

Upcoming Events

*EVBC does not endorse any of the listings and is only involved in events marked as EVBC Events.

2nd Joint Meeting of the German Society for Immunology (DGfI) and the Italian Society of Immunology, Clinical Immunology and Allergology (SIICA)

10–13 September 2019, Munich, Germany

The meeting will cover current topics of basic and translational immunological research in plenary sessions, main symposia and workshops.

Abstract submission deadline: **05 May 2019**

22nd European Society for Clinical Virology's Annual Meeting

11–14 September 2019, Copenhagen, Denmark

ESCV brings together scientists and clinicians throughout Europe and promote public health and advance education, particularly medical education, in clinical and basic virology.

Early registration deadline: **31 May 2019**

Recent Selected Publications by EVBC Members

Inferring HIV-1 transmission networks and sources of epidemic spread in Africa with deep-sequence phylogenetic analysis. *Nat Commun* **10.1038/s41467-019-09139-4**

The Utility of Data Transformation for Alignment, *de novo* Assembly and of Short Read Virus Sequences. *Preprints* **10.20944/preprints201904.0014.v1**

Viral host prediction with Deep Learning. *bioRxiv* **10.1101/575571**

Functional RNA Structures in the 3'UTR of Tick-Borne, Insect-Specific and No-Known-Vector Flaviviruses. *Viruses* **10.3390/v11030298**

RNA Structure Duplication in the Dengue Virus 3' UTR: Redundancy or Host Specificity? *MBio* **10.1128/mBio.02506-18**

BEAST 2.5: An advanced software platform for Bayesian evolutionary analysis. *PLoS Comput Biol* **10.1371/journal.pcbi.1006650**

Viral fitness: history and relevance for viral pathogenesis and antiviral interventions. *Pathog Dis* **10.1093/femspd/ftz021**

Bayesian inference of evolutionary histories under time-dependent substitution rates. *Mol Biol Evol* **10.1093/molbev/msz094**

Interpreting Viral Deep Sequencing Data with GLUE. *Viruses* **10.3390/v11040323**

Misinterpretation of Schmallenberg virus sequence variations: the sample material makes the difference. *Virus Genes* **10.1007/s11262-018-1628-2**

Integrative functional genomics decodes herpes simplex virus 1. *bioRxiv* **10.1101/603654**

Multi-Genome Annotation with AUGUSTUS. *Methods Mol Biol* **10.1007/978-1-4939-9173-0_8**

A Bioinformatics View of Glycan-Virus Interactions. *Viruses* **10.3390/v11040374**

Enhanced bioinformatic profiling of VIDISCA libraries for virus detection and discovery. *Virus Res* **10.1016/j.virusres.2018.12.010**

Vacancies

INEXTVIR - Innovative Network for Next Generation Training and Sequencing of Virome offers 15 PhD positions for 3 years with an attractive stipend, complementary training activities and generous travel and laboratory and research budgets. The successful candidates will be hosted in Belgium, France, Spain, Slovenia or the UK. Details on the different projects and the application process can be found [► here](#).

Application deadline: **May 14, 2019**

PhD-position/Postdoc-position in Bioinformatics at the Institute for Virology and Immunology of the University of Würzburg, Germany. The position is within the DFG-funded research unit "Advanced Concepts in Cellular Immune Control of Cytomegalovirus". Details regarding application can be found [► here](#).

Application deadline: **May 15, 2019**

