

## 37. Posters

### Late Breaking Abstracts (All Fields)

Friday, November 23, 9:00–19:30, Galerie Le Notre

- 37.1 Affinity Purification Using Short Synthetic Peptides: a Tool for the Study of Protein Complex Formation in Neurons  
**F. Buck, M. Christenn, D. Richter and H.-J. Kreienkamp**
- 37.2 ICPL- a New Strategy for Quantitative Proteomics  
**A. Schmidt, J. Kellermann and F. Lottspeich**
- 37.3 Proteomic analysis of plasma proteins of waste incinerating workers  
**D. Sul, H. Hong, S. Oh, H. Im, M. Yang, D. Jo, S. Jee and E. Lee**
- 37.4 Proteomic Analysis of The Human Alcoholic Brain  
**P.R. Dodd, J.M. Lewohl, D.D. Van Dyk, G. Craft and G.R.A. Harris**
- 37.5 Proteomic Analysis of Plasma Proteins of Automobile Emission Inspectors  
**E. Lee, S. Oh, H. Hong, H. Im, M. Yang, D. Jo, E. Oh and D. Sul**
- 37.6 Novel Software for Automatic Protein Identification  
**D. Dalevi, F. Levander, N. Nilsson, T. Rönvaldsson and J. Samuelsson**
- 37.7 Proteome Analysis of Gastric Cancer: Identification of Protein Keys for The Discrimination of Tumor and Normal Tissues  
**C.-W. Lee, E. H. Suh, S. J. Ahn and J. W. Kim**
- 37.8 Expressway™ Choose Your Route for Protein Analysis.  
**W. Kudlicki, J. Fletcher, S. Keppetipola and A. Coffman**
- 37.9 Pharmacoproteomic Screening in Diabetes and Obesity  
**C. Nicolas, L. Marion, P. Bernard, G. Jérôme and P. Pierre**
- 37.10 Isolation of Pure Cells, Nucleic Acids and Proteins from Single Samples  
**M. Bosnes, S. Bergholtz, A. Borgnes, E. Breivold, K. Lycke, A. Keiserud, T. Borgen and D. Lillehaug**
- 37.11 Complexity of Proteomes: Protein Minimization and Quantitative Arrangement  
**J. Klose**
- 37.12 Proteomic Profiling in *E. coli*: Comparison of Peptide Maps for Growth under Heat Shock and Non-Heat Shock Conditions  
**E. Naegele, M. Vollmer and P. Hoerth**
- 37.13 Expression Profile of *E. coli* under Heat Shock Conditions by 2D LC/MS  
**E. Naegele, M. Vollmer and P. Hoerth**
- 37.14 Identification of markers for selection of patients undergoing RCC-specific immunotherapy  
**M. Menig, R. Lichtenfels, B. Mühlenweg, D. Atkins, J. Bukur, T. Halder, A. Ackermann, J. Beck, W. Brenner, S. Melchior, F. Lottspeich and B. Seliger**
- 37.15 Affinity Electrophoresis for Functional Proteomics  
**K. Nakamura**

- 37.16** A Large Scale Survey of Protein Interactions in *C. elegans*  
**A. Nelsen, L. Peppers, D. Cyr, D. Kelly, N. Thota, X. Guan, M. Weiner and R. Hollingsworth**
- 37.17** Bioinformatics for Human Disease Proteome  
**Y. Zhu, G. Cheng, T. Chen, C. Du, S. Wu, P. Wan, D. Li, B. Li and F. He**
- 37.18** ProCorr<sup>®</sup>: Software for Protein Identification from High-Throughput Tandem-MS data  
**G. R. Thomas, J. Vandekerckhove and K. Gevaert**
- 37.19** Identification of Tumor Markers and Tumor-associated Antigens in Human Renal Cell Carcinoma Using Proteome-based Methods  
**B. Seliger, R. Lichtenfels, D. Atkins, M. Menig, J. Bukur, T. Halder, F. Lottspeich, A. Ackermann, J. Beck, B. Mühlenweg, W. Brenner, S. Melchior and R. Kellner**
- 37.20** Galectin-1 Function is Modulated by PTEN  
**D. H. Kim, S.Y. Kim, Y. Y. Bahk and Y. S. Kim**
- 37.21** Clinical Utility of the Plasma Proteome  
**A. Herath, C. Rohlff, T. Patel, J. Bruce, S. McGowan, A. Lyall, R. Townsend and R. Parekh**