

Special Issue

Method Development for Pan-Genome Research on Microbes

Message from the Guest Editor

With the advance of microbiome research, more and more previously unknown species have been realized by researchers worldwide, together with their genome sequences. These species and genomes have constituted a grand pool of new resources for a better understanding of the evolution and ecology patterns of microbes, including virus, bacteria, archaea, fungi, and protists. Faced with the paramount number of species and their accompanying omics data, new tools are also needed. In this Special Issue on “Method Development for Pan-Genome Research on Microbes”, we welcome works that focus on novel methods for mining evolution and ecology patterns of microbes from both pan-genome and metagenome.

Guest Editor

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Deadline for manuscript submissions

closed (30 April 2022)

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Message from the Editor-in-Chief

Genes is central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fast-moving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised. Why not consider *Genes* for your next genetics paper?

Editor-in-Chief

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