype prevalence; second, clinical diagnosis was not confirmed in a uniform way. Apart from

297	differences in diagnosis, this may have contributed to the low recovery of isolates; and
298	finally, data of ribotyping in 027 and 001 strains were mainly extrapolated from primary
299	results of conventional electrophoresis.
300	In conclusion, this is the first report on drug susceptibility of C. difficile ribotypes
301	circulating in Mexico. Ribotypes 027 and 001 were the most frequent and highly drug
302	resistant; furthermore, we found four isolates with reduced susceptibility to fidaxomicin,
303	raising a concern since this drug is unavailable in Mexican Hospitals. The clinical relevance
304	of these findings needs to be addressed to fully understand the epidemiology of CDI in
305	Mexican hospitals.
306	

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309

# 310 Author Disclosure Statement

311 No competing financial interests exist.

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