

# PhyloGroup VII Meeting

Friday 28 September 2007

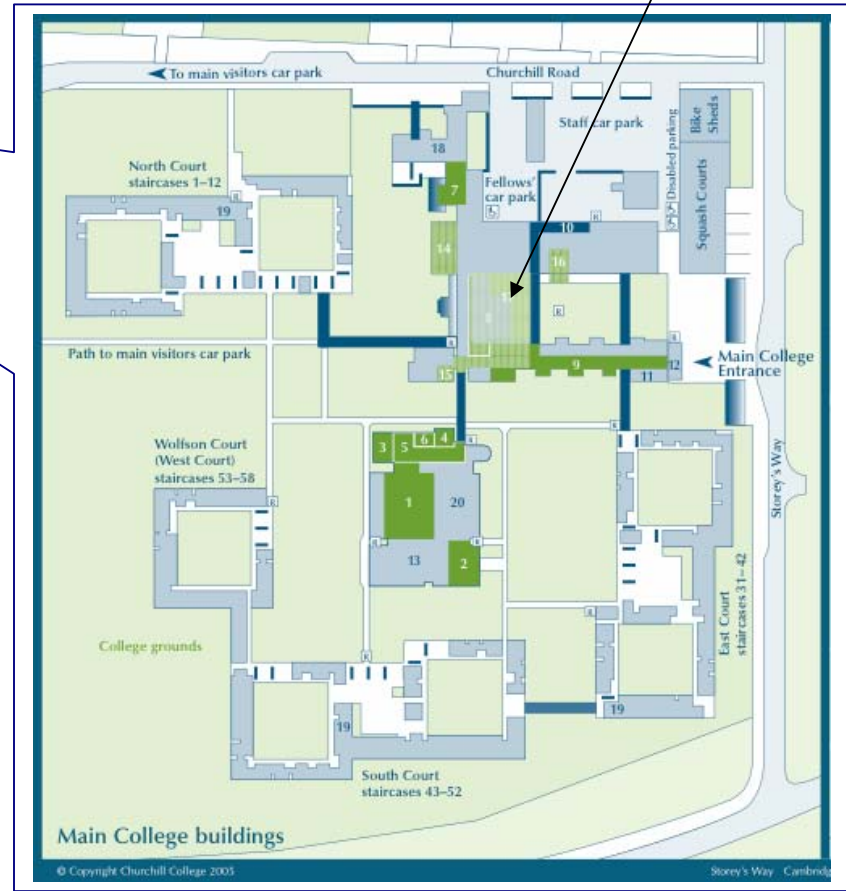
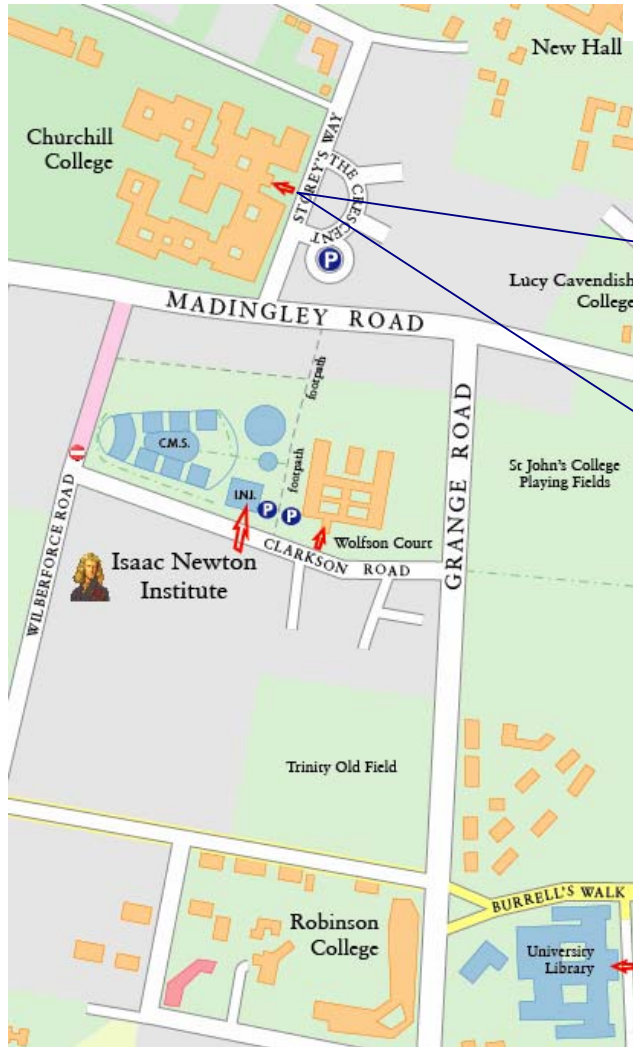
Isaac Newton Institute for Mathematical Sciences, Cambridge

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| <b>10:00</b>         | <b>Coffee at the Newton Institute (Mezzanine Level)</b>   |
| <b>10:29</b>         | <b>Welcome</b>  |
| <b>10:30 – 11:00</b> | <b>Tandy Warnow</b><br><i>Approaching multiple sequence alignment from a phylogenetic perspective</i>   |
| <b>11:00 – 11:30</b> | <b>Philippe Lemey</b><br><i>Recombination detection in human and primate immunodeficiency virus genomes using a quartet scanning approach</i> |
| <b>11:30 – 12:00</b> | <b>Kanchon Dasmahapatra</b><br><i>Can Pleistocene refugia explain current neotropical biodiversity</i>  |
| <b>12:00 – 12:30</b> | <b>Alexei Drummond</b><br><i>Random local clocks in BEAST</i>   |
| <b>12:30 – 13:30</b> | <b>Lunch at Churchill College</b>   |
| <b>14:00 – 14:30</b> | <b>Mario dos Reis</b><br><i>Timing the ancestor of human influenza A</i>  |
| <b>14:30 – 15:00</b> | <b>Simon Whelan</b><br><i>Testing the fit of a Phylogenetic model to observed data</i>  |
| <b>15:00 – 15:30</b> | <b>Martin Lott</b><br><i>Reconstructing the evolutionary history of polyploidy species</i>  |
| <b>15:30 – 16:00</b> | <b>Tea at the Newton Institute (Mezzanine Level)</b>  |
| <b>16:00 – 16:30</b> | <b>Sebastien Roch</b><br><i>Reconstructing phylogenies from very short sequences</i>  |
| <b>16:30 – 16:45</b> | <b>Close - Nick Goldman &amp; Ziheng Yang</b>   |

## List of participants

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# Newton Institute ~ Churchill College Map



Dining Hall