## Monday

8.45am 9.10am 9.50am	Peter Lockhart Welcome Mike Steel Reconstructing phylogenies from characters on a large state space Susan HolmesThe geometry of the space of phylogenetic trees	
10.30am	morning tea	
11.00am 11.40am	Thomas Hagedorn Calculating Invariants for the General Model Rissa Ota Appropriate Likelihood Ratio Tests and Marginal Distributions for Evolutionary Tree Models with Constraints on Parameters	
12.00am	Vincent Moulton Some recent results in phylogenetic combinatorics	
12.30pm	lunch	
2.00pm 2.40pm	Daniel H. Huson <b>Assembly of the Human Genome</b> Lindell Bromham <b>Endogenous retroviruses: data mining, phylogenetics and the risk of cross-species transmission</b>	
3.00pm	afternoon tea	
3.30pm 3.50pm 4.10pm	Anthony Poole Searching genomes for RNA world relics Paul Gardner ncRNA Gene Location Based on RNA Secondary Structure Matthew Goode Overview of the Cebl application environment underlying the Pebble	
Tuesday		
9.00am	Dorit Liebers Pattern of nucleotide substitution and rate heterogeneity in the HVR-I of the mitochondrial control region in large white-headed gulls ( <i>Larus argentatus - fuscus - cachinnans</i> complex)	
9.40am 10.00am	Meg Woolfit Tempo and mode of molecular evolution in island radiations Chris Simon Molecular Clocks, Geology, and the origins of New Zealand Cicadas	
10.30am	morning tea	
11.00am	Rob Smissen A 'total-evidence' approach to knawel (Scleranthus) phylogeny and biogeography	
11.20am	Alexei Drummond The inference of step-wise changes in substitution rates using serial	
11.40am	sequence samples Barbara Holland Median Networks: A visual representation of ancient Adelie Penguin DNA	
12.00am	Peter Lockhart Alpine buttercups	
12.30am	lunch	
2.00pm	Sverker Edvardsson Climatic variations over 1 million years before present from an astrophysical viewpoint: ice ages and global temperatures	
2.20pm	Alan Cooper Complete mitochondrial genome sequences of two extinct moas clarify ratite evolution	
2.40pm	Charles Semple <b>Hybrid Phylogenies</b>	
3.00pm	afternoon tea	
3.30pm	Lynne van Herwerden Extensive mitochondrial introgression in a tropical reef fish	
3.50pm 4.10pm	(Acanthochromis polyacanthus) at a contact zone between colour morphs Andy McKenzie Some Distance and Balance Properties of a Simple Speciation Model Madeleine JH van Oppen Reticulate evolution in reef corals of the genus Acropora	

## Wednesday night

7.30pm Lars S. Jermiin **Mitochondrial DNA Sequences in Ancient Australians: Implications** 

for Modern Human Origins

8.00pm David Penny Another major challenge to the classical Multiregional Model?

## **Thursday**

8.45-9.50am	breakfast at the Chateau
10.00am	Russell Gray Recovering reticulation in human evolution: trees, split graphs, and Pacific language
10.40am	Matthew E. Hurles The Y-chromosomal origins of Oceanic-speaking populations
11.00am	Andrew Rambaut Phylogenetic Tree structure and the origins of HIV 1
11.20am	Allen Rodrigo The complexity of HIV evolution
12.00am	Michael Charleston Principles of co-phylogeny on HIV and primate co-evolution
12.30pm	lunch
2.00pm	Kerryn Slack and Abby Harrison Mitochondrial Genomes and Avian Phylogeny
2.20pm	Matthew Phillips Mammal Phylogeny and Evolutionary Stable Niche Discontinuities
2.40pm	Yu-hsin Owen Lin A Stable Tree for eutherian mammals?
3.00pm	afternoon tea
3.30pm	Alex Grossman Evolutionary individuality of amino acids
4.10pm	Lars Jermiin Jumping Genes and Slithering Proteins: A Tale About Genes and Proteins
	Migrating Between Genomes

## **Friday**

9.00am 9.40am	Tony Larkum <b>Will the hypothesis of a single plastid origin be (dis) proven?</b> David Bryant <b>From gene orders to phylogeny</b>
10.00am	morning tea
10.30am	Nadia El-Mabrouk Reconstructing ancestral genomes by reversals and multiple gene duplication
10.50am	David Penny Reality Check
11.45am	early lunch