



COMBINE 2016: Newcastle upon Tyne, UK

19-23 September

http://co.mbine.org/events/COMBINE_2016/



The “Computational Modeling in Biology” Network (COMBINE) is an initiative to coordinate the development of the various community standards and formats in systems biology and related fields. COMBINE 2016 will be a workshop-style event with oral presentations, posters, and breakout sessions. The five meeting days will include talks about the COMBINE standards and associated or related standardization efforts, as well as presentations of tools using these standards. Oral presentations will be selected from the submitted abstracts. In addition, poster sessions will allow people to inform each other about their software and other projects in a setting that fosters interaction and in-depth discussion.

KEYNOTE SPEAKERS

- **Alfonso Bueno-Orovio** (*Oxford*)
- **Carole Goble** (*Manchester*)
- **Michael Hucka** (*Caltech*)
- **Dagmar Iber** (*ETH Zurich*)
- **Andrew Millar** (*Edinburgh*)
- **Chris J. Myers** (*Utah*)
- **Yujiang Wang** (*Newcastle*)

TOPICS OF INTEREST

- Data exchange and model standards for systems biology
- Graphical notation standards for systems biology
- Standards for sharing and analysing biological pathway data
- Standards for computational biological models

TRAVEL SCHOLARSHIPS

PhD students and postdocs may apply for travel funds to attend COMBINE. More information on the COMBINE website.

KEY DATES

Abstract submission: 8 July 2016
Scholarship application: 8 July 2016
Notification: 15 July 2016
COMBINE: 19-23 September 2016

LOCAL ORGANISERS

Paolo Zuliani & Anil Wipat
ICOS Group
School of Computing Science
Newcastle University

COMBINE COORDINATORS

Gary Bader (*Toronto*)
 Martin Golebiewski (*Heidelberg*)
 Michael Hucka (*Caltech*)
 Nicolas Le Novère (*Babraham*)
 Chris J. Myers (*Utah*)
 David Nickerson (*Auckland*)
 Falk Schreiber (*Monash*)
 Dagmar Waltemath (*Rostock*)

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