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AUTHOR'S CORRECTION

tcdC Genotypes Associated with Severe TcdC Truncation in an Epidemic Clone and Other Strains of *Clostridium difficile*

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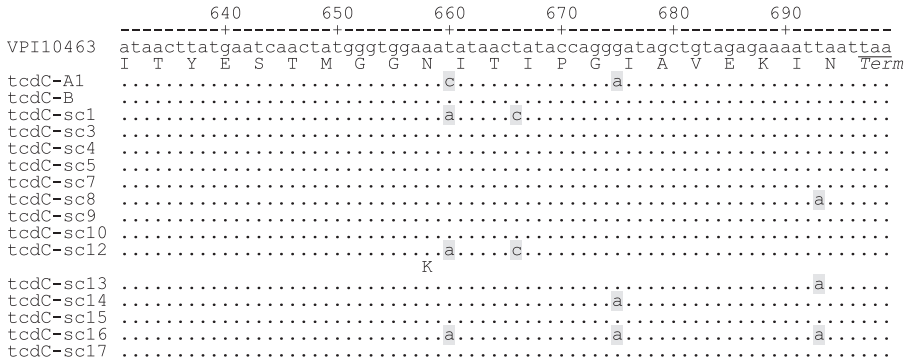
Volume 45, no. 1, p. 215–221, 2007. In our paper, we originally described 14 novel *tcdC* genotypes in *C. difficile* using primers published by P. Spigaglia and P. Mastroantonio (J. Clin. Microbiol. **40:**3470–3475, 2002). Subsequently, we discovered sequencing errors at the 3′ end of *tcdC* in five of these genotypes due to the location of primer C1, which overlaps the last 26 nucleotides of *tcdC*. Sequence analysis with new primers that reside outside *tcdC* identified nucleotide changes in *tcdC-sc2*, *tcdC-sc8*, *tcdC-sc13*, *tcdC-sc14*, and *tcdC-sc16*. Additionally, all three isolates bearing the previously published *tcdC-A* genotype from our collection consistently generated a distinct genotype when sequenced with these primers. Therefore, we are designating this allele *tcdC-A1* (GenBank accession number EF470292). The nucleotide changes described either are untranslated (*tcdC-A1*, *tcdC-sc2*, *tcdC-sc8*, and *tcdC-sc16*) or result in silent mutations (*tcdC-sc13* and *tcdC-sc14*). The GenBank entries for our novel genotypes (GenBank accession numbers DQ861426, DQ861417, DQ861421, DQ861422, and DQ861424) have been corrected accordingly. These revisions illustrate 15 new *tcdC* genotypes.

Page 216: The legend to Fig. 1 should read as follows. "Comparison of TcdC nucleotide and amino acid sequences to the published sequence for the reference strain VPI10463. Dots indicate identical bases, dashes indicate deletions, and shaded letters correspond to nucleotide changes. Only amino acid changes are depicted. Genotype *tcdC-sc2* is not depicted but is identical to *tcdC-sc1* with the exception of an inserted nucleotide (t) between positions 212 and 213 and a nucleotide (t) at position 660."

Page 217: The bottom panel of Fig. 1 should appear as follows.



Page 219, Fig. 2: In the portion of the figure branching up from genotype 17 near the top, the genotype given as "0, B" should be "B." This correction strengthens our original conclusion that *tcdC* genotypes are concordant with MLVA genotypes.

Page 220, column 1: Lines 18 to 22 should read ". . . together by MST of MLVA genotypes. All isolates with identical MLVA genotypes contained identical *tcdC* genotypes."

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