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Community proteogenomics reveals the systemic impact of phosphorus availability on microbial functions in tropical soil

Qiuming Yao¹, Zhou Li¹, Yang Song¹, S. Joseph Wright², Xuan Guo³, Susannah G. Tringe⁴, Malak M. Tfaily⁵, Ljiljana Paša-Tolić⁵, Terry C. Hazen^{1,3}, Benjamin L. Turner², Melanie A. Mayes¹ and Chongle Pan^{1,3*}

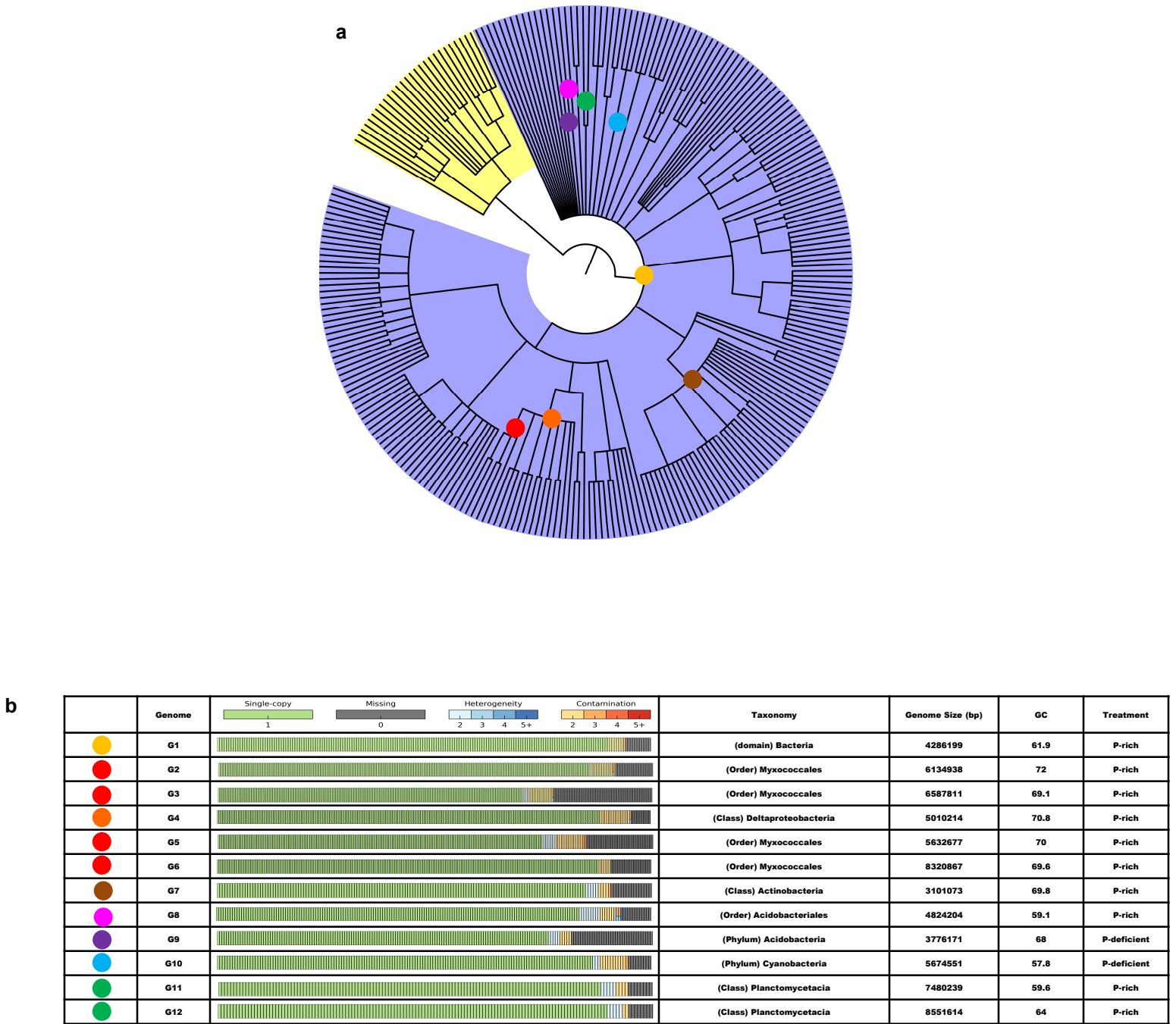
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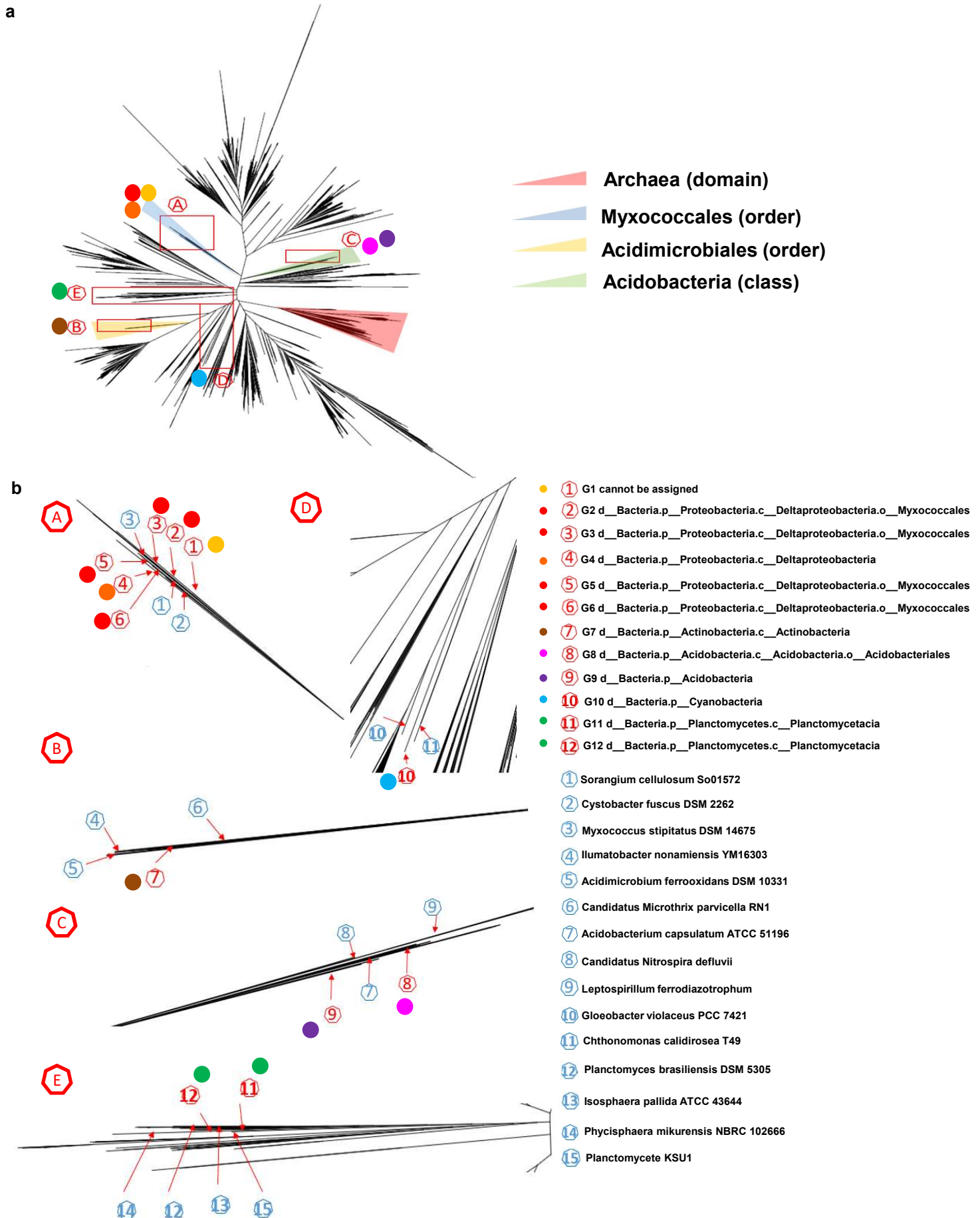
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Supplementary Figure 1: Taxonomy assignments (a) and quality metrics (b) of the near-complete genomes.



Supplementary Figure 2. Phylogenetic analysis of the near-complete genomes. (a) Full phylogeny tree. (b) Selected branches. The new genomes are shown in red numbers and the reference genomes are shown in blue number.