

Corrigendum: Comparative genome sequence analysis of several species in the genus *Tepidimonas* and the description of a novel species *Tepidimonas charontis* sp. nov

Luciana Albuquerque^{1,*}, Nadine Castelhana¹, Pedro Raposo¹, Hugo J. C. Froufe², Igor Tiago³, Rita Severino¹, Inês Roxo¹, Inês Gregório¹, Cristina Barroso^{1,2}, Conceição Egas^{1,2} and Milton S. da Costa¹

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In the published version of this article there were some errors in Table S1 which are corrected in this version.

The title of Table S1 should have read: ‘Differential characteristics of *Tepidimonas* species. 1, strain SPSP-6^T; 2, strain SPSP-18; 3, *Tepidimonas alkaliphilus* YIM 72238^T; 4, *Tepidimonas aquatica* CLN-1^T; 5, *Tepidimonas fonticaldi* AT-A2^T; 6, *Tepidimonas ignava* SPS-1037^T; 7, *Tepidimonas sediminis* YIM 72259^T; 8, *Tepidimonas taiwanensis* I1-1^T; 9, *Tepidimonas thermarum* AA-1^T. +, positive; –, negative; ND, not determined. Strains SPSP-6^T, SPSP-18, *Tepidimonas thermarum* AA-1^T, *Tepidimonas ignava* SPS-1037^T, *Tepidimonas aquatica* CLN-1^T, *Tepidimonas taiwanensis* I1-1^T and *Tepidimonas fonticaldi* AT-A2^T assimilate succinate, L-glutamate and L-glutamine. Strains SPSP-6^T, SPSP-18, *Tepidimonas thermarum* AA-1^T, *Tepidimonas ignava* SPS-1037^T, *Tepidimonas aquatica* CLN-1^T and *Tepidimonas taiwanensis* I1-1^T assimilate lactate, pyruvate, acetate, but do not assimilate D-galactose, D-mannose, trehalose, cellobiose, melibiose, raffinose, D-ribose, D-xylose, D-arabinose, L-arabinose, L-rhamnose, L-fucose, L-sorbose, sucrose, lactose, maltose, ribitol, xylitol, sorbitol, erythritol, D-mannitol, *myo*-inositol, glycerol, benzoate, formate, glycine, L-methionine, L-serine and valine.’

On line 1 of the table ‘Characteristics, 8’ a reference letter was added in strain 8.

The table entries for rows ‘NaCl for growth (%)’, ‘Optimum and Range’ column ‘8’ previously ND, should have read 0.2 and 0–1 respectively.

The table entry for row ‘L-lysine’ column ‘9’ previously – should have read +. The table entries for row ‘L-proline’ columns ‘4, 6 and 9’ previously, – should have read + and entries for columns ‘3, 5 and 7’ previously nd, should have read +.

The table entry for row ‘L-isoleucine’ column ‘5’ previously nd should have read -. The table entries for row ‘L-ornithine’ columns ‘4, 6 and 9’ previously –, should have read +.

The authors apologize for any inconvenience caused.

Author affiliations: ¹Center for Neuroscience and Cell Biology, University of Coimbra, 3004-504 Coimbra, Portugal; ²Next Generation Sequencing Unit, Biocant, BiocantPark, Núcleo 04, Lote 8, 3060-197 Cantanhede, Portugal; ³Center for Functional Ecology, University of Coimbra, 3000-456, Coimbra, Portugal.

*Correspondence: Luciana Albuquerque, luciana@cnc.uc.pt
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