



Bulletin d'inscription

Nom Prénom

Titre Adresse du laboratoire

Ville, Code Postal, Pays

Téléphone Fax

E-mail

Les frais d'inscription au congrès comprennent l'accès au programme scientifique, l'envoi des résumés et de la plaquette d'information, les pauses-café et les déjeuners. Les inscriptions sont à envoyer **avant le 15 mars 2004** à l'Institut de Médecine Légale, à l'attention de M. Sylvain AMORY, 11 rue Humann, 67085 Strasbourg cedex, accompagnées d'un chèque ou d'un virement bancaire à l'ordre de **Monsieur l'Agent Comptable - Université Louis Pasteur compte TP Strasbourg n° 10071 67000 00003005898 83.**

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Nouvelles Technologies pour l'Etude de l'ADN Humain



Strasbourg, France Mai 2004

Frais d'Inscription au congrès :

	1 jour	2 jours	3 jours
Membres IALM	70 €	120 €	150 €
Autres	90 €	160 €	200 €
	<input type="checkbox"/> 12 Mai	<input type="checkbox"/> 13 Mai	<input type="checkbox"/> 14 Mai

DÉLAIS D'INSCRIPTION
15 MARS 2004

Modes de paiement :

- Eurochèque ou chèque international
- Virement bancaire

The SNIp years

12th May

Promises for SNP analysis: Reflections and results

Morning

Welcome presentation

B. LUDES

(Strasbourg)

Perspectives and challenges for the analysis of single nucleotide polymorphism in forensic genetics

P. SCHNEIDER

(Mainz)

SNP analysis technologies in comparison: SNaPshot, qPCR and Mass Spectrometry

W. PARSON (Innsbruck)

SNP typing with multiplex PCR based methods: comparison of SNaPshot, Maldi-Tof MS and Nanogen electrohybridisation

N. MORLING (Copenhagen)

Afternoon

SNP typing for Forensic Genetics

A. CARRACEDO (Santiago de

Compostela)

SNPs and Maldi-Tof MS: tools for individual identification

E. PETKOVSKI

(Strasbourg)

Y- STRs allele distributions inside SNPs defined haplogroups

L. GUSMAO (Porto)

SNP typing by MALDI-TOF MS - The GENOLINK system

D.

NIEMEYER (Leipzig)

Closing remarks

B. LUDES

(Strasbourg)

Degraded and/or ancient DNA

13th May

Degraded and/or ancient DNA

Morning

Welcome presentation	<i>E. CRUBEZY (Toulouse)</i>
General overview on degraded DNA (Toulouse)	<i>E. CRUBEZY</i>
Strategies for degraded DNA analysis (Mainz)	<i>K. BENDER</i>
Bone DNA typing: A pragmatic approach (Oslo)	<i>E. HAGELBERG</i>
Quantitative PCR and ancient DNA (Strasbourg)	<i>C. KEYSER-TRACQUI</i>

Afternoon

Paleomicrobiology: actual aspects and perspectives	<i>M. DRANCOURT (Marseille)</i>
GenomiPhi™: a solution for ancient DNA study? (Strasbourg)	<i>S. AMORY</i>
New technologies: Casework applications (Strasbourg)	<i>T. DELEFOSSE</i>
Anthropology	<i>Contributors from Kunming, Shanghai and Chongqing Medical Universities (China)</i>
Scytho-Siberian tomb at Berel (Kazakhstan) (Paris)	<i>H.-P. FRANCFORT</i>
The Mummies of Taklamakan <i>DEBAINE (Paris)</i>	<i>C.</i>
Closing remarks	<i>B. LUDES (Strasbourg)</i>
<i>Gala dinner: The Chinese table</i>	
Movie projection: The mummies of Taklamakan	

Neurology, Alzheimer's Disease and Cancerology

May 14th

Morning: Neurology - Alzheimer's disease

Welcome presentation

B. LUDES (*Strasbourg*)

G. VINCENDON (*Strasbourg*)

Presidents of Chinese Medical Universities of

Kunming, Shanghai and Chongqing (*China*)

Is there a molecular basis of the neuro-inflammation in Alzheimer's disease

J.-P. LOEFFLER (*Strasbourg*)

Therapeutic perspectives in Alzheimer's disease

F. SELLAL

(*Strasbourg*)

Exploring cognitive functions in central neural system pathologies: examples of visual perception

A. GIERSCH (*Strasbourg*)

Afternoon: Cancerology

STR analysis for quantitative assessment of patient/donor hematopoietic cells origin after allogenic stem cell transplantation

F.

WATZINGER (*Wien*)

Organisation of the Canceropole

P. OUDET

(*Strasbourg*)

Cancerology **Contributors from Kunming, Shanghai and Chongqing Medical Universities** (*China*)

Analysis of STRs for detection of the loss of heterozygosity in malignant neoplasia

F. WATZINGER (*Wien*)

Topoisomerase II Cancer Implications

C. VOEGELI (*Strasbourg*)

Free communications

Closing remarks

B. LUDES (*Strasbourg*)

G. VINCENDON (*Strasbourg*)

Presidents of Chinese Medical Universities of

Kunming, Shanghai and Chongqing (*China*)

Workshops: SNP typing

Sample analysis by Maldi-Tof MS on an ULTRAFLEXTM instrument

Sample analysis by real-time PCR on an ABI PRISMTM instrument