

Phylo-Group Meeting

Garwood Lecture Theatre (South Junction, UCL)

Tuesday 16 September 2003

Organiser: Ziheng Yang (z.yang@ucl.ac.uk)

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

Programme

10:50 Ziheng Yang, Welcome, Introduction, Announcements

11:00 — 11:30 Paul-Michael Agapow, London
Embarrassing Phylogenetics

11:30 — 12:00 Tim Massingham, Hinxton
Using codon models to detect selection

12:00 — 12:30 Lee Bofkin/Nick Goldman, Hinxton
Identifying origins of replication

12:30 — 14:00 Lunch on your own

14:00 — 14:40 Ian Holmes, Oxford
Maximum likelihood estimation of indel rates by expectation maximisation

14:40 — 15:10 Thomas Schlegel, Duesseldorf
Dependencies within Sequences using Phylogeny

15:10 — 15:40 Jochen Kohl, Duesseldorf
Quality control of databases

15:40 — 16:20 Richard Goldstein, Mill Hill, London
Mixture models and their application to the analysis of G protein coupled receptors

16:20 — 16:30 Discussions (about talks and possible future meetings etc.)

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Participants

EBI, Hinxton

Nick Goldman
Simon Whelan
Carolin Kosiol
Ari Loytynoja
Lee Bofkin
Tim Massingham
Pietro Lio

Duesseldorf

Thomas Schlegel
Jochen Kohl
Nicole Scherer
Vinh Le Sy

UCL

Dave Dale
Becky Furlong
Paul-Michael Agapow
Samit Kundu
Wa Yang
Ziheng Yang
Daniel White
Giselle Walker (NHM)

Mill Hill, London

Richard Goldstein

Oxford

Ian Holmes
Alexei Drummond
Rune Lyngsoe