

Phylo-Group Meeting II

Chadwick Lecture Theatre, UCL
Wednesday 18 February 2004
Organiser: Ziheng Yang (z.yang@ucl.ac.uk)

Directions:

<http://abacus.gene.ucl.ac.uk/paul/ucl-guide.html>

<http://www.ucl.ac.uk/CNDA/about/general/map.html>

Programme

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|----------------------|---|
| 10:50 | Ziheng Yang, Welcome, Introduction, Announcements |
| 11:00 — 11:30 | Ingo Ebersberger, Duesseldorf
<i>Comparing human and chimp genomes - What can we learn?</i> |
| 11:30 — 12:00 | Heiko Schmidt, Duesseldorf
<i>Quartet-based combined phylogenetic analysis of incomplete genesets</i> |
| 12:00 — 12:30 | Simon Whelan, Hinxton
<i>Effects of model choice on phylogenetic inference</i> |
| 12:30 — 14:00 | Lunch on your own |
| 14:00 — 14:30 | Beth Shapiro
<i>Population genetics of bison using ancient DNA</i> |
| 14:30 — 15:00 | Alexei Drummond
<i>Nonlinear molecular clocks for evolutionary inference</i> |
| 15:00 — 15:30 | Gerton Lunter, Oxford
<i>An irreversible dinucleotide substitution model, and root placement without outgroups</i> |
| 15:30 — 16:00 | Ziheng Yang, London
<i>Ad hoc rate smoothing for likelihood estimation of species divergence dates</i> |
| 16:00 — 16:20 | General discussions |

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Booked time: 10:00am - 17:00pm

Participants

Cambridge/EBI

Nick Goldman <goldman@ebi.ac.uk>

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Ari Loytynoja

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Beth Shapiro

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Gerton Lunter <lunter@stats.ox.ac.uk>

London

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Ying Cao <cao@ism.ac.jp> (Visitor, Institute of Statistical Mathematics, Tokyo)

Carlos Schrago <guerra@biologia.ufrj.br> (Visitor, Univ. Fed. Rio de Janeiro, Brazil)

Giselle Walker <gisw@nhm.ac.uk> (UCL)

Peter Foster <p.foster@nhm.ac.uk> (NHM)

Richard Goldstein <richard.goldstein@nimr.mrc.ac.uk> (NIMR)

Duesseldorf

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