	22 June, Monday	23 June, Tuesday	24 June, Wednesday	25 June, Thursday

Morning session I.		9:00-9:50
		  Laszlo A. Szekely:
		  Reconstruction and Testing in Phylogenetic combinatorics
		9:50-10:40
		  John A. Rhodes:
		  Identifiability of Phylogenetic Models: When is consistent inference possible?
			9:00-9:50
			  Stephen Willson:
			  Reconstructing regular networks from their trees
			9:50-10:40
			  Vincent Moulton:
			  Blobs, blocks and other cyclic elements
				9:30-9:50
				  Kathi Huber:
				  Encoding of phylogenetic networks of bounded level
				9:50-10:40
				  Elizabeth S. Allman:
				  Two tree mixtures of group-based models

COFFEE	**Registration**	10:40-11:20 *COFFEE*	10:40-11:20 *COFFEE*	10:40-11:20 *COFFEE*

Morning session II.	Informal discussions	11:20-12:10
		  Sagi Snir:
		Quartet MaxCut: Piecing Together Small Trees into the Large Tree of Life
		12:10-13:00
		  Alberto Apostolico:
		Sequence Distances by Exhaustive Substring Composition
			11:20-12:10
			  Miklós Csűrös:
			Counting on phylogenies: ancestral reconstruction of numerical characters
			12:10-13:00
			  Bernard Moret:
			Genomic Evolution: Recent Progress
				11:20-12:10
				  István Miklós:
				Sampling from posterior distributions
				12:10 – 12:30
				  Mareike Fischer
				Hereditary Maximum Parsimony trees – and not so hereditary ones

LUNCH	13:00-14:30 *LUNCH*	13:00-14:30 *LUNCH*	13:00-14:30 *LUNCH*	13:00-14:30 *LUNCH*

Afternoon session I.	14:30-15:20
	  Mike Steel:
	Mixed up trees: The geometry of phylogenetic mixtures
		14:30-14:50
		  Filip Cools:
		Tropical geometry and dissimilarity maps of trees
		14:50-15:10
		  Éva Czabarka:
		Minimizing the number episodes on a species tree - an extension of Gallai's theorem on intervals
		15:10-15:30
		  Simone Linz:
		Computing tree distances: A divide-and-conquer approach
			14:30-14:50
			  Andrew Wei Xu:
			Getting More by Expecting Less: A new approach to the inversion median problem
			14:50-15:10
			  Marc Thuillard
			Minimum Contradiction Method
			15:10-15:30
			  Beáta Faller:
			Combinatorial challenges in conservation biology
				Informal discussions

COFFEE	15:20-16:00 *COFFEE*	15:30-16:00 *COFFEE*	15:30-16:00 *COFFEE*	
Afternoon session II.	16:00-16:50 Benny Chor: *Phylogenetic reconstruction based on whole genomes - initial results*	Informal discussions	Informal discussions	

	17:30-
	  Wine & cheese party
			18:00-
			  Boat tour	