Errata

**Catenulispora rubra** sp. nov., an acidophilic actinomycete isolated from forest soil

Tomohiko Tamura, Yuumi Ishida, Takeshi Sakane and Ken-ichiro Suzuki


On pages 2272 and 2273, the whole-cell sugars are given incorrectly as ribose, mannose, arabinose and glucose. The correct whole-cell sugars should be rhamnose, mannose, arabinose and galactose. In the description on page 2273, the feature of the spore chain is given incorrectly as a hook-like or flexuous manner. The correct feature of the spore chain should be a rectiflexible or straight.

**Mucilaginibacter paludis** gen. nov., sp. nov. and **Mucilaginibacter gracilis** sp. nov., pectin-, xylan- and laminarin-degrading members of the family *Sphingobacteriaceae* from acidic *Sphagnum* peat bog

Timofei A. Pankratov, Brian J. Tindall, Werner Liesack and Svetlana N. Dedysh

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The GenBank/EMBL/DDBJ accession number for the 16S rRNA gene sequence of *Mucilaginibacter gracilis* TPT18<sup>T</sup> is AM490403 (not AM490402, as stated in the paper) and that for the 16S rRNA gene sequence of *Mucilaginibacter paludis* TPT56<sup>T</sup> is AM490402 (not AM490403).

**Description of Pelomonas aquatica** sp. nov. and **Pelomonas puraquae** sp. nov., isolated from industrial and haemodialysis water

Margarita Gomila, Botho Bowien, Enevold Falsen, Edward R. B. Moore and Jorge Lalucat


The GenBank/EMBL/DDBJ accession number for the 16S rRNA gene sequence of CCUG 52575<sup>T</sup> is AM501435 (not AM51435, as stated in the paper).