CORRECTION

Correction: Genomic and Genetic Diversity within the *Pseudomonas fluorescens* Complex

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There are errors in the names of type strains appearing in Figs 1 and 2 and S1 Fig. *P. aureofaciens* LMG 2145 should be *P. aureofaciens* 1245. *P. amygdali* LMG1384 should be *P. amygdali* 13184. *P. resinovorans* LMG 2774 should be *P. resinovorans* LMG 2274. *P. mandelii* LMG 2210 should be *P. mandelii* LMG 21607. *P. luteola* LMG 21607 should be *P. luteola* LMG 7041. Please see the corrected Figs 1 and 2 and S1 Fig here.

OPEN ACCESS

Citation: Garrido-Sanz D, Meier-Kolthoff JP, Göker M, Martín M, Rivilla R, Redondo-Nieto M (2016) Correction: Genomic and Genetic Diversity within the *Pseudomonas fluorescens* Complex. PLoS ONE 11(4): e0153733. doi:10.1371/journal.pone.0153733

Published: April 11, 2016

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Fig 1. Phylogeny of the *Pseudomonas* genus inferred by MLSA. Phylogenetic tree of 451 *Pseudomonas* strains along with 107 type strains based on the concatenated partial sequences of the 16S rDNA, *gyrB*, *rpoD* and *rpoB*, ML method, Tamura-Nei. Only bootstrap values above 75% (from 1,000 replicates) are shown. *Cellvibrio japonicum* Ueda 107 was used as outgroup. Details are found in [S1 Fig](#).

doi:10.1371/journal.pone.0153733.g001

Fig 2. Phylogeny of the *P. fluorescens* complex inferred by MLSA. Phylogenetic tree of 127 sequenced and type strains belonging to the *P. fluorescens* complex based on the concatenated partial sequences of the 16S rDNA, *gyrB*, *rpoD* and *rpoB* genes, the ML method and the Tamura-Nei model. Only bootstrap values above 50% (from 1,000 replicates) are shown. The *P. aeruginosa* type strain was used as outgroup. Bold text as well as superscript ^T indicates type strains. Strains are colored according to the groups established in this work.

doi:10.1371/journal.pone.0153733.g002

Supporting Information

S1 Fig. *Pseudomonas* genus MLSA. MLSA based on partial sequences of 16S rDNA, *gyrB*, *rpoD* and *rpoB* genes from 451 sequenced genomes and 107 type strains (bold), ML method and Tamura-Nei model. *C. japonicus* Ueda 107 was used as outgroup. Only bootstrap values above 75% over 1000 replicates are shown. Bold and ^T indicates type strain.
(PDF)

Reference

1. Garrido-Sanz D, Meier-Kolthoff JP, Göker M, Martín M, Rivilla R, Redondo-Nieto M (2016) Genomic and Genetic Diversity within the *Pseudomonas fluorescens* Complex. PLoS ONE 11(2): e0150183. doi: [10.1371/journal.pone.0150183](https://doi.org/10.1371/journal.pone.0150183) PMID: [26915094](https://pubmed.ncbi.nlm.nih.gov/26915094/)