

BASC Réseau Interactions Biotiques

Meeting « Population genomics : fungi as models »

22 avril 2014

Salle de conférence Bâtiment 362, Université Paris Sud

Programme provisoire

09:00	Ioana Brännström, Uppsala	Lichen genomics
09:20	Gladieux, Berkeley	<i>Neurospora</i> speciation genomics
09:40	Jesper Svedberg, Uppsala	Spore-killer genomics
10:00	Branco, Berkeley	<i>Suillus</i> ecological genomics
10:20	Break	
11:00	Ropars, Orsay Branca, Orsay Rodriguez de la Vega	<i>Penicillium</i> comparative genomics
11:30	Fournier, INRA Montpellier	Magnaporthe comparative genomics
11:50	Grandaubert, MPI Marburg	<i>Zymoseptoria tritici</i> population genomics
12:10	Lunch	
01:40	Pádraic Corcoran	The mating-type chromosome of <i>Neurospora tetrasperma</i>
02:00	Lendenmann, ETH Zurich	QTL mapping with RAD seq markers to identify candidate genes underlying ecologically relevant traits in <i>Zymoseptoria tritici</i>
02:20	Niewenhuis, Uppsala	Sexual dimorphism in <i>Neurospora</i>
03:00	Break	
03:20	Persoons, INRA Nancy	Identification of effectors in <i>Melampsora larici-populina</i> through an original approach: from population genomics to functional genomics
03:40	Ruxandra Molnar	Speciation genomics in <i>Neurospora tetrasperma</i>
04:00	Badouin, Orsay	<i>Microbotryum</i> speciation genomics
04:20	Fisher, IC London	<i>Bd</i> (amphibian-killer fungus) population genomics
04:40	Dutton, Harvard	Cheese ecosystem