

# Lausanne Genomics Days 2012

**Unil**  
UNIL | Université de Lausanne

Dorigny Campus, Génopode Building, Auditorium C (registration not required)

## Thursday February 16, 2012

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| <br><br><br><br><br><br> | <p>9:20 Welcome address</p> <p>9:30 John Moran, University of Michigan, Ann Arbor, USA<br/><b>Studies of a human retrotransposon</b></p> <p>10:15 Nicholas Katsanis, Duke University, Durham, USA<br/><b>Modeling the morbid human genome</b></p> <p>11:00 Coffee Break</p> <p>11:30 Julian Parkhill, Sanger Institute, Hinxton, UK<br/><b>Understanding transmission and evolution of bacterial pathogens through large-scale sequencing</b></p> <p>12:15 Jeffrey Jensen, École Polytechnique Fédérale de Lausanne (EPFL), Lausanne, CH<br/><b>What has genomics taught us about adaptation?</b></p> <p>13:00 LUNCH (sandwiches and drinks provided in Exhibition Hall)</p> <p>14:30 Han Brunner, Radboud University Nijmegen Medical Centre, Nijmegen, NL<br/><b>How next generation sequencing changes medicine</b></p> <p>15:15 Carole Charlier, University of Liège, Sart-Tilman, BE<br/><b>Serial translocation via circular intermediates underlies colour sidedness in cattle</b></p> |
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Organizers: Jacques Beckmann, David Gatfield, Keith Harshman, Félix Naef and Alexandre Reymond

## Friday February 17, 2012

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| <br><br><br><br><br><br><br><br><br> | <p>9:20 Welcome address</p> <p>9:30 Nicolas Galtier, Université Montpellier II, FR<br/><b>Species life-history traits and genome evolution</b></p> <p>10:15 Chris Bowler, Ecole Normale Supérieure, Paris, FR<br/><b>The Tara Oceans Expedition - A holistic approach for marine eco-systems biology</b></p> <p>11:00 Coffee Break</p> <p>11:30 Chris Jiggins, University of Cambridge, UK<br/><b>A butterfly genome offers insights into adaptation and speciation</b></p> <p>12:15 Maarten Koornneef, Max-Planck Institute, Köln, DE<br/><b>Exploiting natural variation in <i>Arabidopsis thaliana</i></b></p> <p>13:00 LUNCH (sandwiches and drinks provided in Exhibition Hall)</p> <p>14:30 Sophien Kamoun, The Sainsbury Laboratory, Norwich, UK<br/><b>Genome evolution in the Irish potato famine pathogen lineage</b></p> <p>15:15 Marc Robinson-Rechavi, University of Lausanne and Swiss Institute of Bioinformatics, Lausanne, CH<br/><b>Evolution and gene expression in vertebrates</b></p> |
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Organizers: Christian Fankhauser, Christian Hardtke and Tadeusz Kawecki