



**Annual Network Meeting**  
**Divani Apollon Palace & Spa, Vouliagmeni, Athens, Greece**  
**9-11 April 2008**

DATE	SESSION	TIME	SPEAKER	AFFILIATION	TITLE
Wednesday	Welcome	15.00-19.00	Arrival & Registration		
		19.30	Evening Reception and Welcome by Jon Austyn - Mythos of the Sea Restaurant		
Thursday 10th April Plenary Room	Session I Clinical	8.30-8.50	Gerold Schuler, Chair	University of Erlangen, DE	Dendritic cell trials
		8.50-9.20	Chris Schmidt	Queensland Institute of Medical Research, AU	Immunological monitoring of immunotherapy trials
		9.20-9.50	Inge Marie Svane	University Hospital Herlev, DK	The Danish experience of DC-based immunotherapy of patients with disseminated breast cancer and renal cell carcinoma – Clinical outcome, biomarkers and immune parameters
		9.50-10.10	Alexander Dohnal	Trimed Biotech, AT	DECAVAREC - An interleukin-12 secreting dendritic cell-based cancer vaccine for the treatment of metastatic renal cell carcinoma
		10.10-10.30	Carl Figdor	Nijmegen Center for Molecular Life Sciences, NL	Towards the next generation of DC vaccines
		10.30-11.00	Coffee break		
Thursday 10th April Plenary Room	Session II Models	11.00-11.15	Muriel Moser, Chair	Free University of Brussels-ULB, BE	The clinical and pre-clinical interface
		11.15-11.35	Markus Manz	Institute of Research in Biomedicine, IT	Human hemato-lymphoid system mice as preclinical in vivo models
		11.35-11.55	Veronique Quillien	Centre Eugène Marquis Rennes, FR	Brain tumour stem cells could be favourable targets for immunotherapy
		11.55-12.10	Cheryl Chiang	University College London, UK	Oxidation by hypochlorous acid breaks tumor tolerance and stimulates T cells which recognize autologous primary tumor
		12.10-12.30	Sebastian Amigorena	Institut Curie, FR	Antigen cross presentation and T cell activation by dendritic cells
		12.30-14.00	Lunch		
Thursday 10th April Plenary Room	Session II Models	14.00-14.20	Marc Gregoire	INSERM U601 Nantes, FR	Sequentially matured human DC with CD4+ T cells as secondary signals favour Th1, CTL and long term memory T cells responses
		14.20-14.40	Gerald Bos	University Hospital Maastricht, NL	The maturation cocktail for DC generation influences induction of MUC1 specific immune responses
		14.40-15.00	Steve Volland	University of Erlangen, DE	T-cell responses in stage IV melanoma patients vaccinated with dendritic cells electroporated with defined RNA
		15.00-15.15	Özcan Met	University Hospital Herlev, DK	High immunogenic potential of p53, surviving and hTERT mRNA-electroporated dendritic cells in primary breast cancer
		15.15-15.30	Ann van Nuffel	Free University of Brussels-VUB, BE	pST1 derived mRNA leads to more and longer protein expression after electroporation: implications of vector choice on activation of monocyte derived DC and antigen presentation
		15.30-16.00	Coffee break		
Thursday 10th April Plenary Room	Session II Models	16.00-16.20	Isabelle Dunand	University of Geneva, CH	Role of microRNA-155 in mature dendritic cells
		16.20-16.40	Éva Rajnavölgyi	University of Debrecen, HU	Modulation of dendritic cell functions by miR-146a and miR-155
		16.40-17.00	Sandra Gessani	Institute of Health - ISS, IT	Role of cytokine environment and cytokine receptor expression in human monocytes in the generation of functionally distinct dendritic cells
		17.00-17.15	Alessandra Mancino	Humanitas Institute, IT	Divergent Effects of Hypoxia on Dendritic Cell Functions
		17.15-17.30	Gabi Theiner	University of Erlangen, DE	Low doses of denileukin difitox (ONTAK) induce survival rather than depletion of CD4+CD25+ regulatory T cells in
		17.30-18.30	General discussion		
		19.30	Buffet dinner		



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 [Continued]

DATE	SESSION	TIME	SPEAKER	AFFILIATION	TITLE
	Session II Models	8.30-8.50	Kees Melief	Leiden University Medical Center, NL	Immunotherapy of established lesions induced by high risk Human Papilloma Virus type 16
		8.50-9.05	Jan Doerrie	University of Erlangen,DE	DC cancer vaccines: development of new strategies for antigen loading
		9.05-9.20	Niels Schaft	University of Erlangen,DE	Many antigen presenting cells, or much antigen per presenting cell: studies on priming and expansion of melanA-specific T cells
		9.20-9.40	Silvano Sozzani	Humanitas Institute	<i>To be confirmed</i>
	Session III Systems	9.40-10.00	Duccio Cavalieri, Chair	University of Florence, IT	Querying a DC "omics" database according to a pathway based logic
		10.00-10.15	Carl Figdor	Nijmegen Center for Molecular Life Sciences, NL	The 'golden experiment'
		10.15-10.30	Luca Beltrame	Frienze / Milan, IT	Pathway Based analysis allows comparison between different datasets and microarray platforms
		10.30-11.00	Coffee break		
Friday 11th April Plenary Room	Session III Systems	11.00-11.20	Sonja Buschow	Nijmegen Center for Molecular Life Sciences, NL	Proteomic analysis of human dendritic cell maturation
		11.20-11.40	Philippe Pierre	CIML-CNRS, FR	Translational profiling of activated monocyte-derived DCs
		11.40-11.55	Lisa Rizzeto	University of Florence, IT	Different forms of <i>Saccharomyces cerevisiae</i> elicit specific responses in dendritic cells
		11.55-12.15	Benny Chain	University College London, UK	Global transcriptomic analysis identifies defects in key functional clusters of genes in dendritic cells derived from leukaemic MUTZ-3 precursors
		12.15-13.45	Lunch		
Friday 11th April Plenary Room	Session III Systems	13.45-14.15	Natalie Pochet	Broad Institute, MIT, USA	A module network approach to the unraveling of the molecular pathways underlying the physiological diversity of the in vivo malaria parasite, <i>Plasmodium falciparum</i> , and its interactions with the human host
		14.15-14.35	Henri de la Salle	INSERM U725, FR	Comparison of the transcriptomes of CD34-derived and human skin Langerhans cells
		14.35-14.55	Gianluca Matteoli	IFOM-IEO, IT	Epithelial cells as modulator of intestinal immuno-homeostasis
		14.55-15.30	Sorin Draghici	Wayne State University, USA	Pathway-based approaches to the analysis of microarray data
		15.30-16.00	Coffee break		
Friday 11th April Plenary Room	Session III Systems	16.00-16.20	Andrea Splendiani	LEAF Biosciences, IT	Use of ontologies in pathways and high-throughput data analysis
		16.20-16.35	Francesca Zollezzi	Genopolis, IT	Improvement of AMDA: the platform for automated data analysis of gene expression experiments
		16.35-16.55	Marco Brandizi	LEAF Biosciences, IT	A collaborative model for representing and managing microarray-related knowledge
		16.55-17.10	Michaela Guendel	LEAF Biosciences, IT	The DC-THERA Directory
	Closing	17.10-18.00	General discussion of the Project, its new initiatives, and future Network sustainability		
		19.30	Farewell dinner		