

ERRATUM

Erratum to: *Haematospirillum jordaniae* gen. nov., sp. nov., isolated from human blood samples

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The author has noticed that some minor details including type strain culture collection codes were missing in the original publication of the article. The corrected protologue is given below.

Description of *Haematospirillum* gen. nov.

Haematospirillum (Hae.ma.to.spi.ril' lum. Gr. n. *haima* -atos, blood; L. n. *spira*, a spiral; N.L. neut.

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dim. n. *spirillum*, a small spiral; N.L. dim. neut. n. *Haematospirillum*, a small spiral from blood).

Cells are Gram-negative, oxidase positive, and catalase positive. Cells are motile, helical in shape and exhibit bipolar flagella. Major fatty acids are C_{16:0}, C_{16:1}ω7c and C_{18:1}ω7c. The predominant ubiquinone is Q-10. The major polar lipids are phosphatidylethanolamine, phosphatidylglycerol and aminolipids. The DNA G+C content of the type strain of the type species is 49.9 %. A member of the order *Rhodospirillales*, family *Rhodospirillaceae*. The type species is *Haematospirillum jordaniae*.

Description of *Haematospirillum jordaniae* sp. nov.

Haematospirillum jordaniae (jor.da'ni.ae. N.L. gen. n. *jordaniae*, named in honour of the late Jean Jordan).

In addition to those given in the genus description, the organism exhibits the following properties. On Heart Infusion Agar supplemented with 5 % rabbit blood, colonies are circular, low convex, entire, grey, and 1 mm in diameter after 48 h of incubation at 35 °C. Cells are approximately 1.6 μm in length and 0.1–0.25 μm in width. Positive for hydrogen sulfide production. Negative for utilisation of urea, nitrate reduction and tryptophan conversion to indole. In the API ZYM test, positive for alkaline phosphatase, esterase (C4), esterase lipase (C8) and acid phosphatase. Negative for lipase (C14), valine arylamidase, cystine arylamidase, trypsin, α-chymotrypsin,

Naphthol-AS-BI-phosphohydrolase, α -galactosidase, β -galactosidase, β -glucuronidase, α -glucosidase, β -glucosidase, *N*-acetyl- β -glucosaminidase, α -mannosidase, and α -fucosidase.

The type strain, H5569^T (=DSM^T 28903 = CCUG 66838^T), was isolated from a human blood sample. The 16S rRNA sequence for strain H5569^T has been deposited in GenBank under Accession Number KM083603.