Erratum

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***Brevibacterium limosum* sp. nov., *Brevibacterium pigmenatum* sp. nov., and *Brevibacterium atlanticum* sp. nov., three novel dye decolorizing actinobacteria isolated from ocean sediments**

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In the article by Pei *et al.* published in Journal of Microbiology 2021; 59, 898–910, the name *Brevibacterium pigmenatum* should be corrected to *Brevibacterium pigmentatum* in the title of the paper, abstract, keywords, introduction, protologue, Figures, and Supplementary data in the article.

The title for this article should have read:

Brevibacterium limosum sp. nov., *Brevibacterium pigmentatum* sp. nov., and *Brevibacterium atlanticum* sp. nov., three novel dye decolorizing actinobacteria isolated from ocean sediments

The keywords should have read:

Keywords: *Brevibacterium limosum* sp. nov., *Brevibacterium pigmentatum* sp. nov., *Brevibacterium atlanticum* sp. nov., genomic analysis, dye decolorization

The last sentence of Abstract for this article should have read:

Based on the phylogenetic, genotypic and phenotypic data, strains o2^T, YB235^T and WO024^T clearly represent three novel taxa within the genus *Brevibacterium*, for which the names *Brevibacterium limosum* sp. nov. (type strain o2^T = JCM 33844^T = MCCC 1A09961^T), *Brevibacterium pigmentatum* sp. nov. (type strain YB235^T = JCM 33843^T = MCCC 1A09842^T) and *Brevibacterium atlanticum* sp. nov. (type strain WO024^T = JCM 33846^T = MCCC 1A16743^T) are proposed.

The last sentence of Introduction should have read:

Through polyphasic taxonomic analysis, strains YB235^T, WO024^T, and o2^T were identified as three novel dye decolorizing bacterial species, for which the names *Brevibacterium pigmentatum*, *Brevibacterium atlanticum*, and *Brevibacterium limosum* are proposed.

The protologue should have read:

Description of *Brevibacterium pigmentatum* sp. nov.

Brevibacterium pigmentatum (pig.men.ta'tum. L. neut. adj. *pigmentatum* pigmented).

Cells are Gram-stain positive, aerobic, non-spore-forming, non-motile and long rod-shaped (approximately 0.2–0.5 µm × 1.5–4.0 µm in size) after incubation for 3 days at 30°C on MA. Colonies are light orange color, opacity, smooth, moist, low convex with middle margins. Growth well on MA, LB and TSA plates, and moderate growth on ISP 4 and ISP 2 plates. Diffusible pigment is produced at MA medium. Growth is observed at 4–45°C, pH 6.0–10.0 and in the presence of 0–20% (w/v) NaCl. Optimum growth occurs at 30–37°C, pH 7.0, and in 3–5% (w/v) NaCl. Oxidase- and catalase- positive. Positive for gelatin hydrolysis and the activities of pyrazinamidase and alkaline phosphatase; negative for nitrate reduction, esculin hydrolysis and the activities of urease, pyrrolidone peptidase, β-glucuronidase, β-galactosidase, α-glucosidase and *N*-acetyl-β-glucosaminidase. Acid is produced from D-glucose, D-ribose, D-xylose, D-malic acid and D-lactose, But not from D-mannitol, D-lactose and D-saccharose. The principal fatty acids were anteiso-C_{17:0} and anteiso-C_{15:0}. The major polar lipids were diphosphatidylglycerol, phosphatidylglycerol and one unknown glycolipid. The diagnostic diamino acid in the cell-wall peptidoglycan is *meso*-diaminopimelic acid. The predominant menaquinone was determined to be menaquinone MK-8(H₂). The DNA G + C base composition is 64.6 mol% and the genome size is 3.91 Mb. The type strain, YB235^T (= JCM 33843^T = MCCC 1A09842^T) was isolated from an ocean sediment of the South Atlantic.

The Figures should have shown:

Fig. 1.

Fig. 3.

The Supplementary data should have shown as the enclosed file.

The authors apologize for any inconvenience that this may have caused.