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ERRATUM

Erratum to: In-depth Characterization via Complementing Culture-Independent Approaches of the Microbial Community in an Acidic Hot Spring of the Colombian Andes

Laura C. Bohorquez · Luisa Delgado-Serrano · Gina López · César Osorio-Forero · Vanja Klepac-Ceraj · Roberto Kolter · Howard Junca · Sandra Baena · María Mercedes Zambrano

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The original version of this article unfortunately contained a mistake found under the Materials and Methods section for the PCR protocol under section Pyrosequencing of V5–V6 Hypervariable Regions", what says:

"PCR amplifications were done as reported at the time [29] in a 25-μL reaction volume containing 2 μl (20 ng) DNA, 0.75 μM of each primer 807 F and 1050R designed by us (Table 1), 2.5 U Pfu Turbo® DNA polymerase (Stratagene, Inc., La Jolla, CA, USA), 1× Pfu reaction buffer, 0.6 mM dNTPs, 5% *v/v* dimethyl sulfoxide using the following PCR conditions: 2 min at 95°C, 30 cycles consisting of denaturation for 30 s at 95°C, a temperature touch down from 60 to 51°C (2°C every six cycles), 72°C for 1 min and a final extension of 72°C for 5 min."

The correct wording should be:

"PCR amplifications were done as reported at the time [29] in a 25-μL reaction volume containing 2 μl (20 ng) DNA, 0.75 μM of each primer 807 F and 1050R designed by us (Table 1), 2.5 U Pfu Turbo® DNA polymerase (Stratagene, Inc., La Jolla, CA, USA), 1× Pfu reaction buffer, 0.6 mM dNTPs, 5% *v/v* dimethyl sulfoxide using the following PCR conditions: 2 min at 95°C, 30 cycles consisting of denaturation for 30 s at 95°C, annealing temperature starting at 60°C and reducing 0.2°C at every cycle, elongation at 72°C for 1 min and a final extension of 72°C for 5 min."

The online version of the original article can be found at http://dx.doi.org/10.1007/s00248-011-9943-3.

L. C. Bohorquez · L. Delgado-Serrano · C. Osorio-Forero · H. Junca · M. M. Zambrano (✉)
Molecular Genetics & Biotechnology, Corporación CorpoGen,
Carrera 5 No. 66A-34,
110231, Bogotá, D.C., Colombia
e-mail: mzambrano@corpogen.org

L. C. Bohorquez · L. Delgado-Serrano · G. López · C. Osorio-Forero · H. Junca · S. Baena · M. M. Zambrano
Colombian Center for Genomics and Bioinformatics of Extreme Environments – GeBiX,
Carrera 5 No. 66A-34,
110231, Bogotá, DC, Colombia

G. López · S. Baena
Unidad de Saneamiento y Biotecnología Ambiental, Departamento de Biología, Pontificia Universidad Javeriana,
POB 56710, Bogotá, DC, Colombia

V. Klepac-Ceraj · R. Kolter
Harvard Medical School,
200 Longwood Avenue,
Boston, MA 02115, USA

Present Address:
V. Klepac-Ceraj
The Forsyth Institute,
245 First Street,
Cambridge, MA 02142, USA