

# CIBB2016: Computational Intelligence methods for Bioinformatics and Biostatistics

1st Day	2nd Day	3rd Day
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08:30	<i>Registration</i>		
09:00	<i>Opening</i>	Dr. Guido Sanguinetti <i>Invited Speaker</i>	Prof. Antonietta Mira <i>Invited Speaker</i>
09:20	Prof. Natalio Krasnogor <i>Invited Speaker</i>		
10:00			
10:20	10:20 Maria Raposo 10:40 Quirina Ferreira	10:00 Kevin Burke 10:20 Daniela Coca 10:40 Gilbert MacKenzie	10:00 Hiroshi Noborio 10:20 Veronica Biga 10:40 Igor Saggese
11:00	<i>Coffee Break</i>		
11:40	11:40 Riccardo Rizzo 11:58 Alexandru Floares 12:16 Antonio Eleuteri 12:34 Javier Palarea 12:52 Daniele Pepe	11:40 Mu Niu 11:58 Umberto Noe  12:16 Ivan Merelli 12:34 Esmail Nouriani 12:52 Anna Paola Carrieri	11:30 Chiara Brombin (Federica Cugnata) 11:48 Marco Bonetti 12:04 Alessandro Nonis 12:22 Andrea Giussani 12:40 Elia Biganzolia 12:58 Clelia Di Serio
13:10	<i>Lunch Break</i>		
14:30	Prof. Mark Beaumont <i>Invited Speaker</i>	Prof. Bud Mishra <i>Invited Speaker</i>	Dr. Daniela Paolotti <i>Invited Speaker</i>
15:30	15:30 Franco Masulli 15:50 Mohammad Kagdi 16:10 Andrew Schaumberg	15:30 Zakaria Benmounah 15:50 Kamel Zeltni  16:10 NVIDIA Guest talk	15:30 Leif Peterson 15:50 Davide Sardina 16:10 Susanna Conde (Xiaoguang Xu)
16:30	<i>Coffee Break</i>		
17:00	17:00 Dario Pescini 17:20 Paolo Cazzaniga 17:40 Vladimir Rogojin 18:00 Riccardo Colombo 18:20 Erin Scott	17:00 Kristy Hassall 17:20 Federico Andreis 17:40 Sanja Brdar 18:00 Daniele Pepe 18:20 Roberto Tagliaferri	17:00 Riccardo Rizzo 17:20 Fabio Cumbo 17:40 Gosue' LoBosco 18:00 Davide Sardina 18:20 Kevin Heffernan
18:40	<i>Closing day</i>		

CIBB Keynote

CIBB Guest Talk

Main Track

Engineering Bio-interfaces And Rudimentary Cells As A Way To Develop Synthetic Biology

Modeling And Simulation Methods for Systems Biology and Systems Medicine

Statistical Inference In Mechanistic Models Of Biological Systems

Biocuration And Integration Of Biomedical Databases

High-performance Computing And Deep Learning Methods For Genomic Data Analysis

Modeling Dependence In Survival Analysis: Methodological Issues And Biomedical Implications