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Evolutionary Genetics of *Streptococcus pneumoniae*

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Deadline for manuscript submissions:

closed (31 July 2019)

Message from the Guest Editors

Understanding the biology of Streptococcus pneumoniae will allow for improved public health policies for a disease that kills up to 1 million children each year globally. We must also understand how this biology changes through time to help prevent pneumococcal infections and deaths; how does S. pneumoniae evolve? Additionally, we can gain insight into general trends of genomic evolution in bacteria using pneumococcus as a genomic model organism due its particular set of characteristics: its natural competence and high rates of recombination; the large number of diverse, sequenced genomes, especially through The Global Pneumococcal Sequencing Project; and the defined selection pressures acting on pneumococcal populations in the form of antimicrobial resistance and vaccine-serotype replacement.

We are interested in examining any aspect of pneumococcus evolution, such as temporal changes and differences between populations, explorations of the evolution of microbial community dynamics, competence, antagonistic interactions, antimicrobial resistance, and serotype replacement and additionally, we welcome explorations into the mechanics of its genome evolution.













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Message from the Editor-in-Chief

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