Errata

_Catenulispora rubra_ sp. nov., an acidophilic actinomycete isolated from forest soil

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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

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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

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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).