

# scioDiscover

highly-parallel protein analysis

Biomarker discovery

Immuno-oncology


Stem cell analysis


Apoptosis and cell cycle analysis

Transcription factor analysis

Organ failure

Predictive toxicology








## Advantages

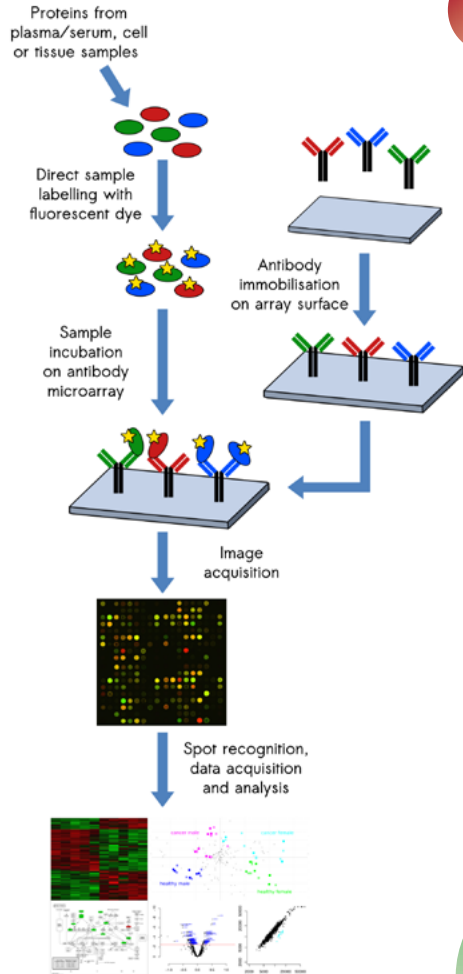
- > High content analysis of 900 proteins
- > Broad spectrum covered such as oncology and organ failure
- > Low sample volumes required (10 µl plasma/serum)
- > Profiling of native samples (non-fractionated/non-depleted)
- > Sensitivity comparable to ELISA assays
- > Fully immuno-based assay (easy translation into diagnostic assays)
- > Reliable dual-colour study design
- > High reproducibility (CV < 10%)
- > Complete sample-to-result service including full data analysis and study report

## Features

- > Parallel analysis with 1130 antibodies
  - > Broad coverage of:
    - Signalling pathways
    - Transcription factors
    - Apoptosis markers
    - Oxidative stress response
    - Cell surface molecules
    - Cytokines & chemokines
  - > Several sample types possible:
    - Plasma/serum
    - Tissue samples
    - Cell culture pellets
    - Cell culture supernatant
    - Cerebrospinal fluid (CSF)
    - Additional sample types on request
  - > Studies possible in human, murine and rat samples
  - > PTM analysis on request
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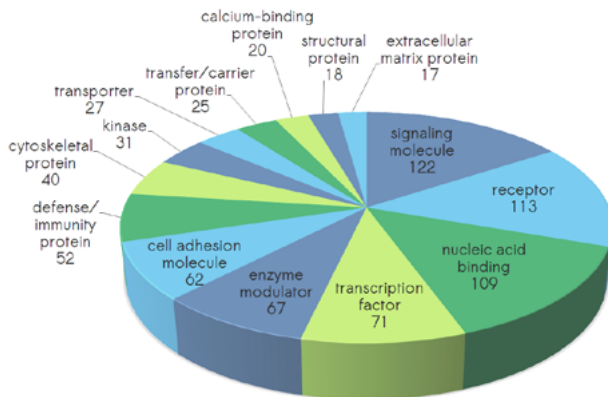
## Applications

- > Biomarker discovery
- > Drug target discovery
- > Cancer pathway screening
- > Signalling pathway profiling
- > Immune activation profiling
- > Disease mechanism profiling
- > Transcription factor analysis
- > Apoptosis and cell cycle analysis
- > Stem cell analysis
- > Ageing research
- > Regenerative medicine

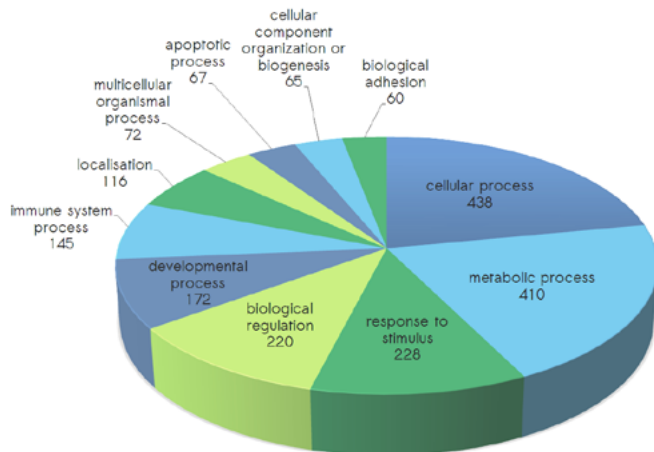




## Protein classes

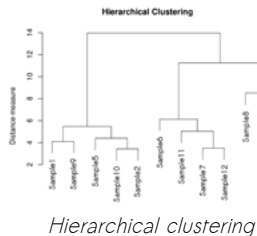


## Biological processes

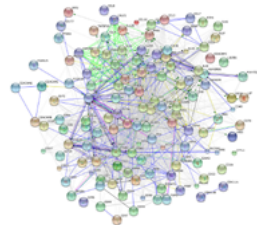


# scioDiscover sample-to-result service

- > Definition of study design
- > Sample preparation
- > Protein extraction
- > Protein concentration measurement
- > Protein quality control
- > Sample labelling
- > Incubation of the samples on scioDiscover antibody microarrays
- > Microarray scanning
- > Raw data acquisition
- > Data normalisation
- > Data analysis including hierarchical clustering analysis
- > Statistical testing for differentially abundant proteins
- > Additional analysis types available on request
- > Comprehensive study report



*Protein interaction analysis*



*Heat map of differentially expressed proteins*



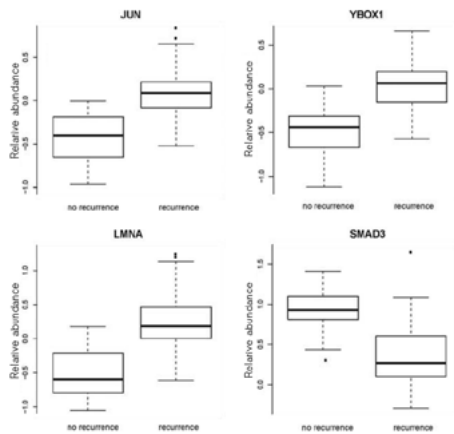
*Principle component analysis*



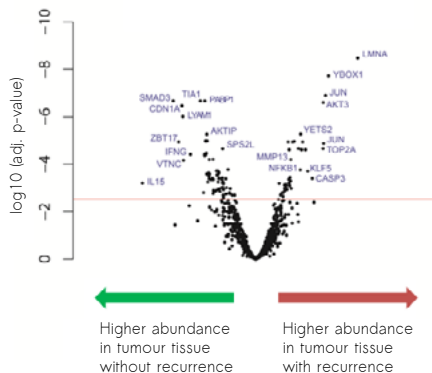
# Case study: Recurrence of bladder cancer

Comparison of protein expression profiles in bladder tumour tissues with and without recurrence in the clinical course.

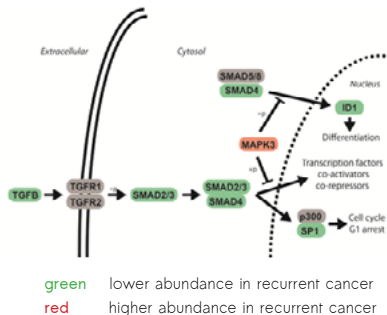
Highly relevant proteins were identified which are specific to the individual tumour subtypes.



Four proteins show a clear differentiation of recurrent and non-recurrent tumours.



## TGFB pathway



## What our customers say

*„The Sciomics protein profiling platform [...] has undoubtedly opened new avenues in our research. We have found the Sciomics team to be extremely helpful and efficient during the whole process, from advising the best strategy to the final interpretation of results.“*

Dr. Leonardo Guasti, Barts and The London, Queen Mary's School of Medicine and Dentistry, London, United Kingdom

*„I was very satisfied by the service offered by Sciomics. Conducting a project with Sciomics did feel like dealing with a partner who pays attention to the smallest details and is very interested in the success of the project.“*

Dr. Gaith Bakdash, Department of Tumor Immunology, RadboudUMC, Nijmegen, The Netherlands

*„Sciomics antibody microarray really helped us to move forward. I am more than pleased to recommend Sciomics Antibody Microarray analysis service. The results were sent fast and ready to publish! If I hit a road block in my experiments, I am sure that their services will help me go through.“*

Dr. José Pedro Castro, German Institute of Human Nutrition (DIfE), Potsdam-Rehbrücke, Germany

## Additional Sciomics products

scioCD	Cell surface marker profiling (available as microarray kit and as a full service)
scioPhospho	Phosphorylation level analysis
scioExtract	Protein isolation buffer for tissue and cell samples

**Sciomics GmbH**  
Im Neuenheimer Feld 583  
D-69120 Heidelberg  
Germany

info@sciomics.de  
www.sciomics.de

T +49 6221 4294830