

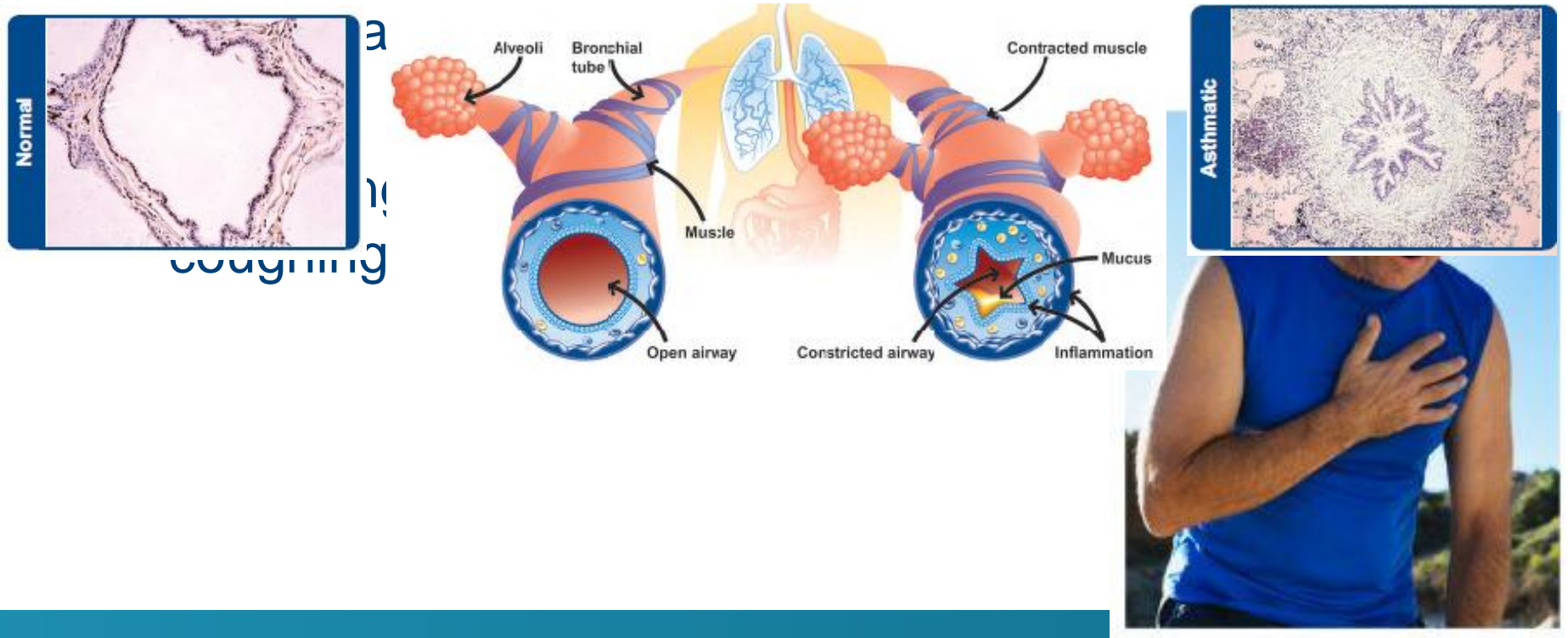


Novel biomarkers of chemical-induced asthma: a murine model

Jeroen Vanoirbeek

Asthma

1. Chronic airway disease:
prevalence: 5-10%, 300.000 people affected world-wide
2. Reversible airflow limitation, airway hyperreactivity &



Asthma

1. Rapid increase of allergy and asthma in past 40 years
2. Changes in gene pool unlikely
3. Changes in “environment”?
 - Hygiene, lifestyle, diet, ...
 - Air pollution (indoor, outdoor)
 - Specific chemicals?
4. Mechanisms?

Occupational asthma

1. Definition:

- occupational asthma is a disease characterized by variable airflow limitation and/or airway hyperresponsiveness due to causes and conditions attributable to a particular occupational environment and not to stimuli encountered outside the workplace

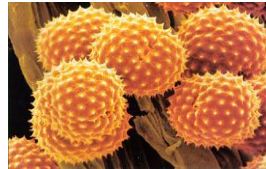
2. 9-15% of all adult asthma is due to exposures on the work floor

3. The most common cause of work-related lung diseases

Occupational agents

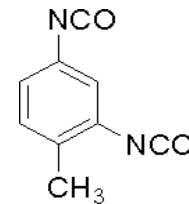
High molecular weight (HMW) compounds (≥ 5 kDa)

- Animal proteins
- Plant proteins
- Enzymes



Low molecular weight (LMW) compounds (< 5 kDa)

- Chemicals
- Metals
- Wood dust
- Pharmaca



Immune
sensitization

No immune
sensitization

IgE mediated

*IgE mediated or
non-IgE mediated*

Non-IgE mediated

HMW allergens

Flour, Lab animals,...

LMW sensitizers

Isocyanates, persulphates,...

LMW irritants

Chlorine, Ammonia,...

Occupational agents

- Diisocyanates
 - Highly reactive, low molecular weight compounds
 - Most common cause of chemical-induced occupational asthma
 - Used for the production of polyurethanes, foams, paints, etc



Skin - lung

- Development of chemical-induced asthma
 - Primary route of exposure and initiation of immune response = respiratory tract
 - Regulation and prevention of OA almost exclusively focusses on airborne exposures
 - Despite reduction in workplace respiratory exposure, diisocyanate asthma continues to occur

Focus on skin exposure

Occupational asthma

Mechanisms ?

- Role of chemical properties ?
 - chemical reactivity
 - irritant properties
- Pathways of sensitization ?
 - via dermal route ?
- Immunological mechanisms ?
 - usually no specific IgE antibodies
 - cellular mechanisms

Implications

- Hazard identification
 - Prediction of asthmogens
- Prevention
 - Avoid skin contact
- Surveillance & diagnosis
 - Identification of sensitized subjects
 - Identification of sensitizing agent in affected subjects

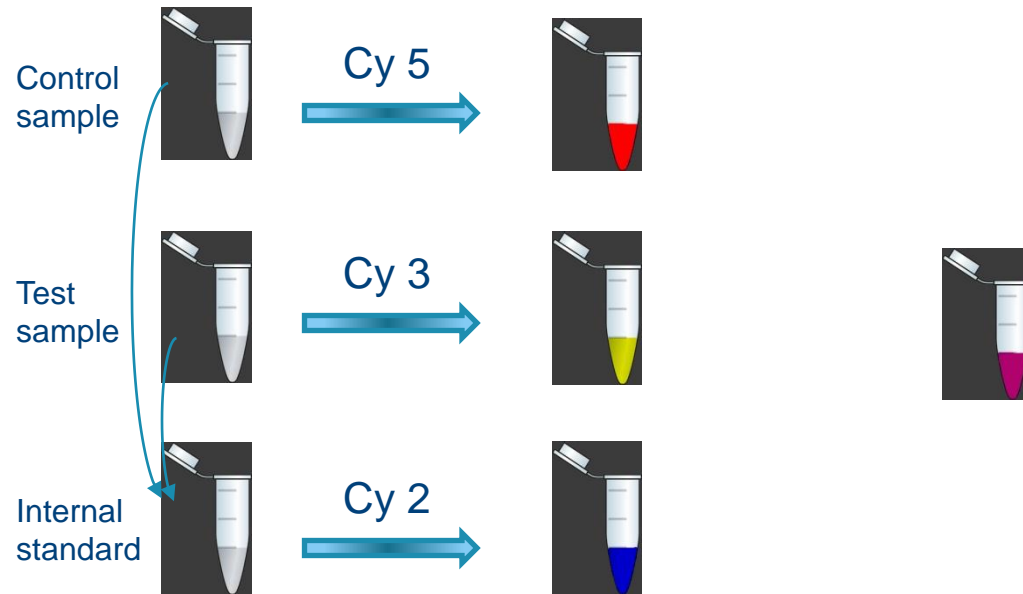
Aim

To identify (early) biomarkers of chemical-induced asthma and sensitization to chemicals using proteomics techniques

Proteomics

- Proteomics is a global strategy in which all proteins (the proteome) derived from a cell, tissue, body liquid or whole organism, are simultaneously visualized and identified

Two-dimensional difference gel electrophoresis: 2D-DIGE



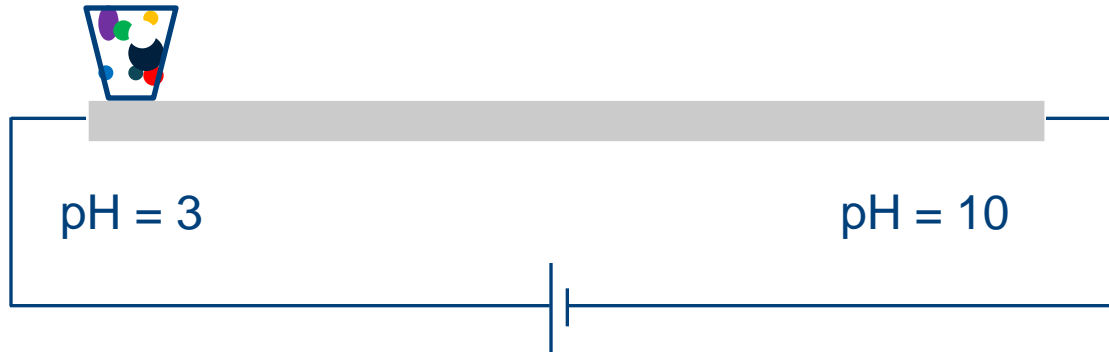
Two-dimensional difference gel electrophoresis: 2D-DIGE



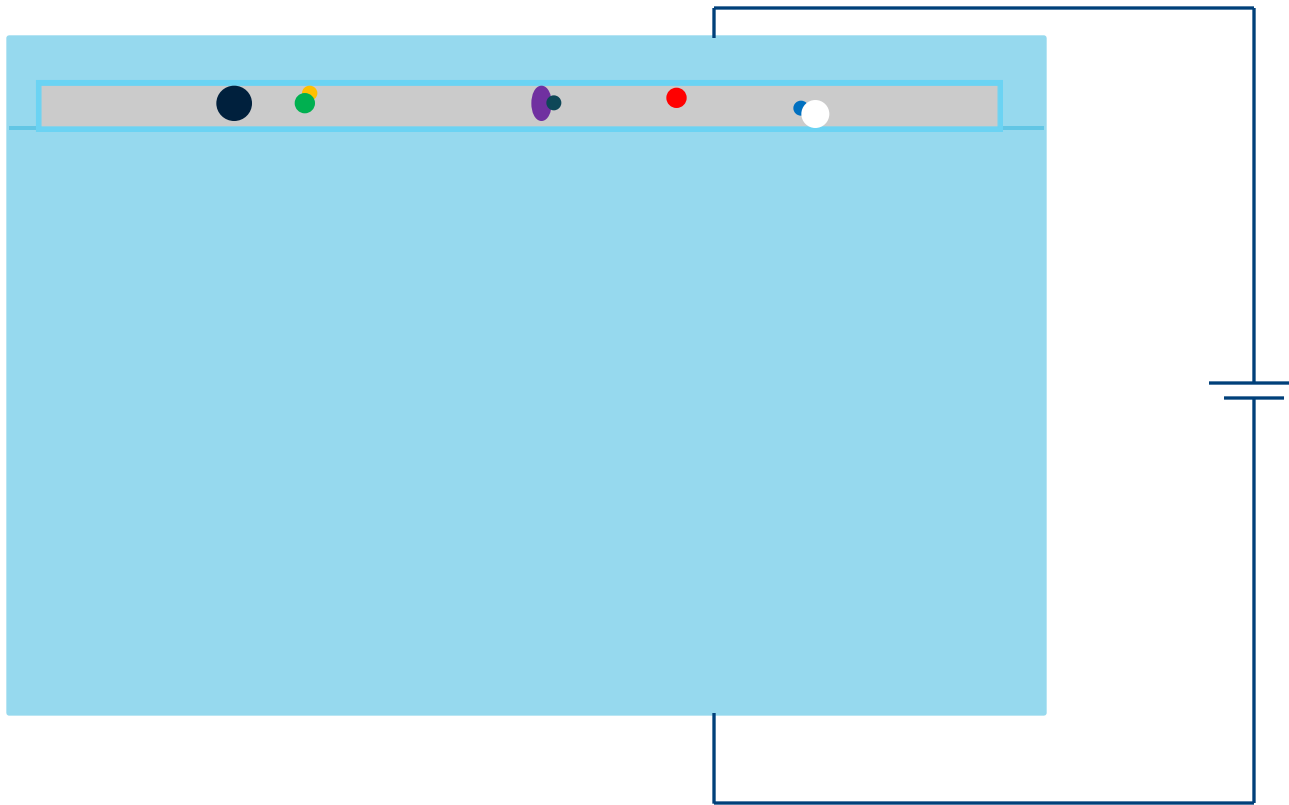
pH = 3

pH = 10

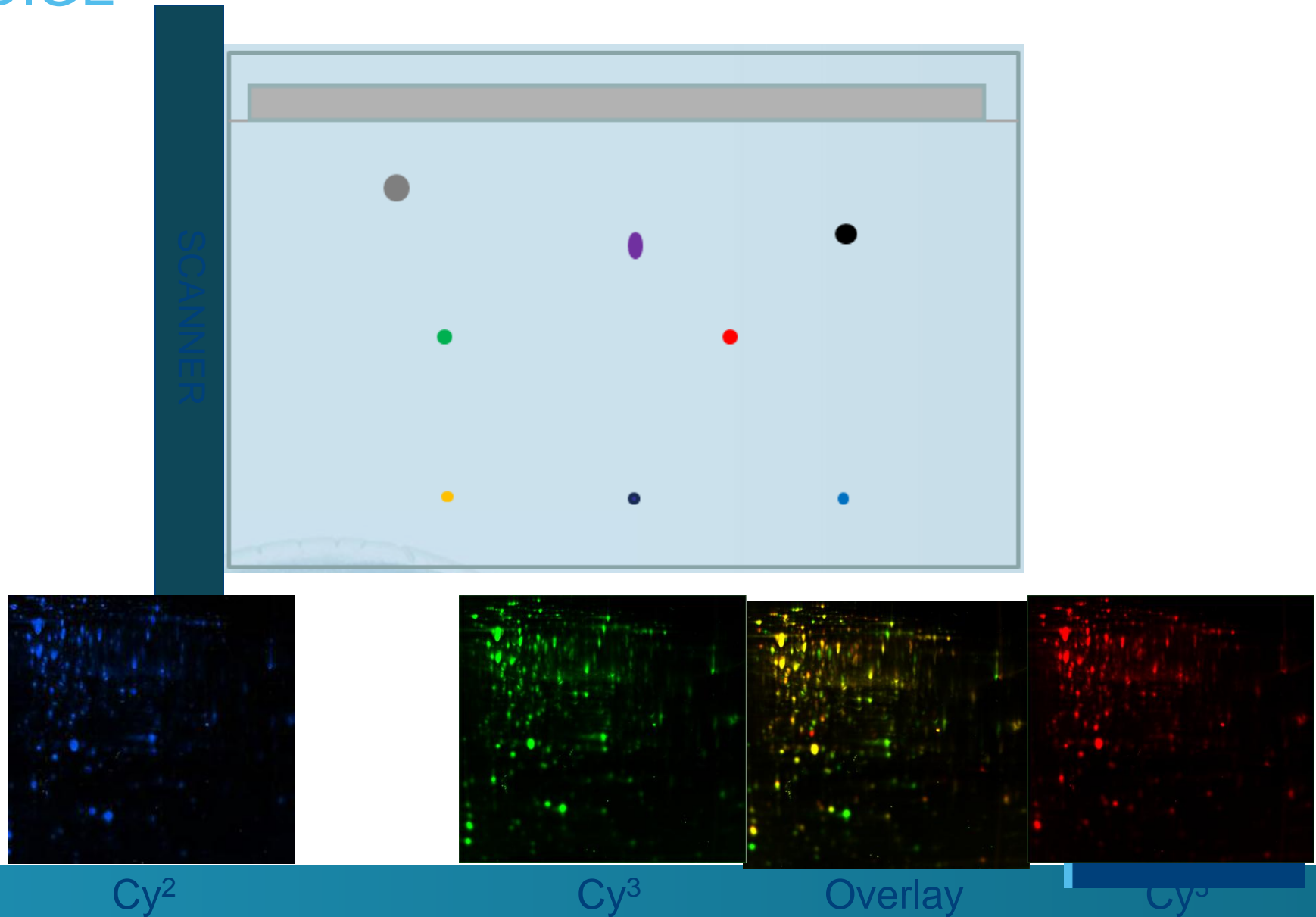
Two-dimensional difference gel electrophoresis: 2D-DIGE



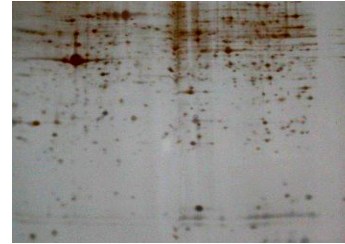
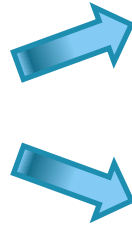
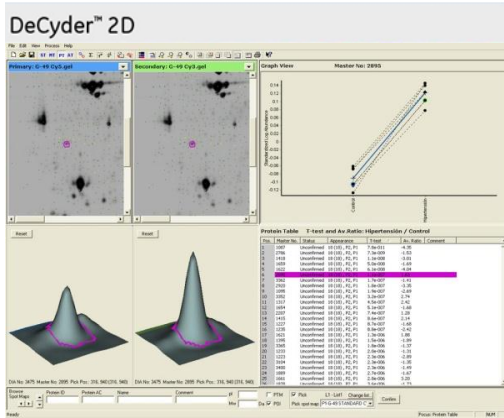
Two-dimensional difference gel electrophoresis: 2D-DIGE



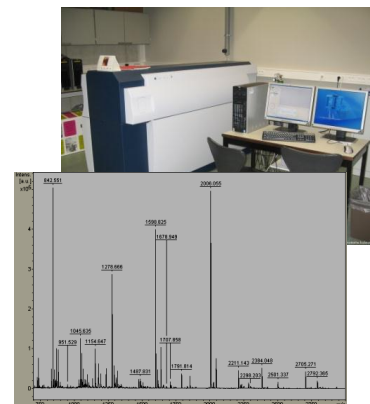
Two-dimensional difference gel electrophoresis: 2D-DIGE



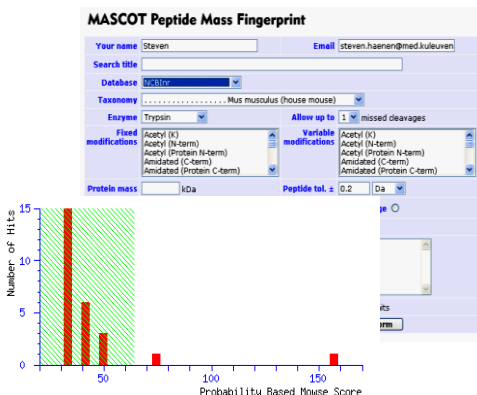
Two-dimensional difference gel electrophoresis: 2D-DIGE



Trypsin Digest



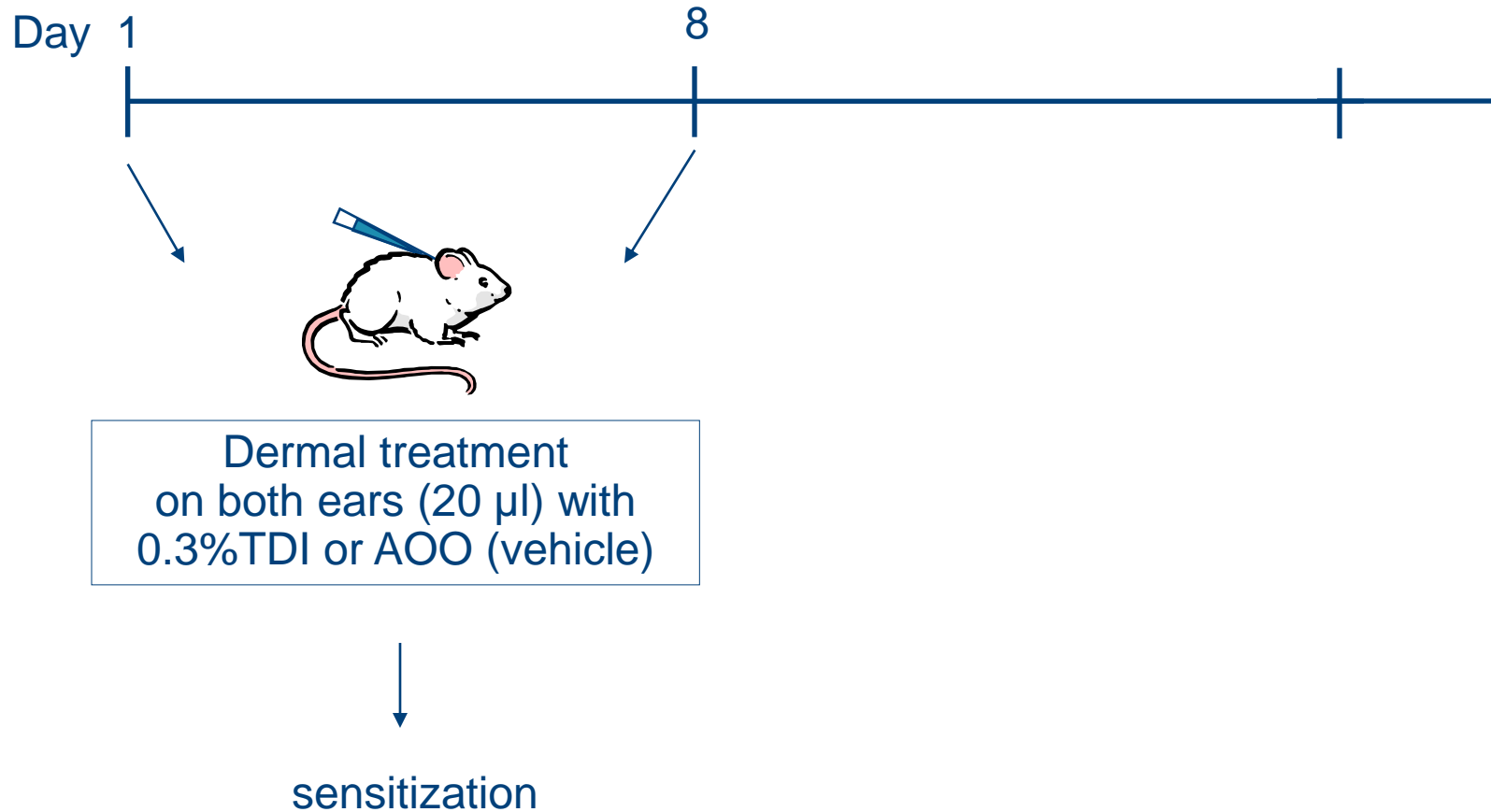
MALDI-TOF MS



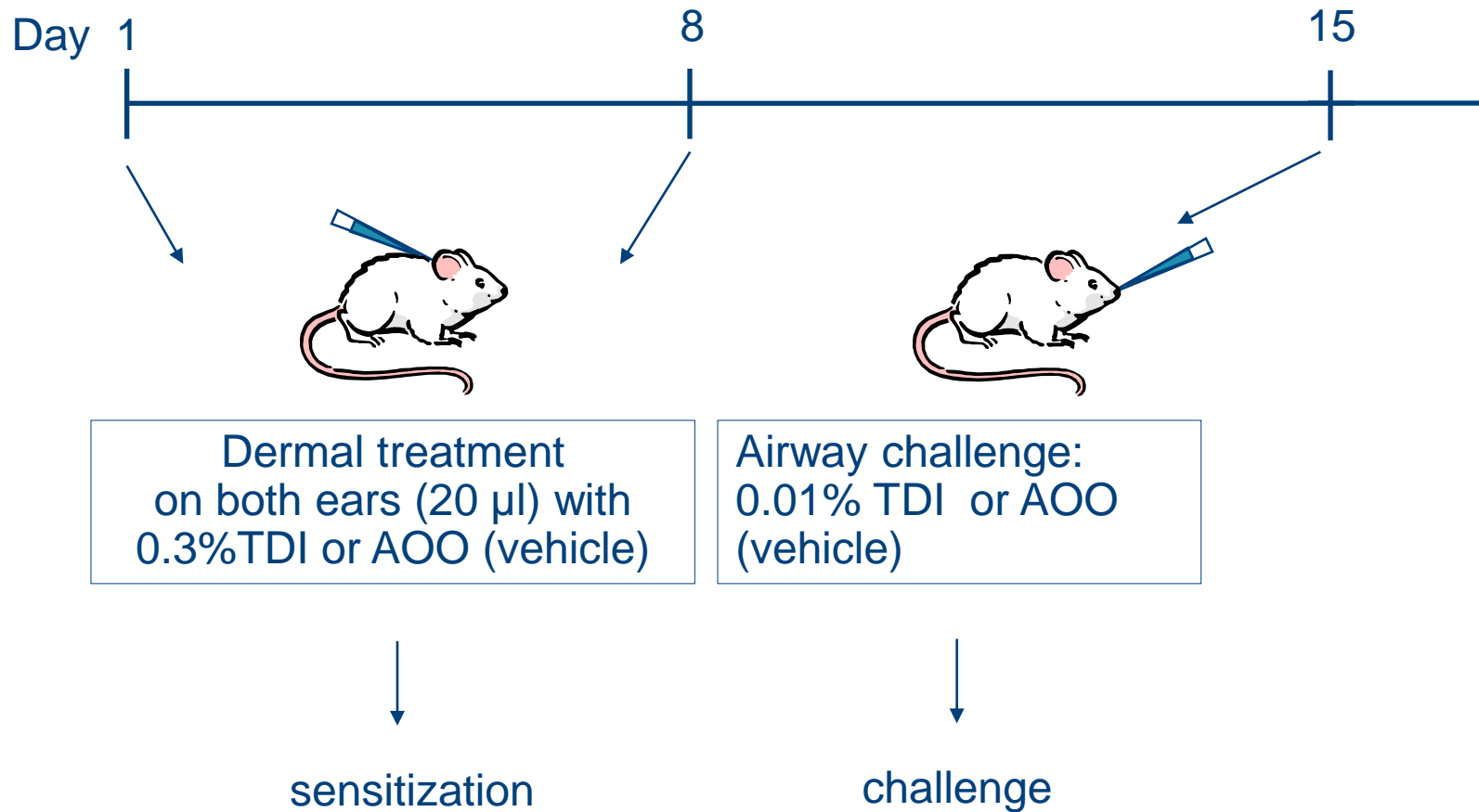
Proteomics in chemical-induced asthma

- Classical research: focus on immune related cells and cytokines, resulting in many insights, but exact mechanisms of OA still unclear
- Development of new & sensitive methods → new approaches

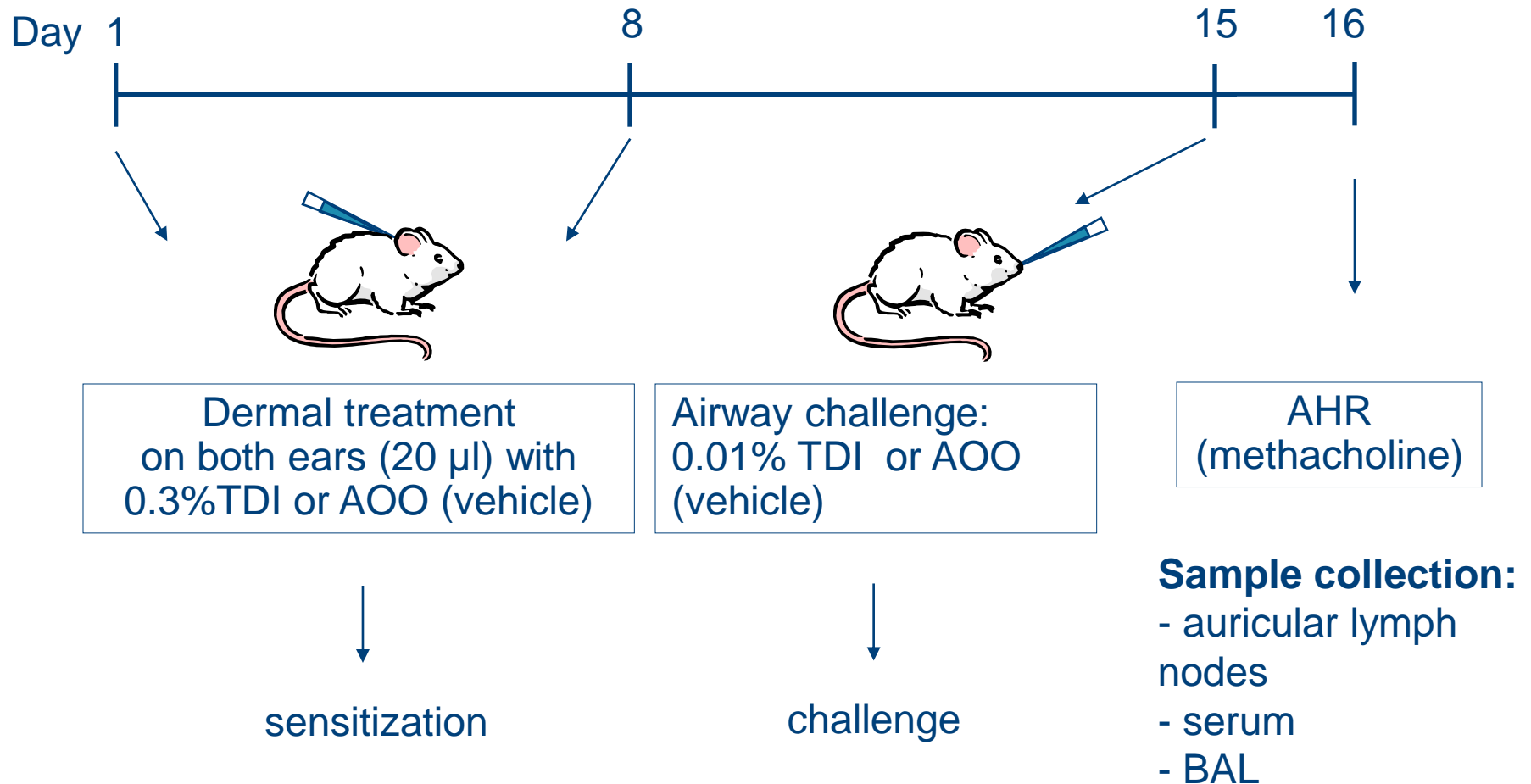
Proteomics in chemical-induced asthma



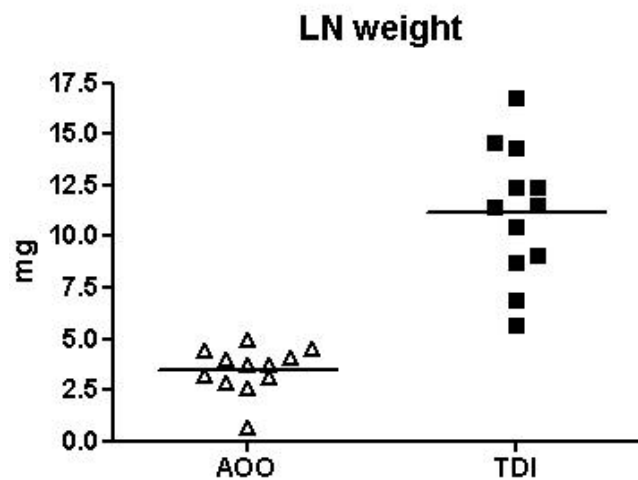
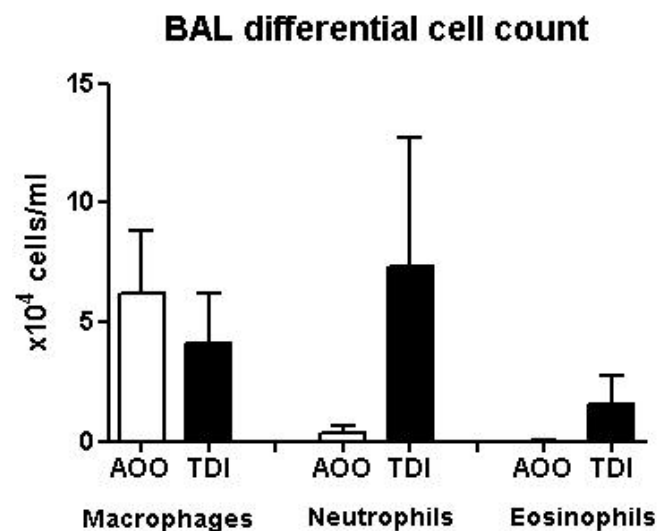
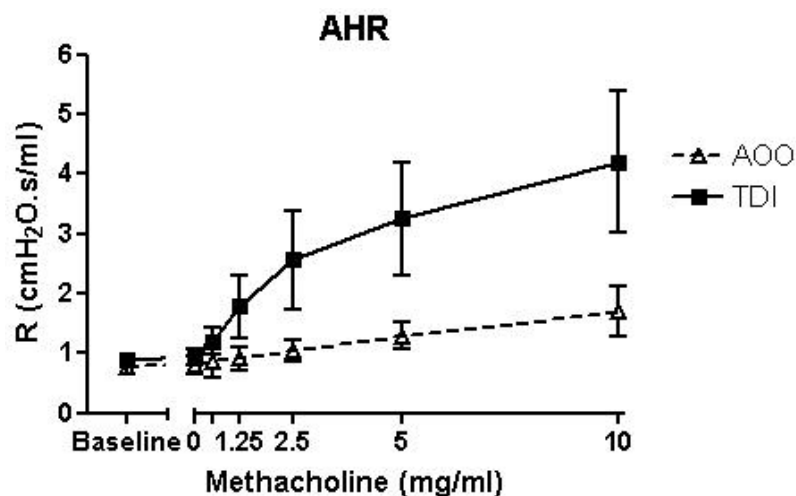
Proteomics in chemical-induced asthma



Proteomics in chemical-induced asthma

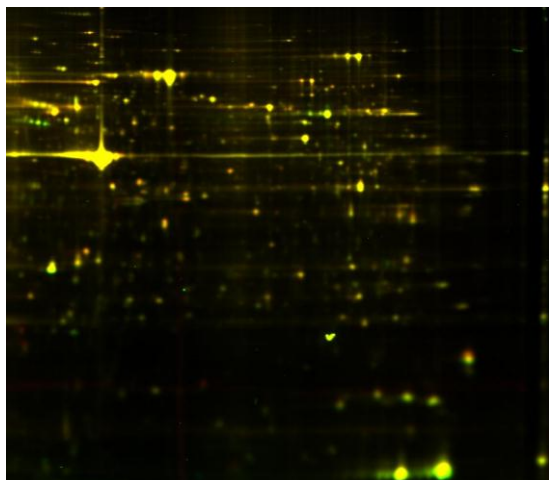


Proteomics in chemical-induced asthma



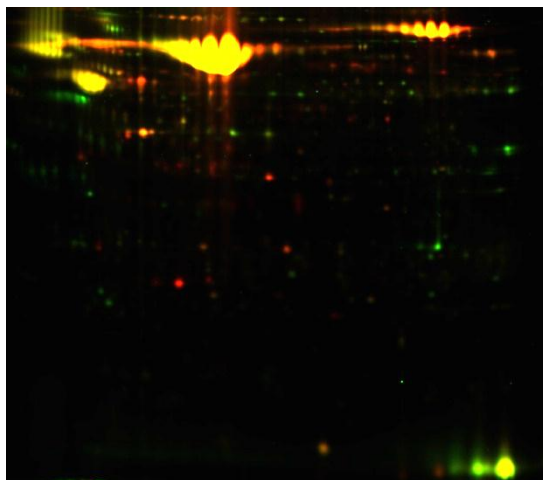
Proteomics in chemical-induced asthma

Auricular lymph nodes



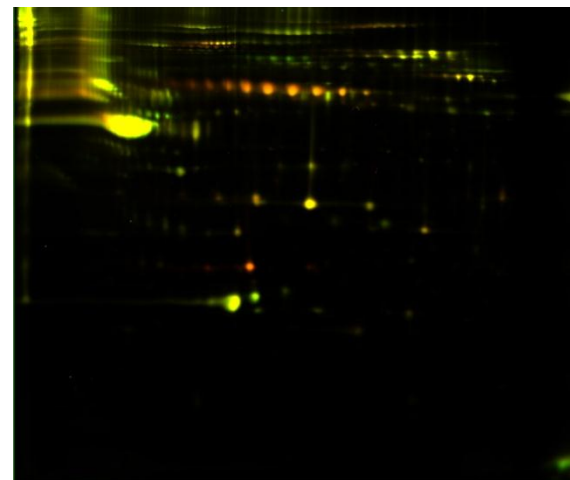
$p < 0.01$	
Diff. Spots	53
Identified	27
Up	14
Down	13

BAL



$p < 0.01$	
Diff. Spots	210
Identified	72
Up	55
Down	17

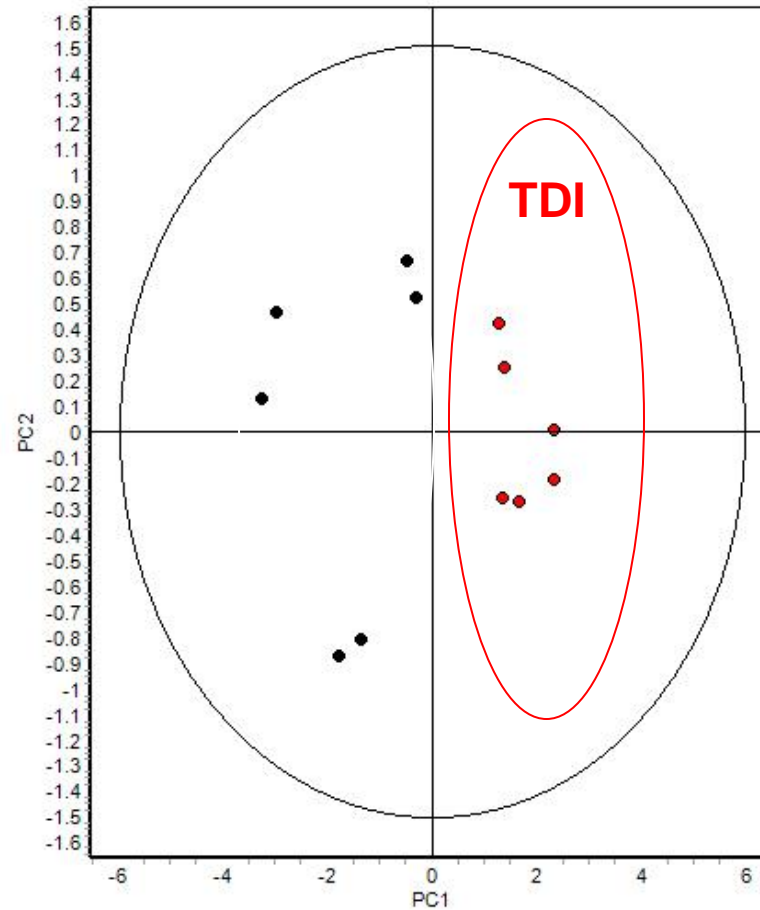
Serum



$p < 0.01$	
Diff. Spots	40
Identified	18
Up	9
Down	9

Proteomics in chemical-induced asthma

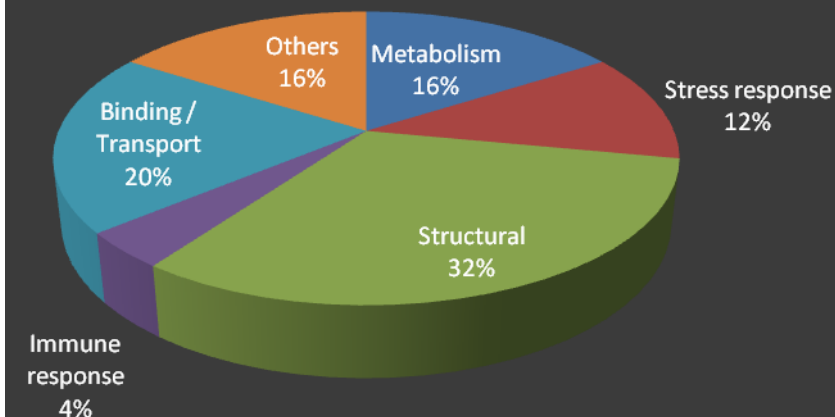
Spot Maps (Score Plot)



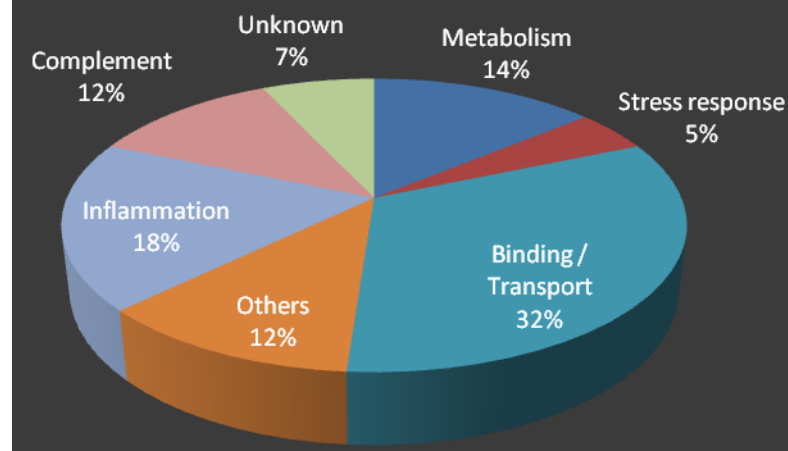
Auricular lymph nodes

Proteomics in chemical-induced asthma

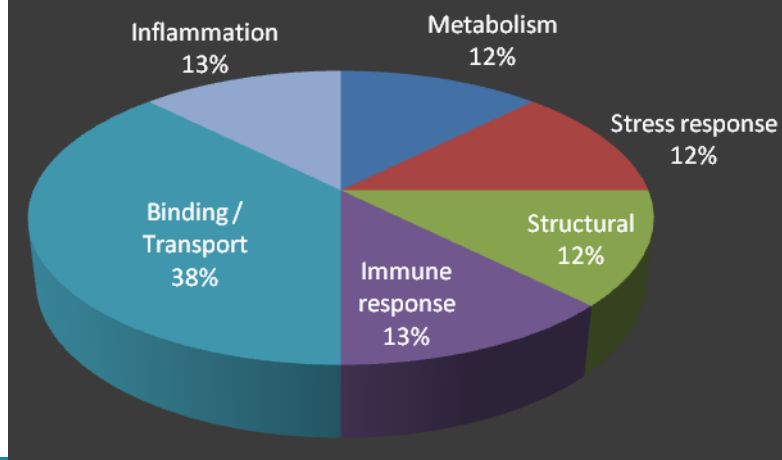
auricular lymph nodes



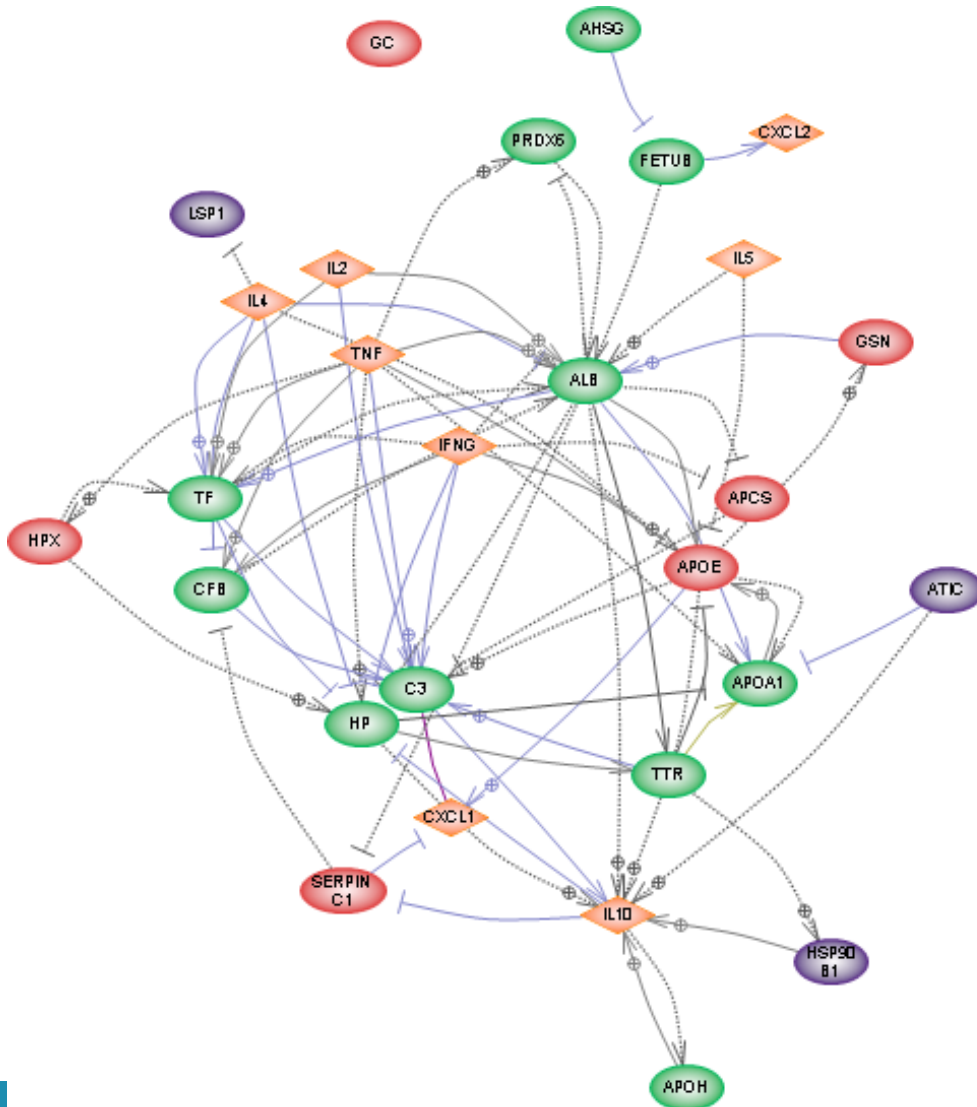
BAL




serum




Proteomics in chemical-induced asthma



auricular lymph nodes

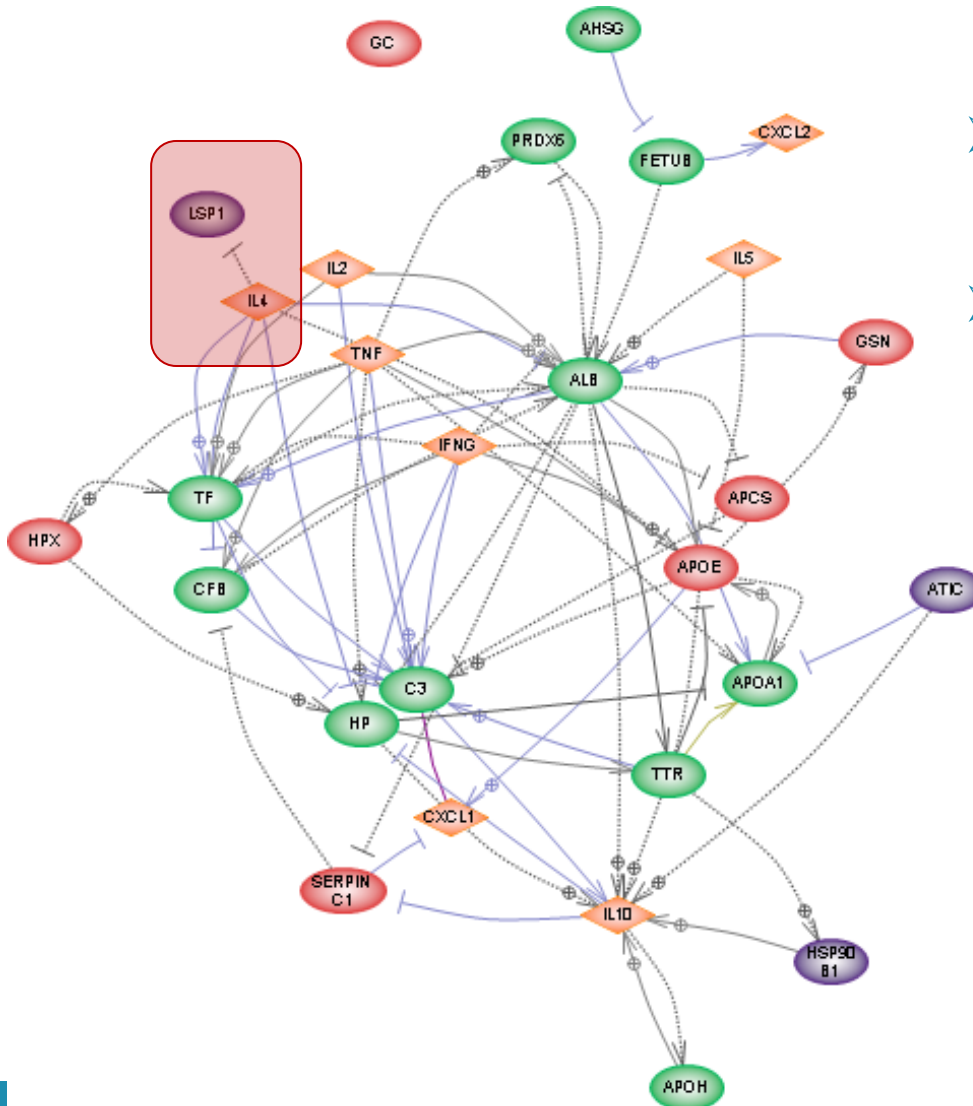


bronchoalveolar lavage



serum

Proteomics in chemical-induced asthma



➤ Lymphocyte specific protein-1 ↓

➤ Literature:

➤ Expression:

- lymphocytes
- neutrophils
- macrophages

➤ Chemotaxis & activation neutro

➤ Overexpression → dysfunction

➤ Regulated by IL-4

Conclusions

1. First systematic & systemic proteomics approach in a model of occupational asthma
2. Dermal sensitization and a single airway challenge leads to profound proteome changes in multiple compartments
3. Physiological and immunological changes are reflected by changes in the proteome

Proteome changes in auricular lymph nodes and serum after dermal sensitization to toluene diisocyanate in mice

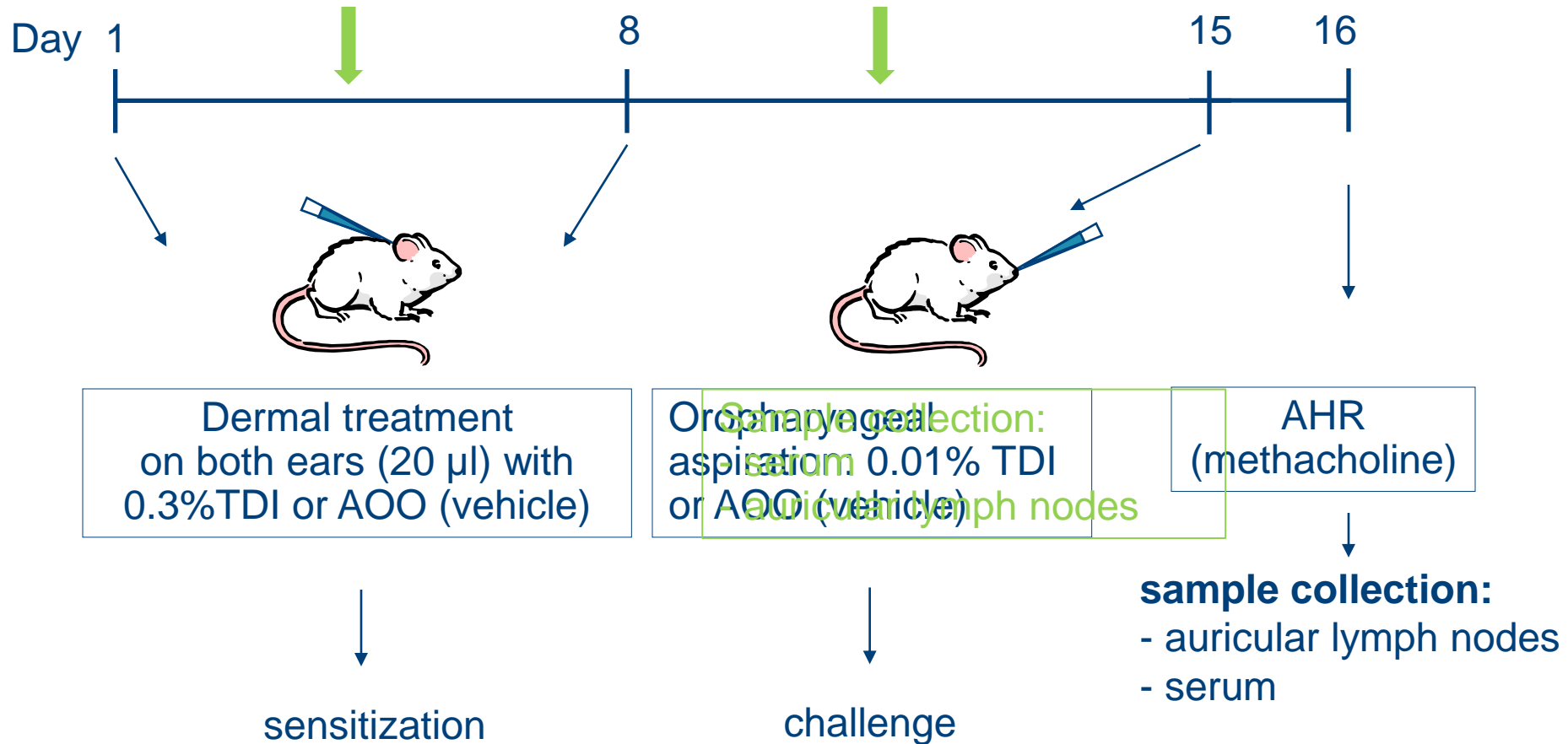
Proteome changes after dermal sensitization

1. So far, no study has focused on early time points
2. Human (early) biomarker research is complicated: only when disease has established

Proteome changes after dermal sensitization

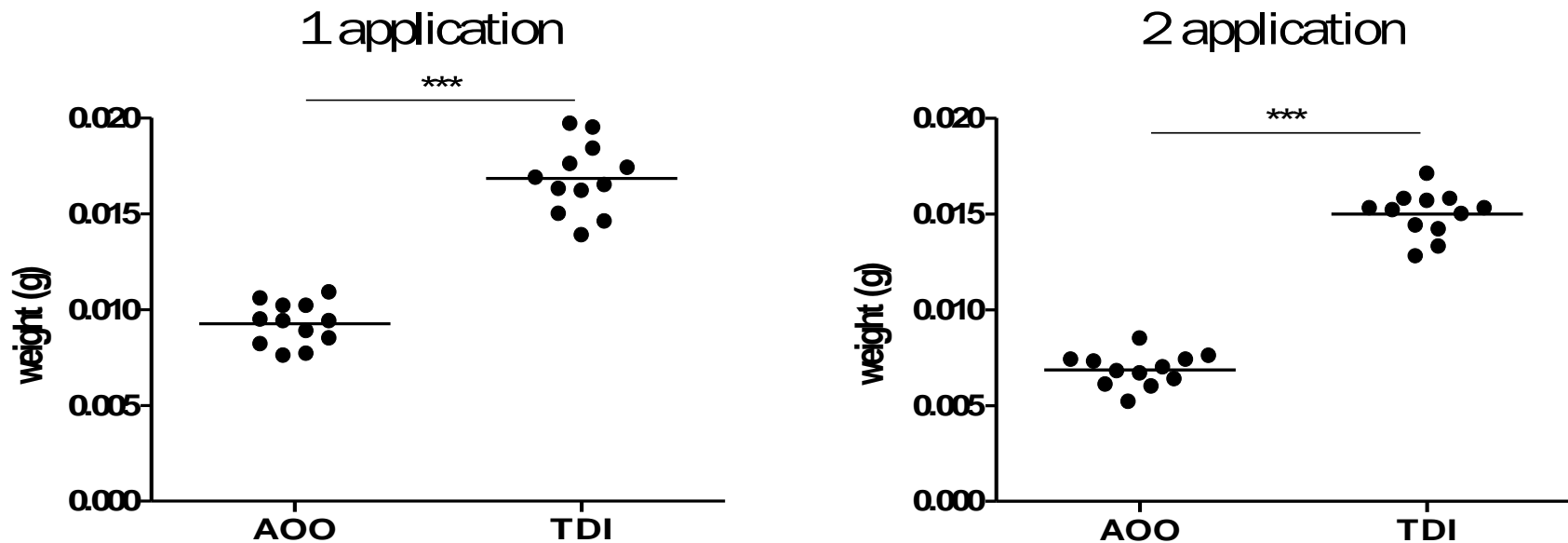
1. So far, no study has focused on early time points
2. Human (early) biomarker research is complicated: only when disease has established
3. Rationale: investigate changes in the proteome during sensitization to identify (early) markers of sensitization

Proteome changes after dermal sensitization



Proteome changes after dermal sensitization

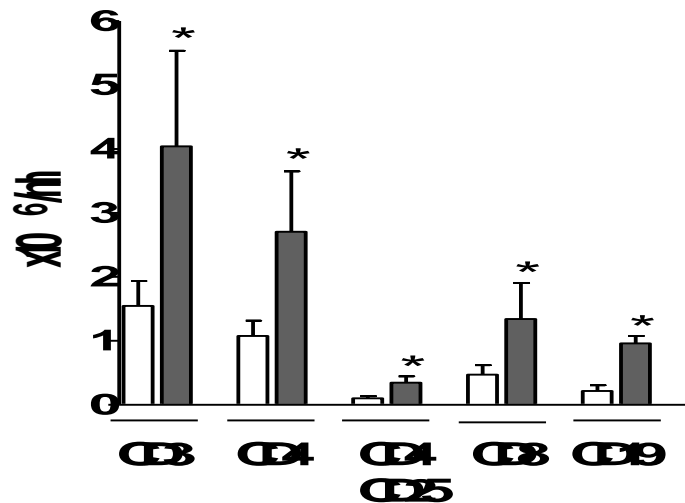
Auricular lymph node weight



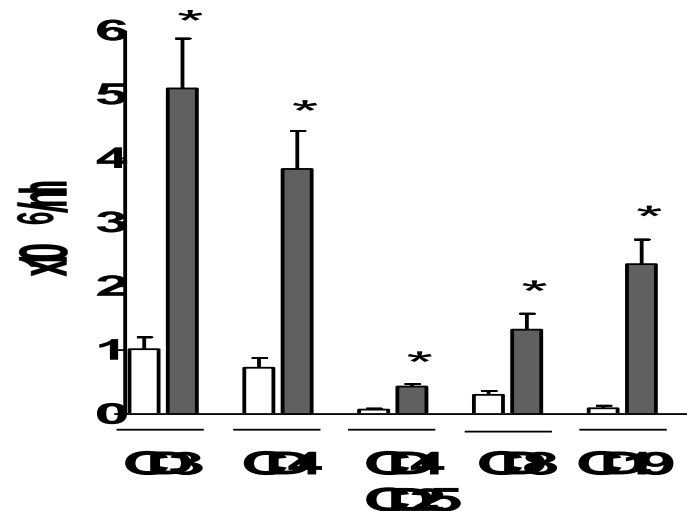
Proteome changes after dermal sensitization

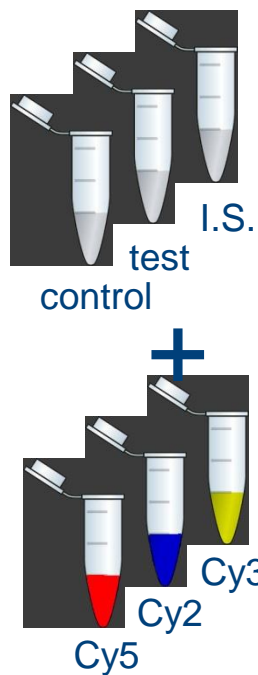
Lymphocyte subpopulations

1 application



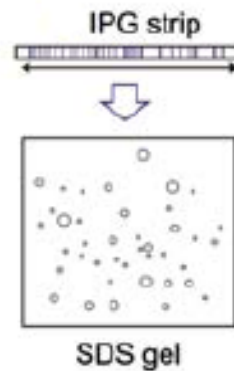
2 applications



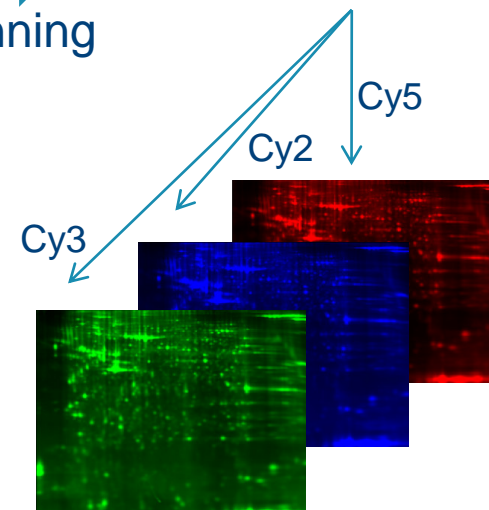


①
labeling

②
separation

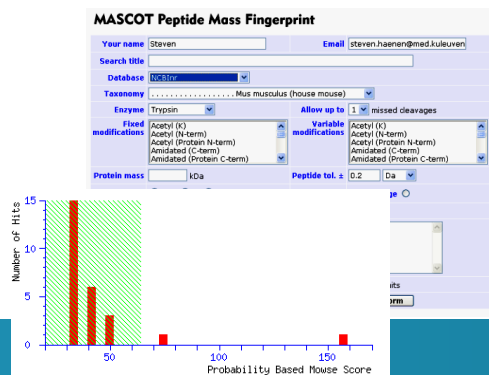


③
scanning

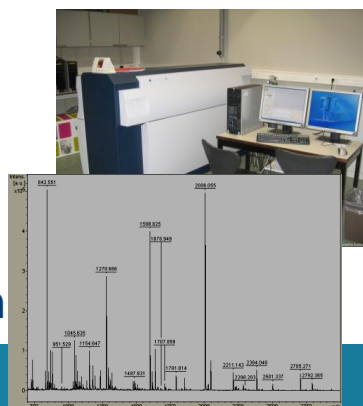


2D-DIGE workflow

Decyder analysis ↓ ④

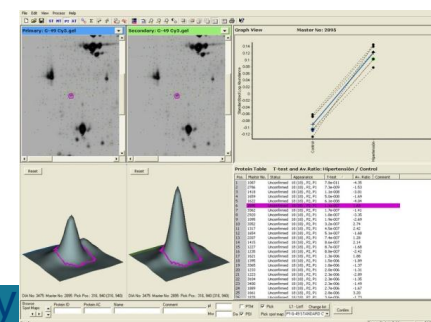


⑥
identification



⑤
mass

spectrometry

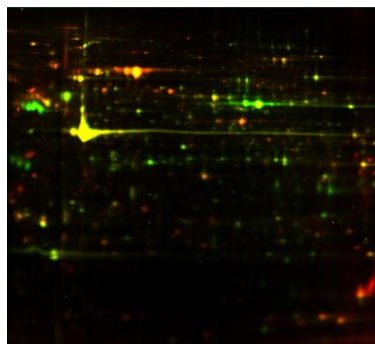
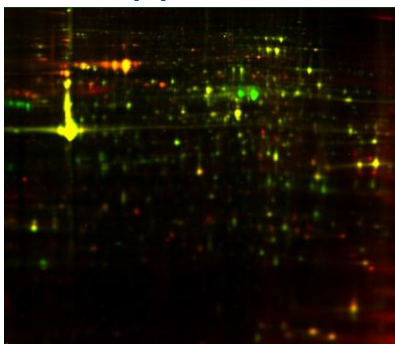


Proteome changes after dermal sensitization

auricular lymph nodes

1 application

2 applications



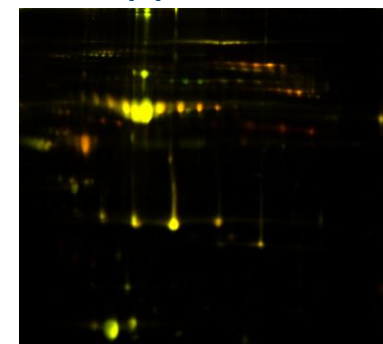
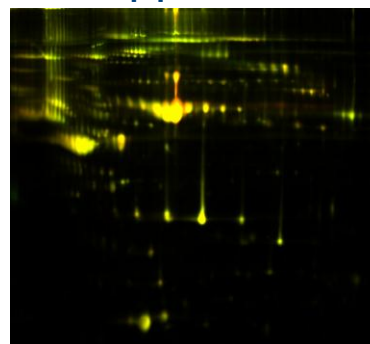
$p < 0.01$	
Diff. spots	38
Identified	26
Up	15
Down	11

$p < 0.01$	
Diff. spots	58
Identified	35
Up	19
Down	16

serum

1 application

2 applications



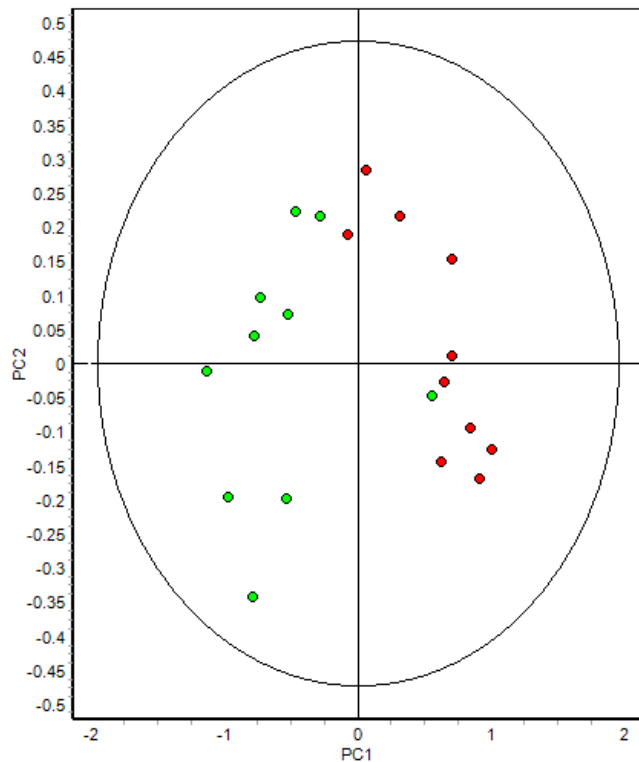
$p < 0.01$	
Diff. spots	7
Identified	3
Up	3
Down	0

$p < 0.01$	
Diff. spots	16
Identified	10
Up	4
Down	6

Proteome changes after dermal sensitization

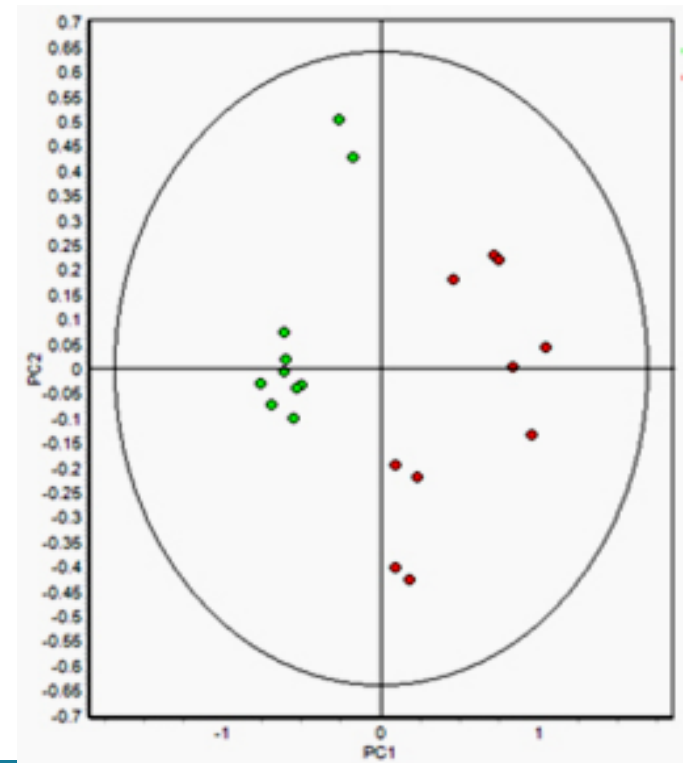
auricular lymph nodes

1 application



● AOO
● TDI

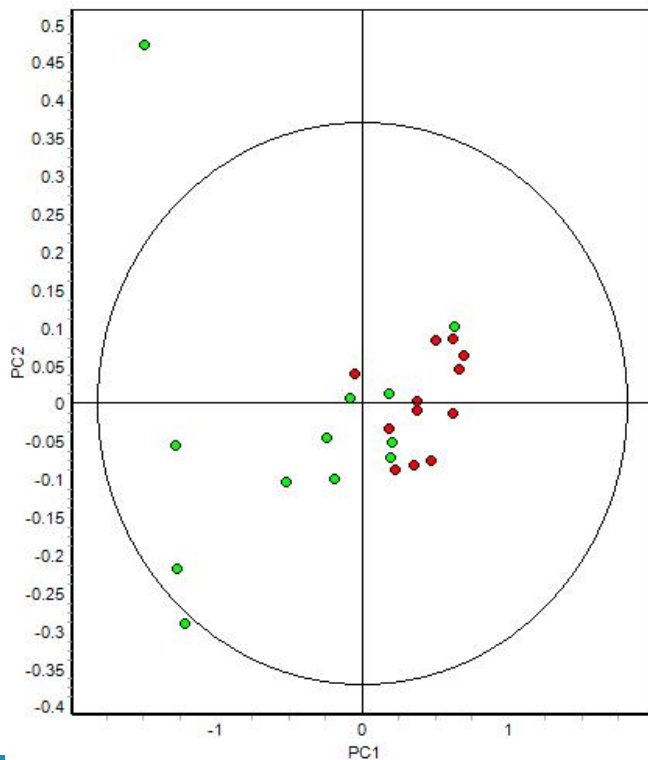
2 applications



Proteome changes after dermal sensitization

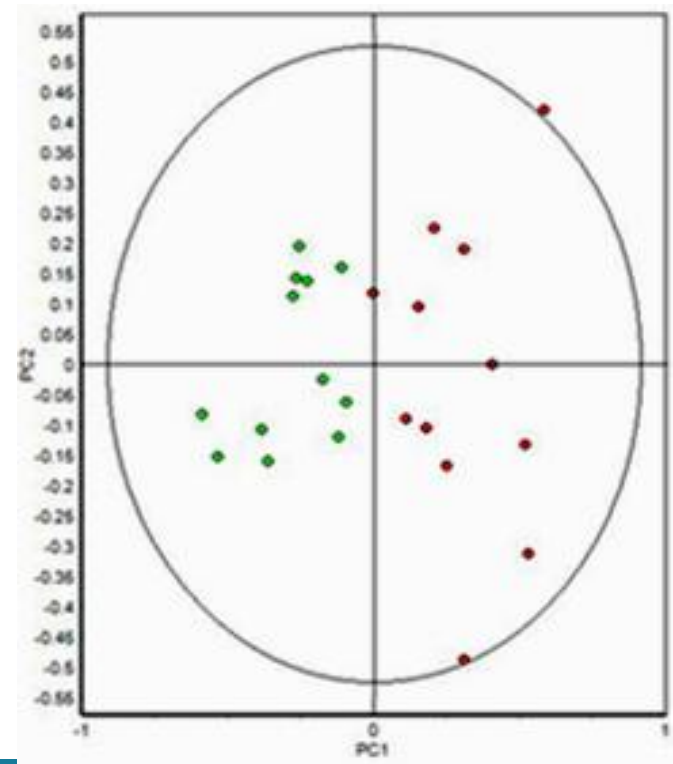
serum

1 application



● AOO
● TDI

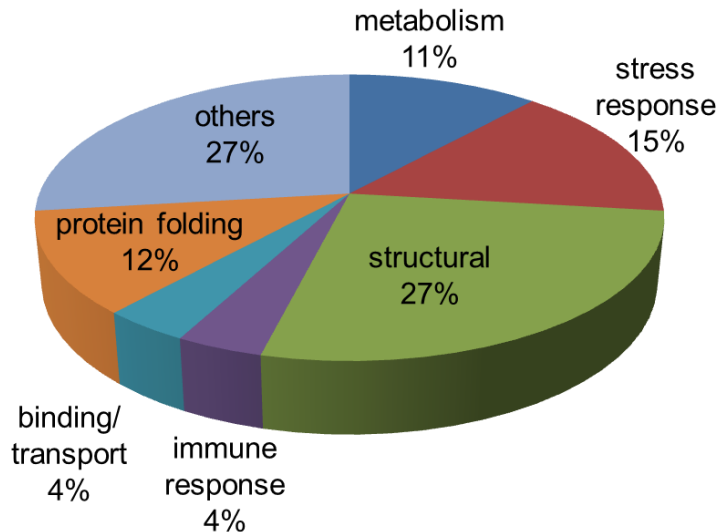
2 applications



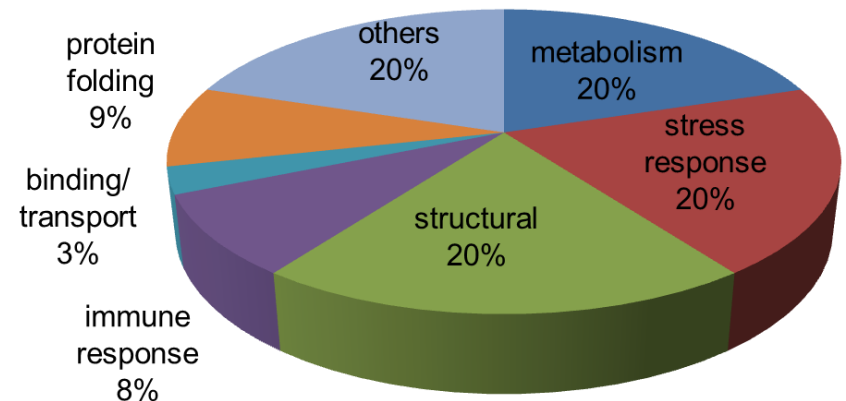
Proteome changes after dermal sensitization

auricular lymph nodes

1 application



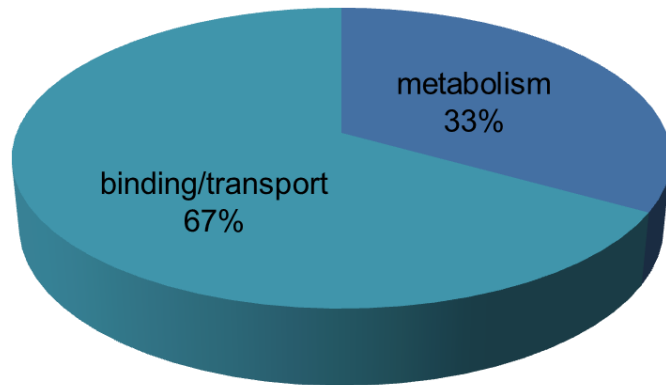
2 applications



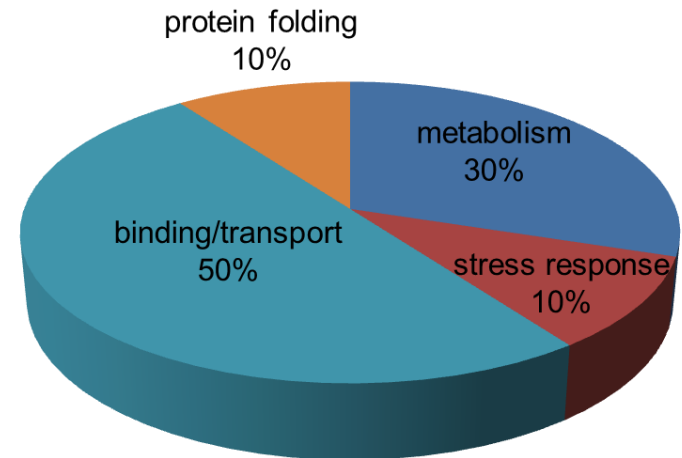
Proteome changes after dermal sensitization

serum

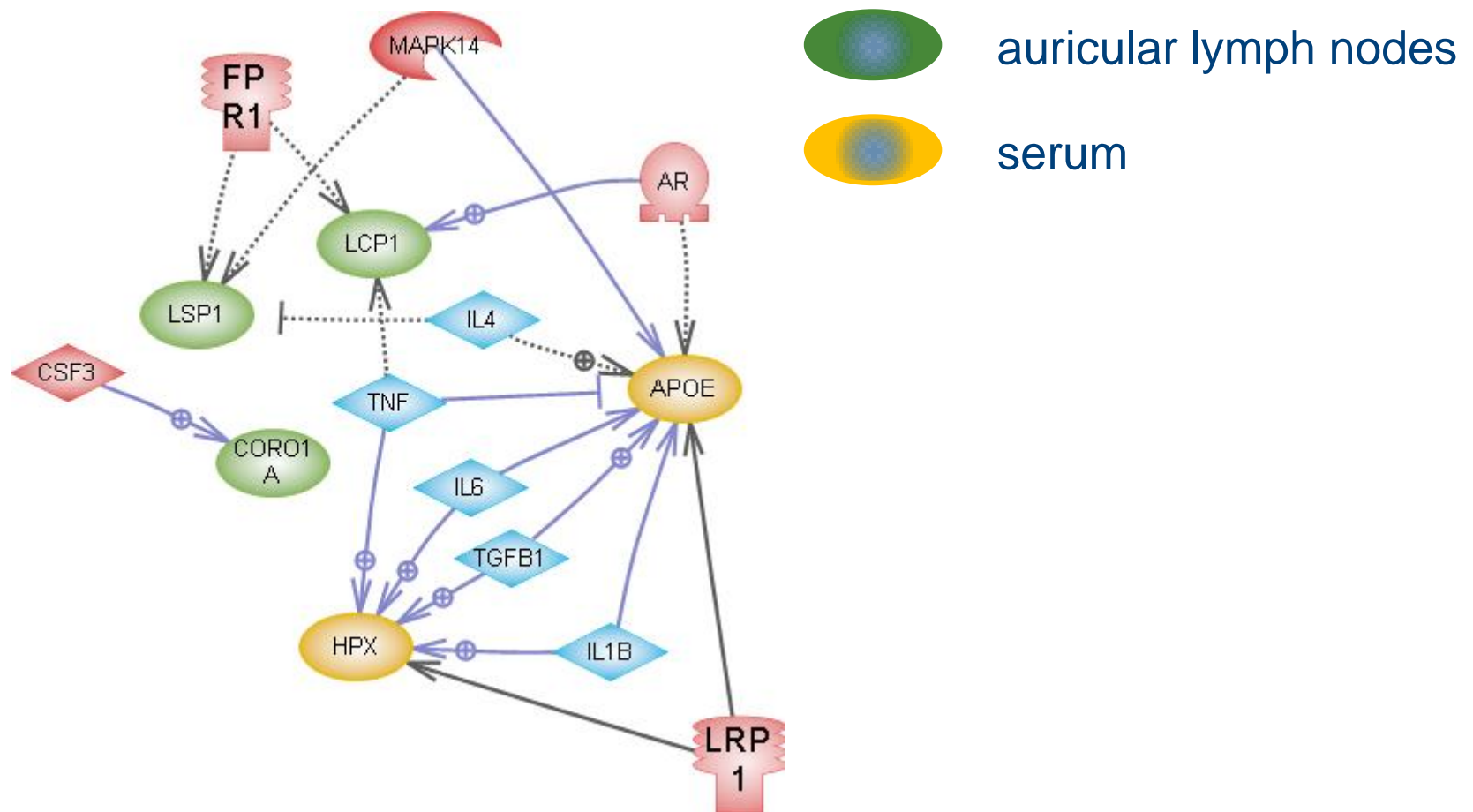
1 application



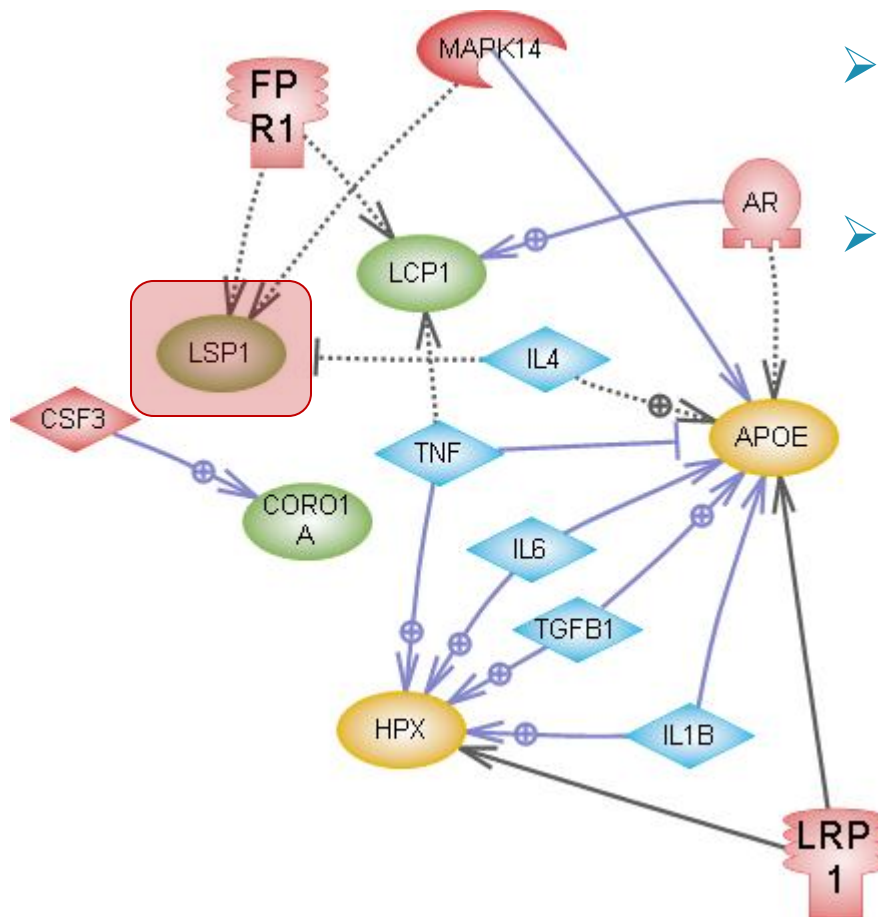
2 applications



Proteome changes after dermal sensitization



Proteome changes after dermal sensitization



➤ Lymphocyte specific protein-1

➤ Literature:

➤ Expression:

- lymphocytes
- neutrophils
- macrophages

➤ Chemotaxis & activation neutro

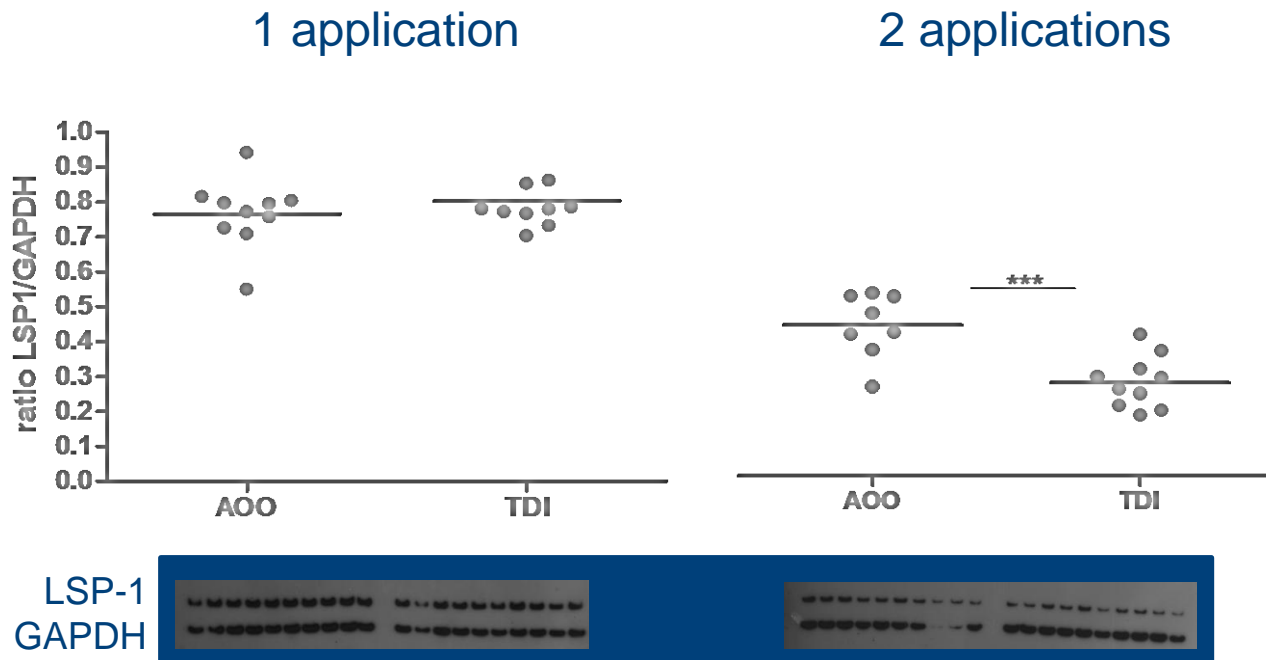
➤ Overexpression → dysfunction

➤ Regulated by IL-4

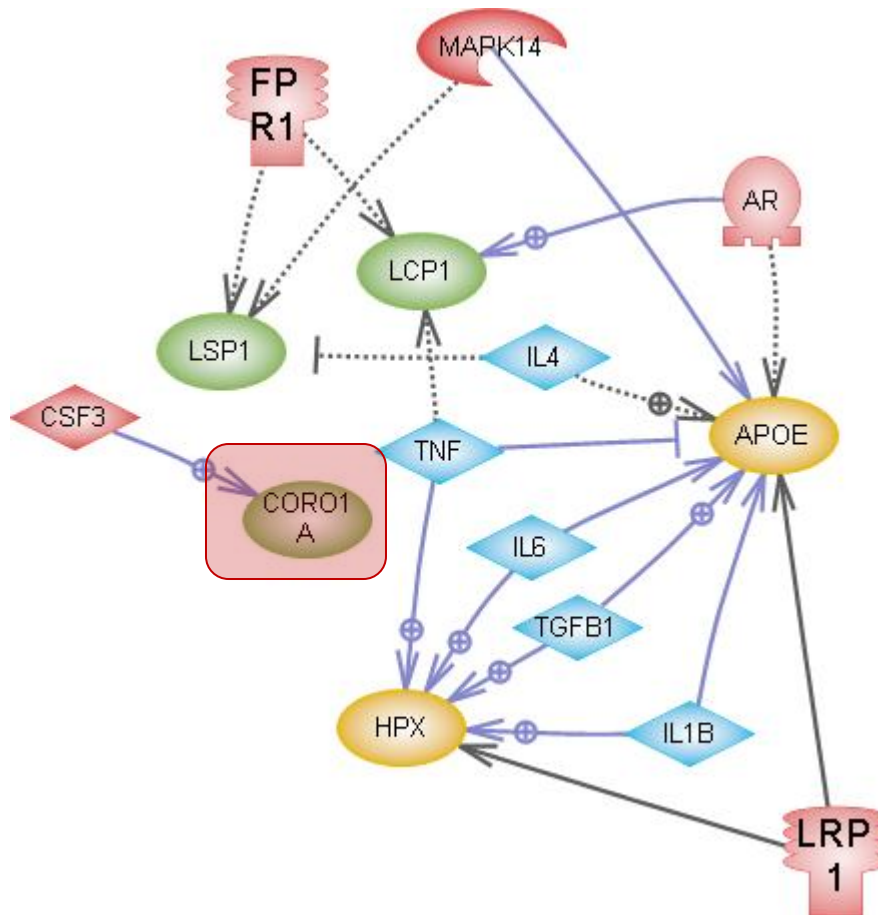
➤ Down regulated in complete model (*Haenen et al., 2010*)

Proteome changes after dermal sensitization

LSP-1



Proteome changes after dermal sensitization



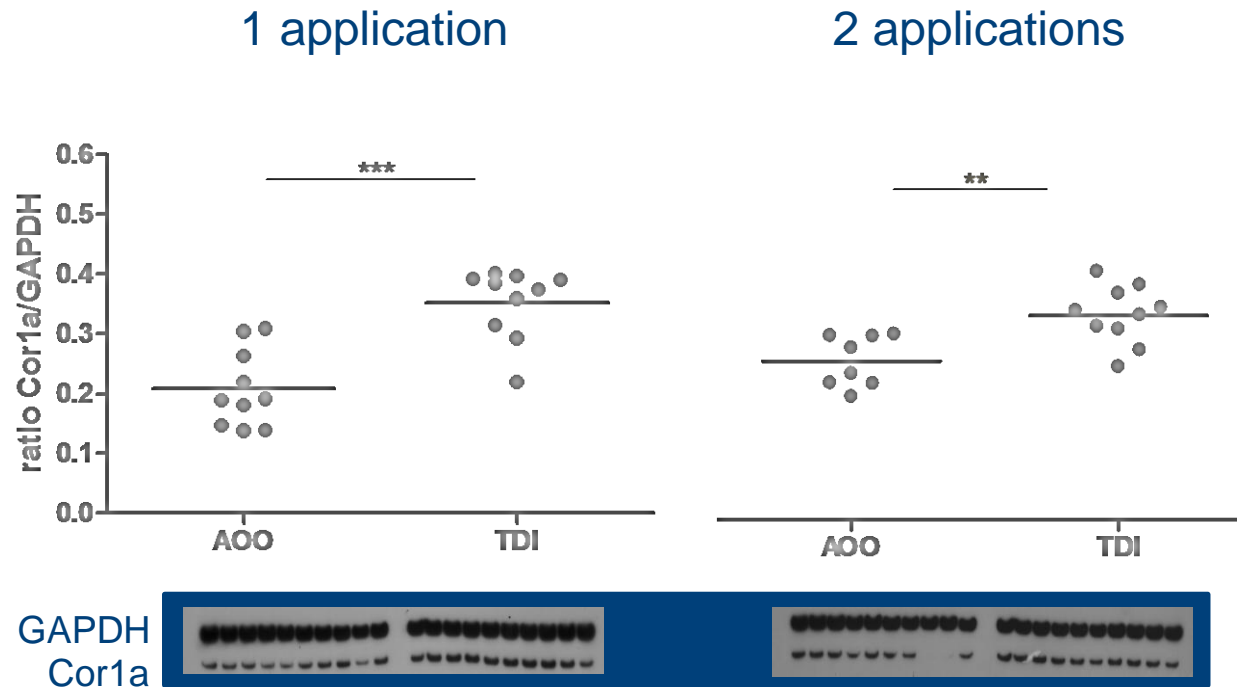
➤ Coronin 1a

➤ Literature:

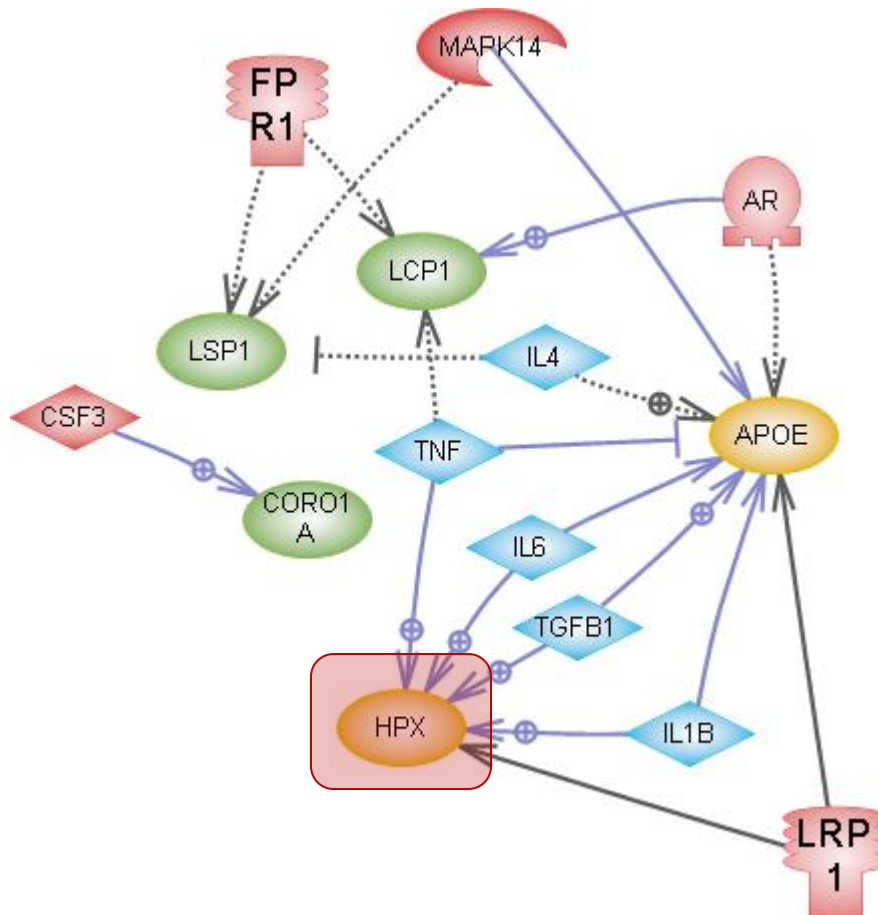
- Member of well-conserved family of coronin proteins
- Mediates Ca^{2+} release
- Involvement T cell activation and proliferation

Proteome changes after dermal sensitization

Cor 1a



Proteome changes after dermal sensitization



➤ Hemopexin

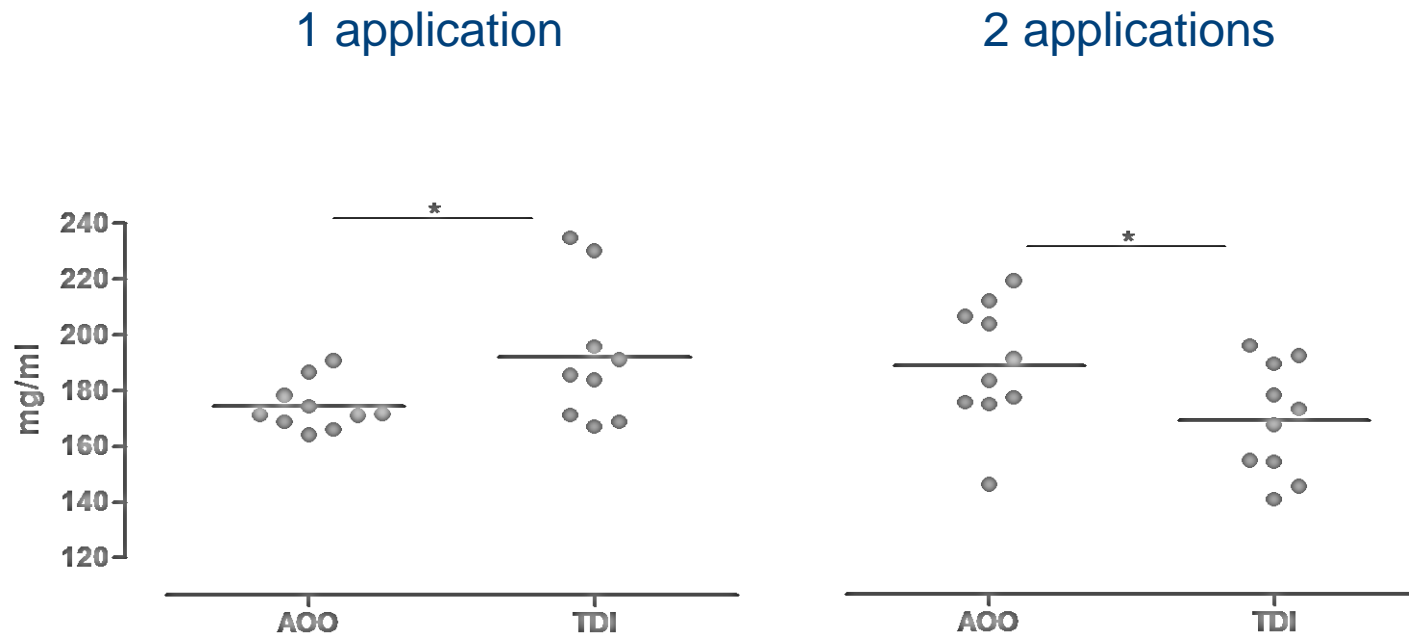
➤ acute phase protein

➤ Literature:

- Involved in innate body defense
- Upregulated during inflammation
- Also in complete model (BAL and serum)

Proteome changes after dermal sensitization

Hemopexin



Conclusions

1. Sensitization causes profound differences in the proteome of TDI-sensitized mice compared to control mice
2. A subset of proteins were confirmed in an independent set of mice
3. Validation needed in human exposed workers

Acknowledgements

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Evelyn Maes

Prometa

Dr. Geert Baggerman