



## Bacteriophage Genomes and Genomics: News from the Wild

Guest Editor:

**Dr. Marie-Agnès Petit**

Micalis Institute, INRA,  
AgroParisTech, Université Paris-  
Saclay, Jouy-en-Josas, France

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### Message from the Guest Editor

Dear Colleagues,

In the jungle of the phage world, genomes analyses have started to bring some order: Genomics becomes a reliable tool to classify phages, and ICTV has recently begun implementing taxonomic classifications, based primarily on genomic data. The raising interest in phage therapy has launched phage discoveries for a variety of bacterial species, allowing to reduce the over-representation of model phage-like coliphages in databases, and to reach a more equilibrated view on the phage world. Teaching on its side has opened an unprecedented view on mycophages, and new teaching projects now tackle other genera, such as Bacilli or Streptomyces. Finally, (meta)viromics and single whole genome amplification are also bringing to the shore a wealth of new genomes. As such, the time is ready to collect the most striking features of the recent discoveries into a single issue.

Dr. Marie-Agnès Petit

*Guest Editor*





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### Dr. Eric O. Freed

Director, HIV Dynamics and  
Replication Program, Center for  
Cancer Research, National  
Cancer Institute, Frederick, MD  
21702-1201, USA

## Message from the Editor-in-Chief

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*Viruses* Editorial Office  
MDPI, Grosspeteranlage 5  
4052 Basel, Switzerland

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