Bioinformatics Australia 2009 (BA2009)

Melbourne Convention and Exhibition Centre,
Melbourne, Australia, 29-30 October 2009

Conference Chair

Phoebe Chen (Deakin)

National Organization Committee

Phoebe Chen (Chair) (Deakin)
Catherine Abbott (Flinders)
Matthew Bellgard (Murdoch)
Bruno Gaëta (UNSW)
Dominique Gorse (QFAB)
Tim Littlejohn (IBM Australia)
Annette McGrath (AGRF)
Mark Ragan (UQ)
Shoba Ranganathan (Macquarie)

Local Organization Committee

Phoebe Chen (Chair) (Deakin)
Phil Batterham (Melbourne)
Ross Coppel (Monash)
Vladimir Likic (Bio21)
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Tony Papenfuss (WEHI)

Sponsorship Co-Chairs

Matthew Bellgard
(Bioplatforms Australia)
Tim Littlejohn (IBM Australia)
Mark Ragan (UQ)

Publication Co-Chairs:

Shoba Ranganathan (Macquarie)
Matthew Bellgard (Murdoch)

Overview:

The Bioinformatics Australia 2009 conference will be the forth national meeting held by the Bioinformatics Australia Society. This annual meeting will showcase Australia's scientific activity in the key areas of bioinformatics and computational biology. The previous meetings of Bioinformatics Australia were held in Sydney 2006, Brisbane 2007 and Gold Coast 2008 to provide a national forum for both academia and industry researchers.

In addition to the scientific content, this meeting will provide networking opportunities with bioinformatics and biotechnology stakeholders in Australia for students, users and developers of bioinformatics resources as well as ICT and life scientists interested in bioinformatics.

The Bioinformatics Australia 2009 conference will be held concurrently with the AusBiotech 2009 National Conference and sharing events, exhibition and networking space. The conference will include scientific sessions, panel sessions and discussion forum.

Call for abstracts

Participants are invited to submit abstracts for oral and poster presentations for the Bioinformatics Australia 2009 conference, for review. All accepted abstracts will be published in the BA2009 Proceedings.

Oral Presentations are invited in the following six symposium areas:

1. Computational Biology
2. Methods and Tools in Bioinformatics
3. Proteomics and Structural Bioinformatics
4. Comparative Genomics and Evolution
5. Systems Biology, Medical and Agricultural Applications
6. Emerging Areas, including Metabolomics and Immunoinformatics

Poster presentations are invited in all bioinformatics-related topics.

Closing dates for submission of abstracts:

Oral Presentation: **31 August 2009**
Poster Presentation: **15 September 2009**

Submission of abstracts via easychair at conference web site.

All participants need to be registered by **28th September 2009** 23:59 AEST to be included in the conference proceeding.