Response to Rodriguez-Cavallini's et al. letter to the editor

Why were molecular studies not considered to classify *Clostridium difficile* isolates implicated in an outbreak?

## To the Editor:

We are pleased that Rodríguez-Cavallini and colleagues have recognized our paper, "Clostridium difficile outbreak in Costa Rica: control actions and associated factors," for broadening the literature on C. difficile associated disease (CDAD) epidemiology and prevention in Latin America (1).

As to the contradiction mentioned with regards to the study results, it is important to consider that a diverse array of antibiotics can be associated with CDAD. Our study results are in agreement with the Clinical Practice Guidelines for Clostridium difficile Infection in Adults: 2010 Update (2), which includes restriction of antimicrobial use as cephalosporins, with high recommendation strength based on quality of evidence.

For our study, samples were collected from all affected patients rather than from a subset of cases during the outbreak. When comparing our results with those cited in existing literature in Latin America and in Costa Rica, it is important to consider that the local epidemiological profile of *C. difficile* at other centers may vary considerably from that of the hospital in this study. Also, it should be noted that the main objective of the study was to describe the infection control strategies implemented during the outbreak and to determine factors associated with *C. difficile* infection. The determination of specific molecular features of *C. difficile* was not a main focus of the investigation.

Finally, we commend the invaluable effort of the Laboratorio de Investigación en Bacteriología Anaerobia (LIBA) of the University of Costa Rica in contributing to the existing knowledge of autochthonous *C. difficile* pulsotypes. Given the inherent diversity of Costa Rican health centers, we feel it is extremely important to incorporate the work of LIBA into the country's routine surveillance of *C. difficile* to increase knowledge regarding this critical pathogen.

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