

ABSTRACT

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Prof. Dr. Sebastien Calvignac-Spencer

(Helmholtz Institute for One Health, Greifswald)

Pathogen evolution in flux (Evolutionary genomics of pathogens)

Host: Tobias Lenz

Molecular clock models have become a familiar fixture of molecular phylogenetics. Here, I will briefly review their application to the study of pathogen evolution, using examples that illustrate the surprising variations of substitution rates during pathogen evolution.