CORA Cork Open Research Archive
Cartlann Taighde Oscailte Chorcaí

Title	Comparative genomic analysis reveals a diverse repertoire of genes involved in prokaryote-eukaryote interactions within the Pseudovibrio Genus
Authors	Romano, Stefano;Fernàndez-Guerra, Antonio;Reen, F. Jerry;Glöckner, Frank O.;Crowley, Susan P.;O'Sullivan, Orla;Cotter, Paul D.;Adams, Claire;Dobson, Alan D. W.;O'Gara, Fergal
Publication date	2016-03-30
Original Citation	Romano, S., Fernàndez-Guerra, A., Reen, F. J., Glöckner, F. O., Crowley, S. P., O'Sullivan, O., Cotter, P. D., Adams, C., Dobson, A. D. W. and O'Gara, F. (2016) 'Comparative genomic analysis reveals a diverse repertoire of genes involved in prokaryote-eukaryote interactions within the Pseudovibrio Genus', Frontiers in Microbiology, 7, 387 (20pp.). doi: 10.3389/fmicb.2016.00387
Type of publication	Article (peer-reviewed)
Link to publisher's version	http://journal.frontiersin.org/article/10.3389/fmicb.2016.00387/full - 10.3389/fmicb.2016.00387
Rights	© 2016, Romano, Fernàndez-Guerra, Reen, Glöckner, Crowley, O'Sullivan, Cotter, Adams, Dobson and O'Gara. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) or licensor are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms - https://creativecommons.org/licenses/by/4.0/
Download date	2026-10-16 17:24:40
Item downloaded from	https://hdl.handle.net/10468/4114

University College Cork, Ireland
Coláiste na hOllscoile Corcaigh

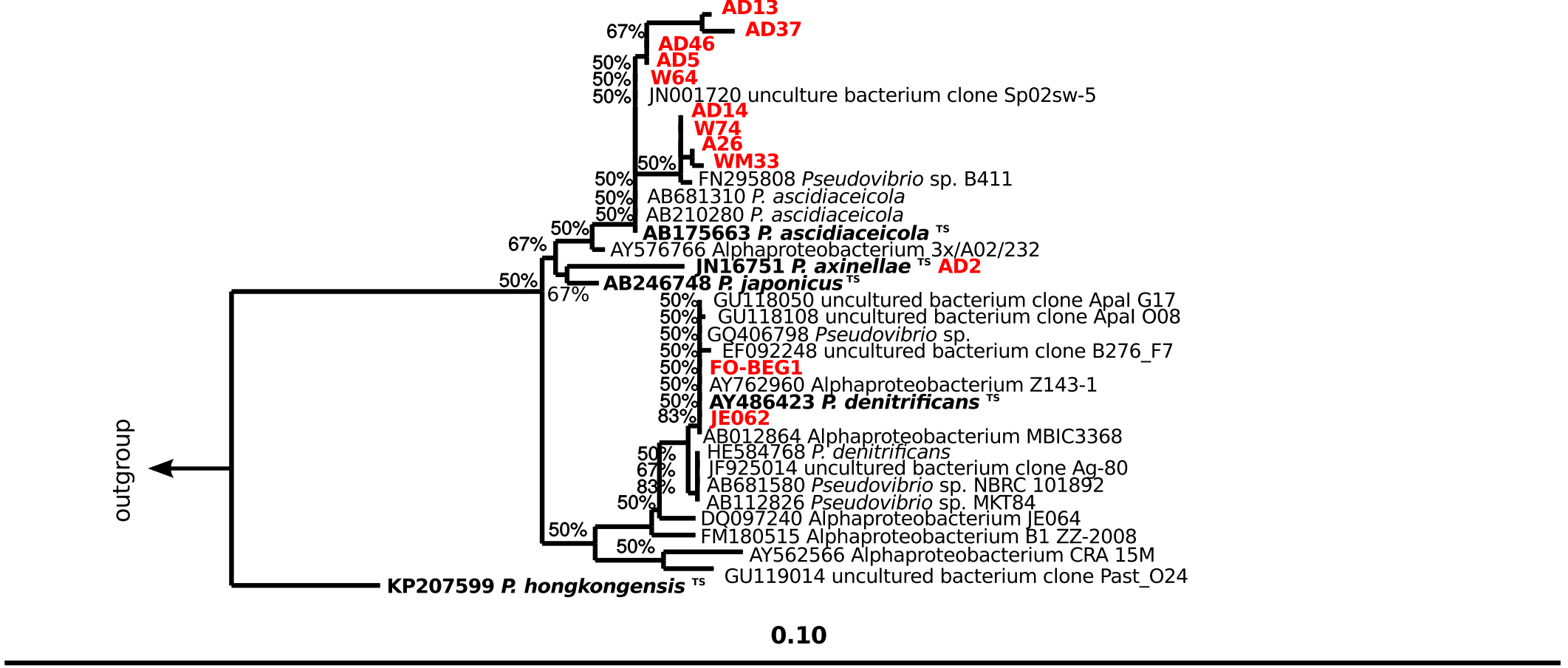

Figure S1 16S rRNA genes phylogenetic analysis. Consensus phylogenetic tree constructed using the 16S rRNA genes of *Pseudovibrio* related bacteria. The values at the nodes indicate the bootstrap obtained during the reconstruction of the consensus tree performed with the build-in function in the ARB program. Strains investigated in the present work are indicated in bold red. Type stains are reported in bold black with the superscript "TS"