# FORECASTING EVOLUTION ?

**JULY 8 – 11, 2015**
**CALOUSTE GULBENKIAN FOUNDATION, LISBON**

**ORGANIZERS:**

Isabel Gordo (Instituto Gulbenkian de Ciência)
Michael Lässig (University of Cologne)
Ville Mustonen (Wellcome Trust Sanger Institute)

Graphics: human influenza tree 2012–15 | Picture credit: Marta Łuksza

**INVITED SPEAKERS** include

Dan Andersson (Uppsala University)
Trevor Bedford (Fred Hutchinson Cancer Research Center)
Jesse Bloom (Fred Hutchinson Cancer Research Center)
Arup Chakraborty (Massachusetts Institute of Technology)
Sarah Cobey (University of Chicago)
Michael Desai (Harvard University)
Michael Doebeli (University of British Columbia)
Daniel Fisher (Stanford University)
Marco Gerlinger (Institute of Cancer Research London)
Michael Hochberg (Centre National de la Recherche Scientifique Montpellier)
Christopher Illingworth (Cambridge University)
Roy Kishony (Harvard University)
Natalia Komarova (University of California Irvine)
Richard Lenski (Michigan State University)
Stanislas Leibler (Rockefeller University)
Marta Łuksza (Institute for Advanced Study Princeton)
Luke Mahler (University of Toronto)
Berenike Maier (University of Cologne)
Leonid Mirny (Massachusetts Institute of Technology)
Richard Neher (Max Planck Institute Tübingen)
Armita Nourmohammad (Princeton University)
Colin Russell (University of Cambridge)
Sohrab Shah (University of British Columbia)
Boris Shraiman (University of California Santa Barbara)
Olivier Tenaillon (Inserm Paris)
Aleksandra Walczak (Ecole Normale Supérieure)

**REGISTRATION** preferably by March 31st.
Travel grants for junior scientists available.
www.sfb680.uni-koeln.de/forecasting_evolution.html

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MOLECULAR BASIS OF
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