CORRECTION

Correction to: *Duganella rivus* sp. nov., *Duganella fentianensis* sp. nov., *Duganella qianjiadongensis* sp. nov. and *Massilia guangdongensis* sp. nov., isolated from subtropical streams in China and reclassification of all species within genus *Pseudoduganella*

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Correction to: Antonie van Leeuwenhoek (2020) 113:1155–1165
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In the published version of the article, the title should have read ‘*Duganella rivi* sp. nov., *Duganella fentianensis* sp. nov., *Duganella qianjiadongensis* sp. nov. and *Massilia guangdongensis* sp. nov., isolated from subtropical streams in China and reclassification of all species within genus *Pseudoduganella*’.

A newly proposed name in the paper, *Duganella rivus*, was incorrectly formed. The name *Duganella*

rivus was changed to *Duganella rivi*, with the following description:

Description of *Duganella rivi* sp. nov.

Duganella rivi (ri’vi. L. gen. n. *rivi* of a rivulet).

Cells are Gram-stain-negative, facultative anaerobic, rod-shaped, motile with several flagella, 2.4–3.1 µm in length and 0.9–1.1 µm in width. The colonies are pale and convex. Growth occurs at 4–34 °C (optimum 24 °C), pH 4.0–10.5 (optimum 7.0–7.5) and NaCl 0–0.5% w/v (optimum 0%). It is positive for oxidase, catalase and urease, but negative for nitrate reduction, H₂S production, phenylalanine deaminase, Voges-Proskauer test, methyl red test and indole production. It hydrolyses Tweens 20 and 40, aesculin and starch, but not Tweens 60 or 80, casein or gelatin. It utilises maltose monohydrate, D-lactose monohydrate, D-glucose, D-fructose, sucrose, L-arabinose and starch, but not D-trehalose dihydrate, glycerol, mannitol, D-sorbitol, L-glutamic acid, L-valine, glycine, L-serine, L-aspartic acid, L-proline, sodium pyruvate, sodium oxalate, sodium formate dihydrate, sodium acetate anhydrous or disodium fumarate. It is negative for esterase (C4), esterase lipase (C8), lipase (C14), cystine arylamidase, trypsin, α-chymotrypsin, α-galactosidase, β-glucuronidase, β-glucosidase, α-mannosidase and α-fucosidase, positive for alkaline phosphatase, leucine arylamidase, valine arylamidase,

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acid phosphatase, naphthol-AS-BI-phosphohydrolase, β -galactosidase, α -glucosidase and N-acetyl- β -glucosaminidase. The major fatty acids are C_{16:1} ω 7c, C_{16:0}, C_{18:1} ω 7c, C_{12:0} and C_{10:0} 3OH. The polar lipids include phosphatidylethanolamine, phosphatidylglycerol and one unidentified phospholipid. The G + C content of genomic DNA is 61.7%.

Type strain FT55W^T (= GDMCC 1.1675^T = KACC 21467^T) was isolated from a subtropical stream in China. The GenBank/EMBL/DDBJ accession numbers for the 16S rRNA gene sequence and the

draft genome of strain FT55W^T are MN865823 and WWCK00000000, respectively.

The authors apologize for any inconvenience caused.

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