## GIVV 2008

http://mlaa.com.au/giw2008

Gold Coast, Australia

1-3 December

incorporating Bioinformatics Australia 2008 Conference

Sean Grimmond

Sequencing the transcriptome in toto

Institute for Molecular Bioscience The University of Queensland

Eugene Koonin

Evolutionary genomics in the age of 10,000 genomes

National Center for Biotechnology Information National Institutes of Health, Bethesda

Ming Li

Keynote Speakers

Modern homology search

Canada Research Chair in Bioinformatics University of Waterloo

Yi-xue Li

Modeling human genome-wide combinatorial regulatory networks initiated by transcription factors and microRNAs using forward and reverse engineering

Center for Bioinformation Technology Chinese Academy of Sciences Shanghai Jiaotong University

John Mattick

Misunderstanding the mammalian genome

**ARC Federation Fellow** Institute for Molecular Bioscience The University of Queensland

Eric Schadt

Constructing cross-tissue networks to elucidate disease circuits

CSIRO

Executive Scientific Director Genetics Rosetta Inpharmatics/Merck Research Labs







Mark Ragan

University of Queensland, Australia

Jonathan Arthur

University of Sydney, Australia

See-Kiong Ng

Institute for Infocomm Research, Singapore

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