

Erratum to: *Pontibacter locisalis* Sy30T sp. nov. isolated from soil collected from an abandoned saltern

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Published online: 1 July 2016
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**Erratum to: Antonie van Leeuwenhoek (2016)
109:415–420
DOI 10.1007/s10482-016-0646-0**

Subsequent to the publication of the above paper, it has been noticed that some details need correcting, including the title and one of the type strain culture collection codes.

The correct title should read:

Pontibacter locisalis sp. nov. isolated from soil collected from an abandoned saltern

The last line of the Abstract should read:

The type strain is Sy30^T (=KCTC 42498^T=CCTCC AB 2015060^T).

The corrected protologue is given below:

The online version of the original article can be found under doi:[10.1007/s10482-016-0646-0](https://doi.org/10.1007/s10482-016-0646-0).

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Description of *Pontibacter locisalis* sp. nov.

Pontibacter locisalis (lo.ci.sa'lis. L. n. *locus* place, locality; L. gen. n. *salis* of salt; N.L. gen. n. *locisalis* from a place of salt)

Cells are Gram-stain negative, non-motile, aerobic and rod-shaped with the size of 0.4–0.6 μm in width and 2.0–7.0 μm in length. Colonies are red, non-luminescent, and circular with entire edges on 2216E agar. Flexirubin-type pigments are absent. Growth occurs at 15–40 °C (optimum, 30–33 °C), pH 6.5–9.0 (optimum, 7.0–8.0) and in presence 0.5–5.0 % of NaCl (optimum, 2.0–2.5 %). Requires NaCl for growth. Can hydrolyze gelatin, aesculin, alginate and starch but not casein, cellulose and agar. Nitrate reduction, lysine decarboxylase, ornithine decarboxylase, H₂S production, citrate utilization, Voges–Proskauer reaction, indole production, glucose acidification, arginine dihydrolase and urease are negative. Positive for catalase, alkaline phosphatase, esterase (C4), esterase lipase (C8), leucine arylamidase, valine arylamidase, cystine arylamidase, trypsin, acid phosphatase, naphthol-AS-BI-phosphohydrolase and N-acetyl-β-glucosaminidase but negative for oxidase, lipase (C14), α-chymotrypsin, α-galactosidase, β-galactosidase, β-glucuronidase, α-glucosidase, β-glucosidase, α-mannosidase and α-fucosidase. Assimilates sucrose, D-lactose, D-glucose, D-galactose, D-fucose, L-rhamnose, D-melibiose, D-arabitol, D-trehalose but not D-mannose, L-alanine and sodium lactate. The predominant fatty acids are iso-C_{15:0}, and summed

feature 4 (iso-C_{17:1} I and/or anteiso-C_{17:1} B). The major menaquinone is MK-7. The major polyamine is *sym*-homospermidine. The DNA G+C content of the type strain is 47.7 mol %. The polar lipids comprise major amounts of phosphatidylethanolamine, a phosphoaminolipid and two unidentified polar lipids, and minor amounts of two unidentified polar lipids and two phospholipids.

The type strain Sy30^T (=KCTC 42498^T = CCTCC AB 2015060^T), was isolated from dry soil of an abandoned marine solar saltern at Weihai, China. The GenBank accession number for the 16S rRNA gene sequence of strain Sy30^T is KR080555.