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### *Corrigendum to*

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Corrigendum to: "Comparative genomics and pan-genomics of the Myxococcaceae, including a description of five novel species: *Myxococcus eversor* sp. nov., *Myxococcus*

*llanfairpwllgwyngyllgogerychwyrndrobwlllantysiliogogochensis* sp. nov., *Myxococcus vastator* sp. nov.,

*Pyxidicoccus caerfyrddinensis* sp. nov. and *Pyxidicoccus trucidator* sp. nov." [Genome Biol. Evol. 12(12) (2020) 2289-2302] . *Genome Biology and Evolution*, 13(4). <https://doi.org/10.1093/gbe/evab040>

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## Corrigendum

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**Corrigendum to “Comparative genomics and pan-genomics of the Myxococcaceae, including a description of five novel species: *Myxococcus eversor* sp. nov., *Myxococcus llanfairpwllgwyngyllgogerychwyrndrobwlllantysiliogogochensis* sp. nov., *Myxococcus vastator* sp. nov., *Pyxidicoccus caerfyrdinensis* sp. nov. and *Pyxidicoccus trucidator* sp. nov.” [Genome Biol. Evol. 12(12) (2020) 2289-2302]**

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In the species descriptions of the species *Myxococcus llanfairpwllgwyngyllgogerychwyrndrobwlllantysiliogogochensis* sp. nov., *Myxococcus vastator* sp. nov., and *Pyxidicoccus caerfyrdinensis* sp. nov. the acronyms of the culture collections to which the type strains were deposited contained typographic errors. The corrected species descriptions are provided below:

### SPECIES DESCRIPTIONS

*Myxococcus llanfairpwllgwyngyllgogerychwyrndrobwlllantysiliogogochensis* sp. nov.

*Myxococcus llanfairpwllgwyngyllgogerychwyrndrobwlllantysiliogogochensis*, (llan.fair.pwll.gwyn.gyll.gog.er.ych.wyrn.-dro.bwllll.ant.ysil.iog.ogogoch.en’sis. N.L. masc. adj. llanfairpwllgwyngyllgogerychwyrndrobwlllantysiliogogochensis, pertaining to llanfairpwllgwyngyllgogerychwyrndrobwlllantysiliogogoch, reflecting its isolation from soil collected in that parish [gridref 53.22°N 4.19°W]).

Vegetative cells are Gram-negative bacilli tapering slightly at the ends, measuring 0.4-0.6 µm x 4.0-7.0 µm in electron micrographs. Colonies exhibit swarming motility and appear pale brown on VY-2 agar (w/v 0.5% dried baker’s yeast, 0.1% CaCl<sub>2</sub>·2H<sub>2</sub>O, 1.5% agar). Fruiting bodies are irregular spheroids, orange in colour. Aerobic growth was observed at 30 °C and 35 °C, and at pH 5.0-9.0. Growth was unaffected by the addition of 1-4% NaCl. Hydrolyses esculin, gelatine, p-nitrophenyl-β-D galactopyranoside and urea. Assimilates N-acetyl-glucosamine, adipate, arabinose, glucose, malate, maltose, mannitol, mannose and phenyl acetate. Cells prey with low efficiency upon *Escherichia coli* TOP10, *Clavibacter nebraskensis* DSM 7483 and *Ustilago maydis*. DNA GC content is 68.7 mol%. The draft genome sequence of AM401<sup>T</sup> is available from GenBank (Accession VIFM00000000). The type strain (AM401<sup>T</sup> = NBRC 114351<sup>T</sup> = NCCB 100770<sup>T</sup>) was isolated from soil collected in the parish of Llanfairpwllgwyngyllgogerychwyrndrobwlllantysiliogogoch, United Kingdom [gridref 53.22°N 4.19°W].

*Myxococcus vastator* sp. nov.

*Myxococcus vastator* (vas.ta’tor L. masc. n. *vastator* the ravager, after its ability to devastate colonies of prey cells).

Vegetative cells are Gram-negative bacilli tapering slightly at the ends, measuring 0.6-0.7 µm x 3.0-6.0 µm in electron micrographs. Colonies exhibit swarming motility and appear pale brown on VY-2 agar (w/v 0.5 % dried baker’s yeast, 0.1% CaCl<sub>2</sub>·2H<sub>2</sub>O, 1.5 % agar). Fruiting bodies are irregular spheroids, orange in colour. Aerobic growth was observed at 30 °C,

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and at pH 8.0-9.0. Growth was unaffected by the addition of 1 % NaCl. Hydrolyses arginine and urea. Assimilates malate. Cells prey with low efficiency upon *Escherichia coli* TOP10, *Clavibacter nebraskensis* DSM 7483 and *Ustilago maydis* DSM 14603.

DNA GC content is 69.9 mol%. The draft genome sequence of AM301<sup>T</sup> is available from GenBank (Accession JAAIYB000000000). The type strain (AM301<sup>T</sup> = NCCB 100768<sup>T</sup> = NBRC 114352<sup>T</sup>) was isolated from soil collected in the parish of Llanfairpwllgwyngyllgogerychwyrndrobwllllantysiliogogoch, United Kingdom [gridref 53.22°N 4.19°W].

*Pyxidicoccus caerfyrddinensis* sp. nov.

*Pyxidicoccus caerfyrddinensis* (caer.fyrdd.in.en'sis N.L. masc. adj. *caerfyrddinensis* from Caerfyrddin, reflecting its isolation from soil sampled near Carmarthen (the Anglicised name for Caerfyrddin), Wales [51.86°N 4.31°W]).

Vegetative cells are Gram-negative bacilli tapering slightly at the ends, measuring 0.7-0.8 µm x 3.0-8.0 µm in electron micrographs. Colonies exhibit swarming motility and appear pale brown on VY-2 agar (w/v 0.5 % dried baker's yeast, 0.1% CaCl<sub>2</sub>·2H<sub>2</sub>O, 1.5 % agar). Fruiting bodies are irregular spheroids, orange in colour. Aerobic growth was observed at 30-40 °C, and at pH 6.0-9.0. Growth was unaffected by the addition of 1-3 % NaCl. Hydrolyses arginine, esculin, gelatine, p-nitrophenyl-β-D galactopyranoside and urea. Cells prey efficiently on *Clavibacter nebraskensis* DSM 7483, and with low efficiency upon *Escherichia coli* TOP10 and *Ustilago maydis* DSM 14603.

DNA GC content is 70.2 mol%. The draft genome sequence of CA032A<sup>T</sup> is available from GenBank (Accession JAAIYA000000000). The type strain (CA032A<sup>T</sup> = NCCB 100776<sup>T</sup> = NBRC 114353<sup>T</sup>) was isolated from soil collected in Carmarthen, United Kingdom [gridref 51.86°N 4.31°W].