

CMSB 2020 brings together computer scientists, biologists, mathematicians, engineers, and physicists interested in a system-level understanding of biological processes. In particular, the conference is open to theoretical works with potential applications to modelling and systems biology. Contributions are welcome on all aspects of the analysis of biological systems, networks, data, and corresponding application domains.

Topics of interest include, but are not limited to:

Formalisms for modelling biological processes
Models and their biological applications
Frameworks for model verification, validation,
analysis and simulation of biological systems
Methods for synthetic biology and biomolecular
computing

Machine learning, model inference from experimental data

Model integration from biological databases Multi-scale modelling and analysis methods Collective behaviour

High-performance computational systems biology and parallel implementations

## PC co-chairs:

Alessandro Abate, University of Oxford (UK)
Tatjana Petrov, University of Konstanz (Germany)
Verena Wolf, Saarland University (Germany)

## **Confirmed Invited Speakers:**

Grégory Batt, Inria and Institute Pasteur, France lain Couzin, Max Planck Institute of Animal Behaviour & University of Konstanz, Germany Domitilla Del Vecchio, MIT, USA

## Format:

Papers should be submitted to one of the following categories:
Regular papers (15 pages, LNCS style)
Tool papers (6 pages, LNCS style)

## Important dates:

Tutorial abstract submission: April 1, 2020 Regular paper submission: April 15, 2020 Tool paper submission: April 15, 2020 Presentation submission: April 15, 2020 Poster abstract submission: June 15, 2020



Post-proceedings will appear as a volume in Springer LNCS/LNBI.