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Integrated Data Analysis of *In Vitro* Testing Approaches Advancing BioTech Product Development and Safety.

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Sens-it-iv Consortium





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A few clarifications

- 'My' BioTech products:
 - Industrial enzymes
 - Pharma proteins and peptides



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- 'My' Expertise:
 - Irritation
 - Carcinogenesis
 - Sensitization



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A few clarifications

- 'My' BioTech products:
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- 'My' Expertise:
 - Irritation
 - Carcinogenesis
 - Sensitization

- Specific issues:
 - Potency is defined by "number of patients" not "concentration"
 - Hazard and risk assessment "relative" to historical data



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Outline of the presentation

- Ensuring safety of technical enzymes without using animals
 - Historical data, data waiving and read-across
 - Solid basis for developing ITS for relative hazard identification

- Future hazard identification and risk management of industrial enzymes:
 - Computational approaches
 - *In vitro* characterization
 - Implementation of the Sens-it-iv toolbox
 - Integration of data

- Publicity:
 - Congress on In Vitro Sensitization



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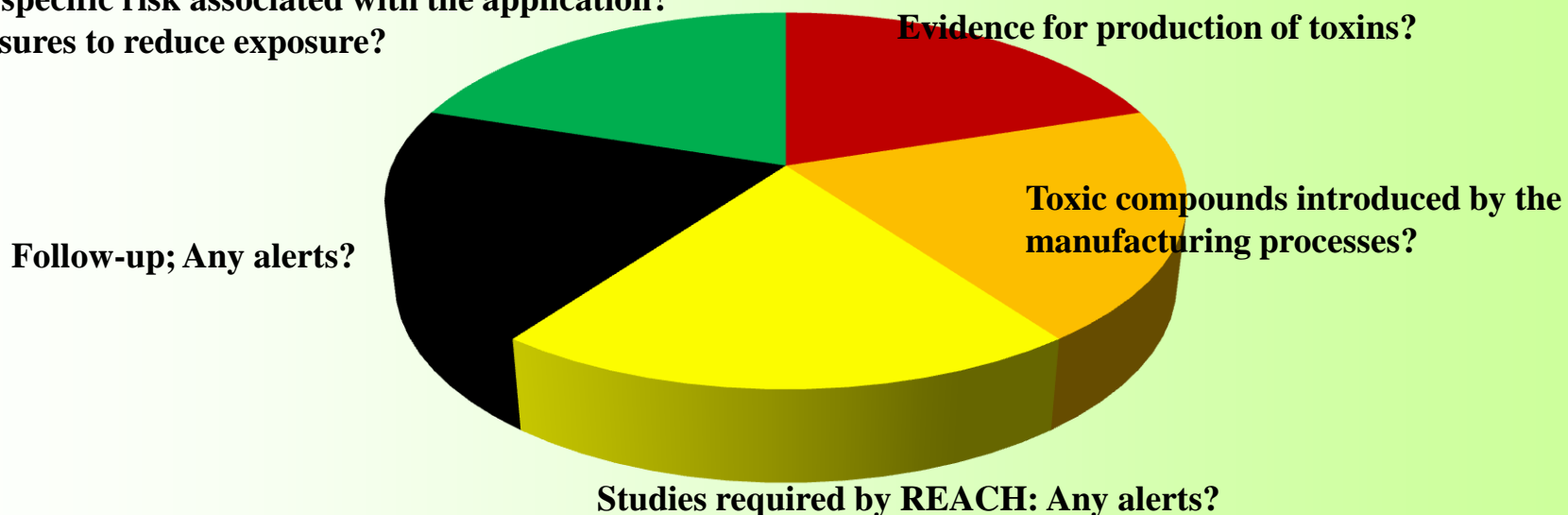
A proposed strategy for ensuring safety of detergent enzymes without using animals in the context of REACH



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Risk assessment of industrial enzymes in the past

- Any specific risk associated with the application?
- Measures to reduce exposure?



- Production strain
- Constituents
- Animal data
- Human data: Clinical and occupational follow-up
- Exposure



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Historical data, data waiving and read-across in the REACH context

- 40 of experience and substantial documentation on
 - The safety of production strains
 - Good standard manufacturing processes
 - Characterization of the enzyme test materials
 - High quality studies for all relevant (REACH) endpoints
 - *In vivo* as well as *in vitro* studies
 - Measurements for eliminating exposure (granulates)
- Solid basis for applying read-across and data waiving without compromising safety (applied for REACH registration)
- Solid basis for development of ITS for relative hazard identification (product development and safety)





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Future strategy for ensuring safety of industrial enzymes without using animals

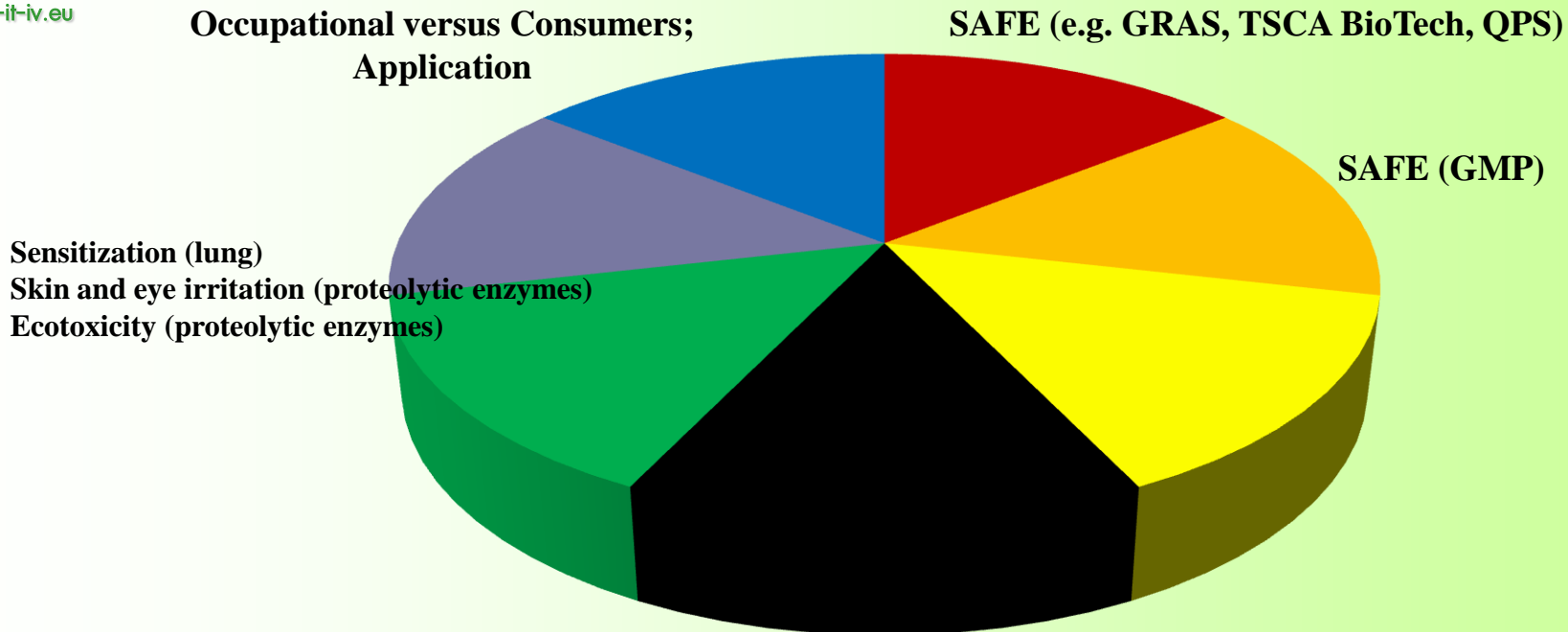


SIXTH FRAMEWORK
PROGRAMME



Future hazard identification strategy

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- Production strain
- Constituents
- Historical animal data
- Human data: Clinical and occupational
- Data from computational methods
- Data from in vitro methods
- Exposure



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1. Computational approaches:

From the protein point of view ...



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The computational approach for assessing protein characteristics

- Size (1-40 kDa) and structure (?)
- Physico-chemical characteristics
 - Hydrophobicity/-philicity; sheets; loops; turns; flexibility; accessibility; ...
- Specific sequences that are recognized by T and B lymphocytes (epitopes)
 - Linear (T, B) or conformational (B)
 - Databases and prediction tools available
- Impact of the protein on the viability/functionality of mammalian (human) cells and cellular processes (under construction)
 - Enzyme activity; substrate binding (incomplete); receptor-ligand like interactions





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Comparison of the predictivity of various prediction models (B cell epitopes)

Protein structure	Prediction model		PPV (%)
Primary and secondary	Single scale	hydrophilicity	48-61
		polarity	54-58
		accessibility	50-58
		flexibility	50-71
		turns	34-72
	Combination of scales		55-65
	Machine learning tools	ABCpred, Bepi-pred	60-68
		SPA	<35-65
Primary and 3D-structure		Structural similarities	Considered too low
3D-structure	Combination of scales (see above)	CEP, DiscoTope, BEpro	66-71
	Using amino acid sequences and motifs	Pepitope	41-54
		Pep-3D-Search	70
		EMT	79-100

Roggen (2008) In *Immunogenicity of Biopharmaceuticals* (van de Weert M and Møller EH) pp 75-95, AAPS Press

Roggen (2008) *Drug Discovery Today: Technologies* 5, 49-55



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2. Cellular processes:

Implementation of the selected tests from the Sens-it-iv toolbox





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Overall objective of the *Sens-it-iv* project

- Development of tools that are ready for prevalidation.
 - Physiologically relevant
 - Cell culture conditions supporting relevant cell phenotype, cell-cell interactions and cell-compound interactions
 - Mechanism of action
 - -omics
 - Addressing specific questions
 - Solid knowledge about *in vivo* processes required
 - SOP(s)
 - Including detailed performance criteria
- Tools that can be incorporated in ITS



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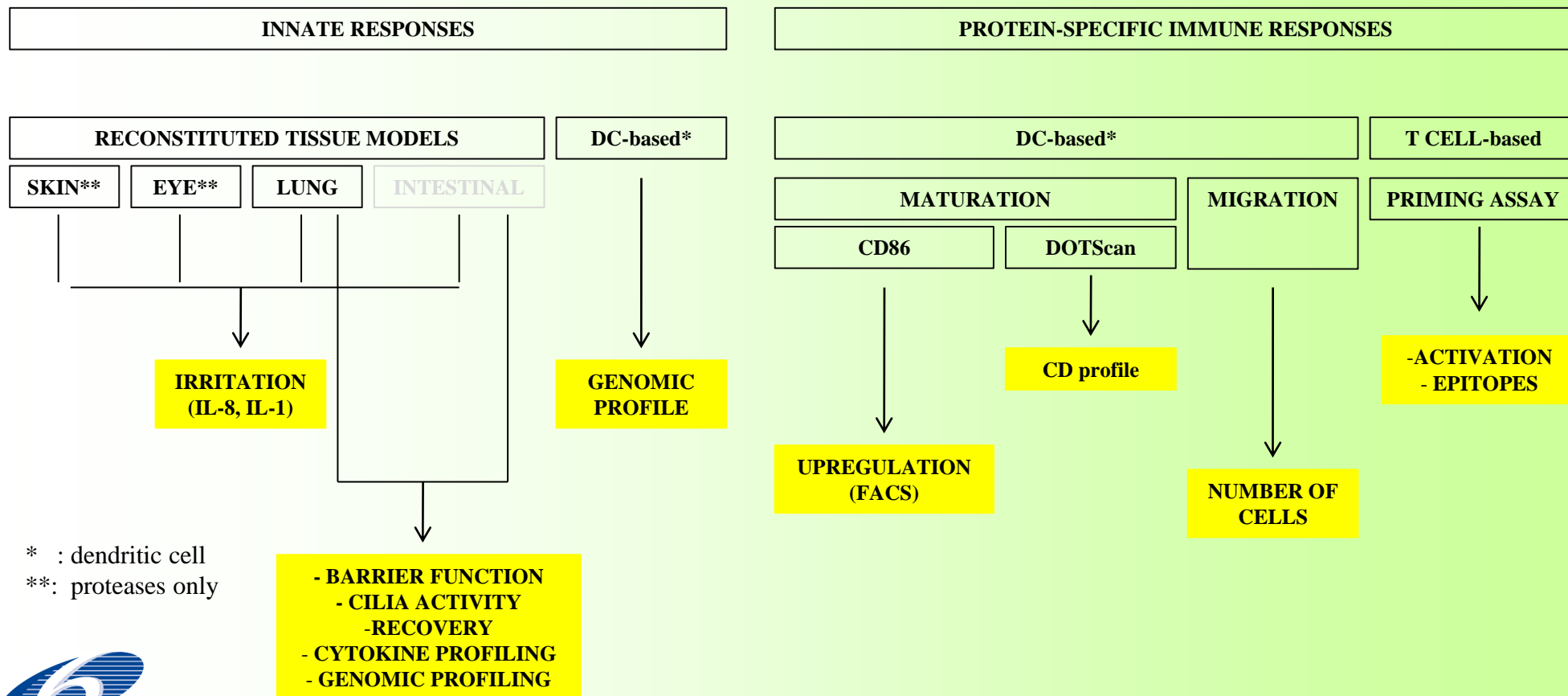
“The Sens-it-iv Toolbox”

	Protein binding
Keratinocytes	NCTC2544 test
	Human reconstituted skin
Lung EC	Precision cut lung slices
	Human reconstituted <u>alveolar</u> epithelium *
	Human reconstituted <u>bronchial</u> epithelium *
	Specific sensitizer genomic profil *
DC	Genomic Allergen Rapid Detection (GARD) test *
	Maturation #1 (CD86, CD54, IL-8, ...) *
	Maturation #2 (DotSCan) *
	Migration *
T-cells	Primary T-cell stimulation *
Bioactivation	Neutrophil - THP-1 metabolization tests
	Proteomics marker profile (combined list)



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Measuring the impact of a protein on cells and cellular processes *in vitro*

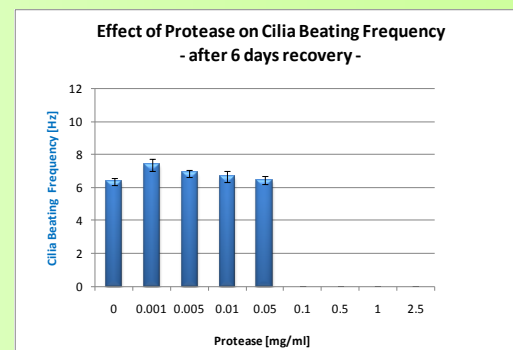
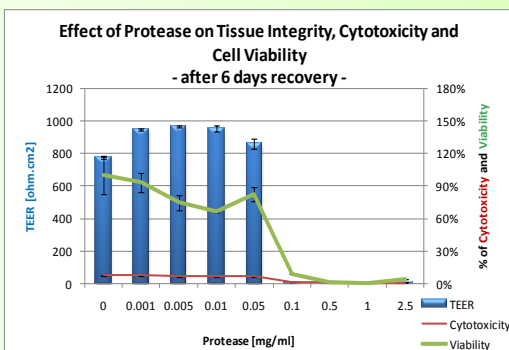
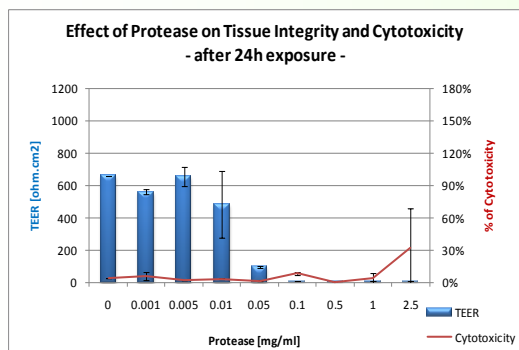
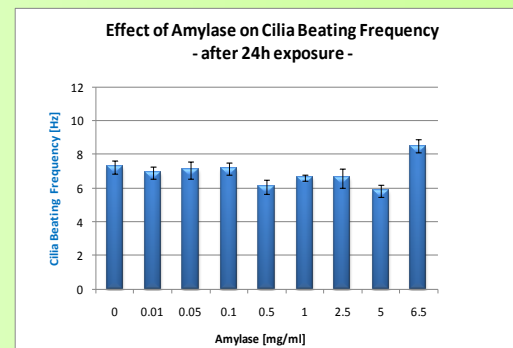
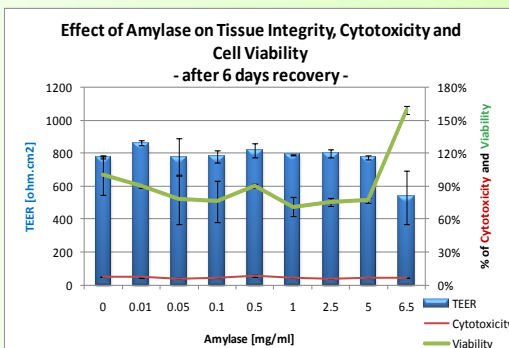
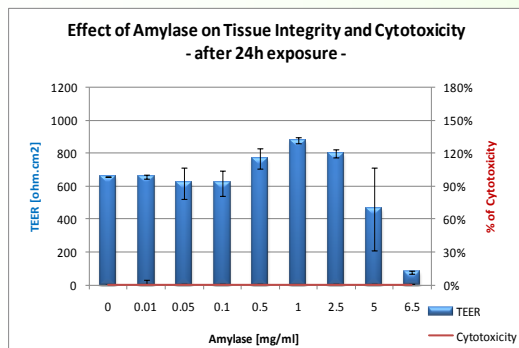


* : dendritic cell
**: proteases only



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Reconstituted human bronchial tissue: Barrier function, cilia beating and recovery

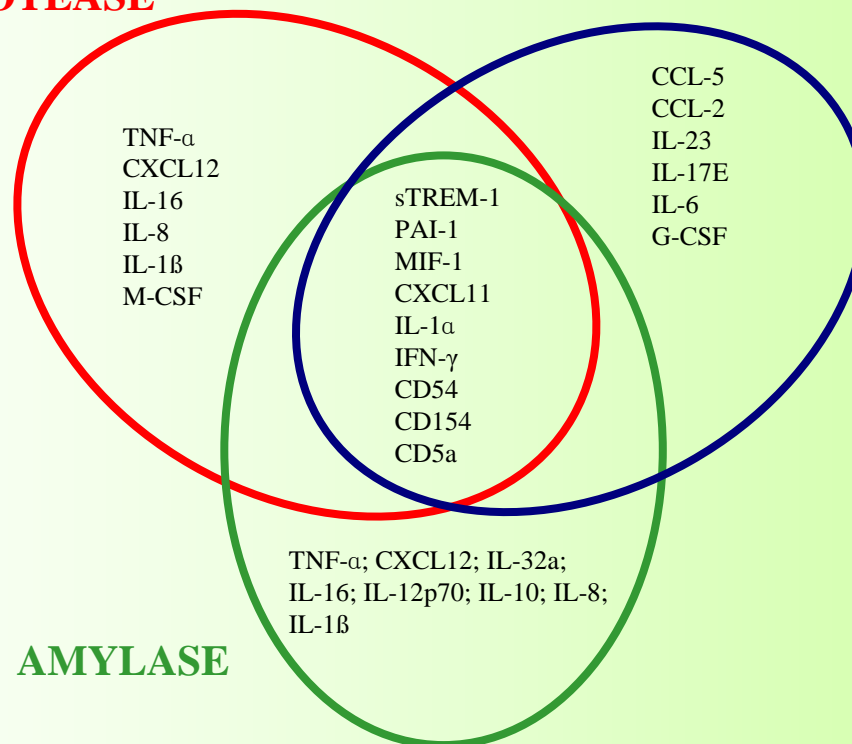




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Reconstituted human alveolar tissue: Cytokine profiling

PROTEASE



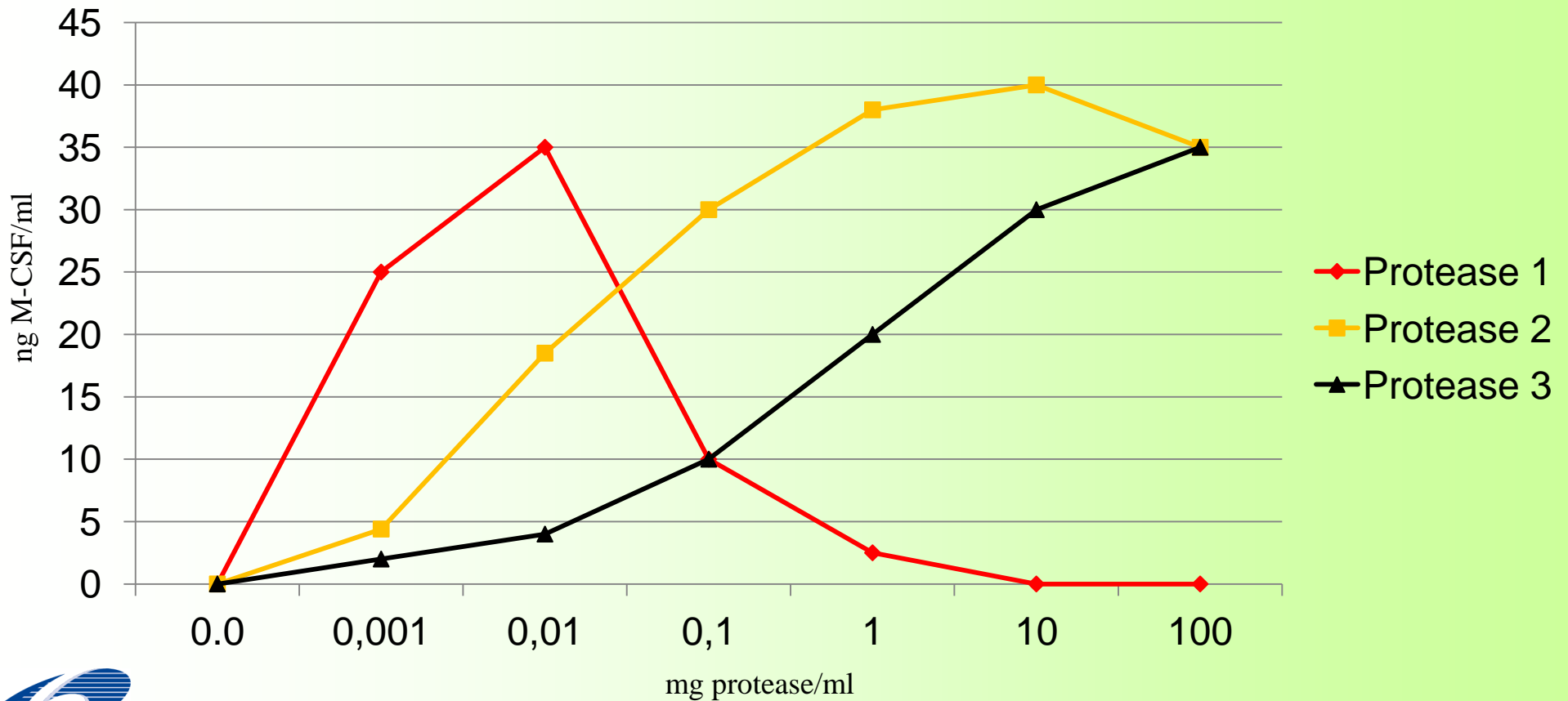
LIPASE

AMYLASE



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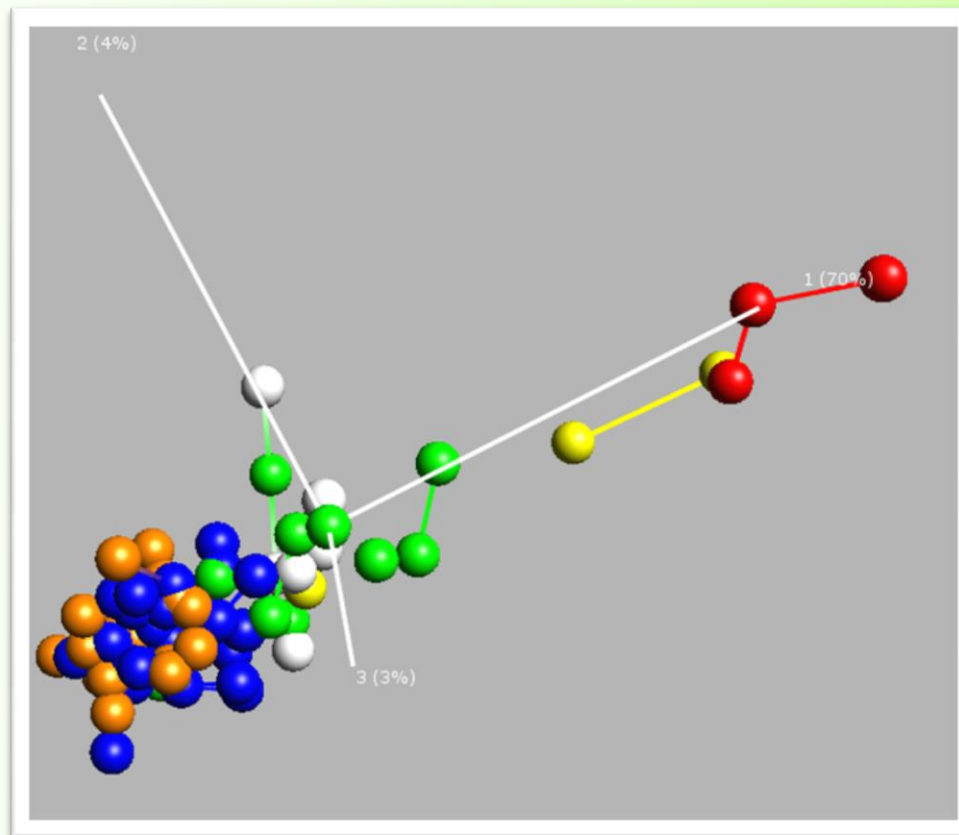
Ranking of enzymes according to potential sensitizing potency: an example





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Genomic Analysis of MUTZ-3 Cells after exposure



C. Borrebaeck et al. Lund University (S)

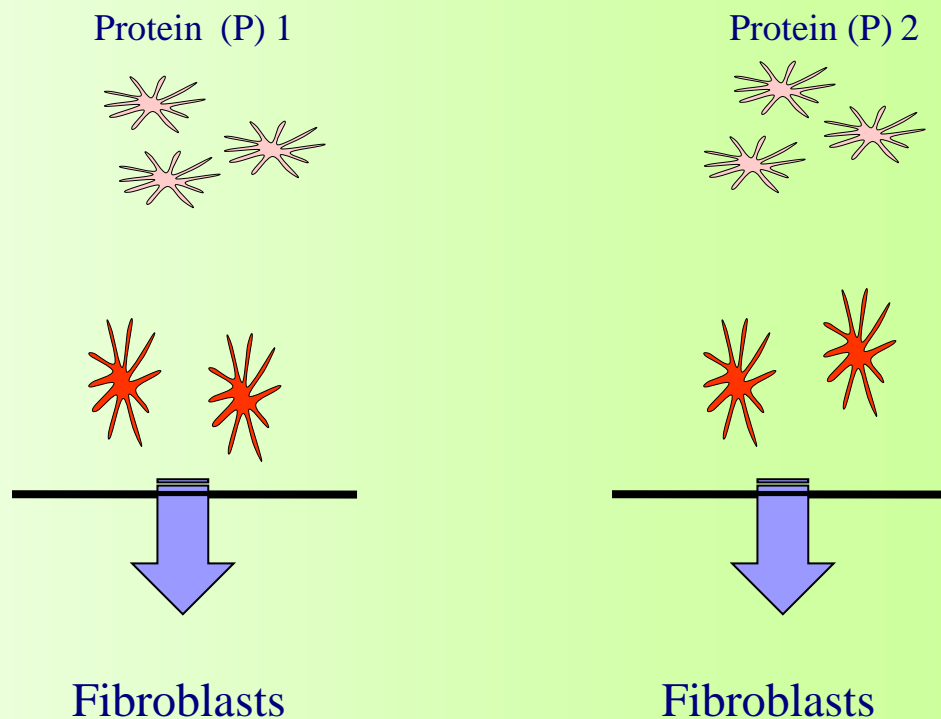


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Transwell DC migration assay: the principle

CSFE-labeled MUTZ-LC

The read-out of the assay is the
number of cells migrating
towards the lower compartment.





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Integration of the data is ongoing

Enzyme activity	Variant / class	In vivo data		Computational		Innate immune responses					Adaptive immune responses		
		Animal	Human	B cell	T cell	Barrier function (mg/ml)	Cilia beating (mg/ml)	Recovery	Cytokine profile (mg/ml)	Genomic profile	DC maturation	DC migration	T cell priming
Protease	1	Animals		Epitope lists available		0.01	0.01	no	0.001	Analysis in progress	Analysis in progress		Epitopes identified
	2	Guinea pig, Rat, Mouse		Overlaps and differences identified		0.1	0.1	no	0.1				
	3	Serological data				10	10	yes	10				
Amylase	A	Immunochemical characterization		Epitope lists available		10	10	yes		Analysis in progress	Analysis in progress		
	B	Humans		Overlaps and differences identified		No effect	No effect	--					
Others (Lipase)		Clinical studies, Occupational data		Epitope lists available		No effect	No effect	--					
		Serological data											



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		Serological data											

Seems to be confirmed by animal studies

Also observed in animals and humans



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Summary

- Set-up still 'unstable'
 - Sufficient/adequate testing?
 - Avoid 'overkill' for the sake of time and money
 - Integration of historical data, computational data and in vitro data to be optimised
 - The weight of each input still to be established
- Ambition to be able to perform 'relative' hazard identification and risk assessment
- Learnings from industrial enzymes should help to establish testing strategies for BioPharmaceuticals

