

PROBABILITY, POPULATION GENETICS and EVOLUTION
C.I.R.M., June 11–15 2012
Program

| | Lundi 11 | Mardi 12 | Mercredi 13 | Jeudi 14 | Vendredi 15 |
|-------|-----------------|-----------------|--------------------|-----------------|--------------------|
| 8:45 | Wakeley | Barton | Metz | Coop | Ewens |
| 9:35 | Evans | Taylor | Champagnat | Pfaffelhuber | Hermisson |
| 10:20 | <i>Coffee</i> | <i>Coffee</i> | <i>Coffee</i> | <i>Coffee</i> | <i>Coffee</i> |
| 10:50 | Griffiths | Pennings | Tran | Vatutin | Sagitov |
| 11:40 | Moehle | Krone | Véber | Klebaner | Bansaye |
| 12:30 | <i>Lunch</i> | <i>Lunch</i> | <i>Lunch</i> | <i>Lunch</i> | <i>Lunch</i> |
| 14:00 | | | | | Winter |
| 14:50 | | | | | Delmas |
| 15:40 | | | | | Bertoin |
| 16:30 | <i>Tea</i> | <i>Tea</i> | | <i>Tea</i> | <i>End</i> |
| 17:00 | Ralph | Kurtz | | Joyce | |
| 17:50 | Kersting | Popovic | | Blath | |
| 18:40 | Lambert | Birkner | | Berestycki | |
| 19:30 | <i>Dinner</i> | <i>Dinner</i> | <i>Dinner</i> | <i>Dinner</i> | <i>Dinner</i> |

List of the speakers in alphabetic order, with titles

Vincent Bansaye, Palaiseau

Small positive values for supercritical branching processes in random environment

Nick Barton, Wien

Mathematical problems in population genetics

Julien Berestycki, Paris

Branching processes and selection

Jean Bertoin, Zürich

The cut-tree of large Galton-Watson trees and the Brownian CRT

Matthias Birkner, Mainz

Ancestry in the face of competition

- Jochen Blath, Brelin
An ancestral recombination graph for diploid populations with skewed offspring distribution
- Nicolas Champagnat, Nancy
Adaptive dynamics in an individual-based, multi-resources chemostat model
- Graham Coop, Davis
Moving towards a general model of the coalescent with linked selection
- Jean-François Delmas, Marne-la-Vallée
Record sur les arbres de Lévy ou paradoxe sur les mutations favorables et neutres dans les CSBP
- Steven N. Evans, Berkeley
Go forth and multiply?
- Warren Ewens, Philadelphia
On the deterministic theory of population genetics and its possible stochastic extensions.
- Robert Griffiths, Oxford
The Lambda-Fleming-Viot process and a connection with Wright-Fisher diffusion
- Joachim Hermisson, Wien
Evolutionary rescue in structured populations
- Paul Joyce, Idaho
Characterizing the Distribution of Lysis Time and Burst Size in Lytic Phage
- Götz Kersting, Frankfurt
The total external branch length of evolving Kingman coalescent trees
- Fima Klebaner, Melbourne
The Long Run Age Structure of Population-Dependent General Branching Processes in Environments with A High Carrying Capacity
- Steven Krone, Idaho
Antibiotic resistance plasmids and spatial structure

- Tom Kurtz, Madison
Filtering and models in population biology
- Amaury Lambert, Paris
Coalescent point processes and phylogenies
- Hans Metz, Leiden
Conflict between alleles and modifiers in the evolution of genetic polymorphisms
- Martin Moehle, Tübingen
On Compound Poisson Population Models
- Pleuni Pennings, Harvard
Quantifying the evolution of drug resistance in HIV
- Peter Pfaffelhuber, Freiburg
Selective sweeps in structured populations
- Lea Popovic, Concordia
Stochastically induced bistability in Density Dependent Population Processes on Multiple Scales
- Peter L. Ralph, Davis
Exploring recent relatedness – IBD and biparental ancestry
- Serik Sagitov, Göteborg
Interspecies correlation for Brownian traits
- Jesse Taylor, Tempe
Gene Flow and the Population Genetics of Human Infectivity in East African Sleeping Sickness
- Chi Viet Tran, Lille
Limit theorems of historical processes in population dynamics
- Vladimir Vatutin, Moskow
Critical branching process with two types of particles evolving in asynchronous random environments
- Amandine Véber, Palaiseau
On the usefulness of genealogical trees

John Wakeley, Harvard

Gene genealogies within a fixed pedigree, and the robustness of Kingman's coalescent

Anita Winter, Essen–Duisburg

Evolving genealogies of spatial Λ -Cannings dynamics