

Pregramme des sessions

1-5 févr. 2021

Phylogeny and inference: from models to data and back

Afternoon

Institut Henri Poincaré, Amphithéâtre Hermite
11 rue Pierre et Marie Curie 75005 Paris

jeu. 4 février

13:00

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Session | **Site:** Institut Henri Poincaré, Amphithéâtre Hermite, 11 rue Pierre et Marie Curie 75005 Paris

13:30 - 14:00 **Amaury Lambert - The genomic view of diversification**

14:00 - 14:30

Eduardo Rocha - Gene flow, innovation and the invention of microbial functions

14:30 - 15:00 **Benny Chain - Capturing T cell receptor similarities**

16:00 - 16:30 **Martin Weigt - Wandering through sequence space**

16:30 - 17:00 **Michael Desai - Evolution as a stochastic algorithm**

17:00 - 17:30

Alison Feder - Inferring spatially-embedded genealogies in expanding populations

17:30 - 18:00

Daniel Weinreich - A high-throughput approach to study epistasis in proteins

19:00