

Proteomics in Cancer Research 3rd Rudbeck Symposium

November 11-12, 2004

Rudbeck Hall, The Rudbeck Laboratory, Uppsala University

Open to all

Organizers: Kenneth Nilsson and Fredrik Ponten

Thursday, November 11th, 2004

9:00 Kenneth Nilsson
Introduction

Molecular pathology – present methodology

9:15 Olli-Pekka Kallionemi, Turku,
Integration of microarray technologies for
diagnostics and cancer therapy development

10:00 Coffee

10:30 Anne-Lise Børresen, Oslo
The use of microarrays for molecular
profiling of breast and ovarian cancer

11:15 Jan Dumanski, Uppsala
Full-coverage genomic microarrays-
applications in research and diagnostics

12:00 Lunch

Proteomics in human cancer 1

13:30 Samir Hanash, Seattle
The use of proteomics for the diagnosis of cancer

14:15 Mark Rubin, Boston
Prostate cancer

15:00 Coffee

Proteomics - present methodology

15:30 Peter James, Lund,
Large-scale proteomics approaches
for defining cancer biomarkers

16:15 Hanno Langen, Basel
Proteomics as a tool for drug development (2D)

Friday November 12, 2004

9:00 Carl Borrebaeck, Lund,
Design of antibody microarrays for
high-throughput proteomics

Proteomics in human cancer 2

9:45 Samir Hanash, Seattle
Mapping of signaling pathways
related to tumorigenesis in human cancer

10:30 Coffee

11:00 Julio Celis, Copenhagen
Towards discovery-driven research
in breast cancer

11:45 Lunch

13:15 Gert Auer, Stockholm
Colorectal cancer proteomics

The Human Proteome Project

14:00 Mathias Uhlén, Stockholm
Antibody-based tissue proteomics

14:45 Fredrik Pontén, Uppsala
High throughput tissue profiling

15:30 Closing of the symposium

Combine with the
"Functional Genomics"
meeting in Stockholm Nov. 9-10!

For more information contact:
inger@biotech.kth.se

For further information please contact Elisabth.Sandberg@genpat.uu.se



UPPSALA
UNIVERSITET

XPR