Errata

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

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International Journal of Systematic and Evolutionary Microbiology (2007), 57, part 10, 2272–2274

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

International Journal of Systematic and Evolutionary Microbiology (2007), 57, part 10, 2349–2354

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

International Journal of Systematic and Evolutionary Microbiology (2007), 57, part 11, 2629–2635

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