



Diversity and Evolution of Phage Genomes

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Message from the Guest Editors

The increasing problem of multidrug-resistant bacteria has renewed interest in phages as biological control agents. Despite its enormous potential, phage therapy is far from being widely accepted. Medical doctors' and veterinarians' principal concerns about the use of phages are due to its viral nature and potential for dissemination of toxins and virulence determinants. Furthermore, bacterial resistance mechanisms may render phage therapy useless soon if not properly applied. While the microbiome is now dominating the headlines, it is clear that in every microbial environment, we will find at least an equivalent number of viral particles; however, our current understanding of viral diversity and its coding potential is still in its earliest development.

As can be seen, both research areas converge on the need to characterize phage genomes and their insights into phage diversity and evolution. We aim to generate a collection of articles with recent advances in phage diversity and evolution through genomic sequencing that are paving the way for both basic and applied research in phage biology.





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Message from the Editor-in-Chief

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