

Ad hoc meeting with Industry _23 June 2021

Allergenicity assessment

GMO Unit



Allergenicity assessment



GUIDELINES FOR PREDICTION

Codex Alimentarius 2003-2009

EFSA GMO Panel - 2010

- 2011

<u>- 2017</u>

CEP Panel - 2009

FEEDAP Panel - 2008

- 2017

NDA Panel - 2016



Foods derived from modern biotechnology

2003-2009

Second edition



WoE approach

Allergenicity assessment – Weight of evidence



The information in the WoE includes:

- Source of the protein (IgE and non-IgE)
- Amino acid sequence comparison
- In vitro degradation studies (mainly pepsin test)
- Specific serum screening
- Cell based / in vivo assays

On a case-by-case basis



Foods derived from modern biotechnology

Second edition

Protein safety – Allergenicity



Complex questions raised decades ago:

What makes a protein an allergen?
How much is too much?



- Whole foods as well as individual proteins to be risk assessed
- The risk assessment (RA) should take into account:
 - not only cross-reactivity of a reaction
 - but also *de novo* sensitisation
- Different sources of information taken into account in the RA
- No validated method available Weight-of-evidence (WoE)

Future innovative foods





Novel food sources: Food 2030 - Innovative EU research ensures food system is future-ready

https://ec.europa.eu/knowledge4policy/publication/food-2030-innovative-eurosearch-ensures-food-system-future-ready_en

Nutrition for more sustainable and healthy diets; Climate resilience and environmental sustainability; Circularity and resource efficiency; and finally, Innovation and the empowerment of communities











Protein safety – Allergenicity



Complex questions still remains:

What makes a protein an allergen?
How much is too much?



 An urgent need to improve and better modernise the allergenicity assessment prediction



 EFSA activities with the focus on allergenicity assessment: Stakeholders involvement











EU projects

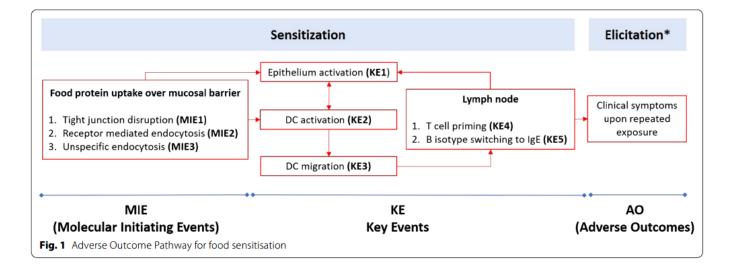


Verhoeckx et al. Clin Transl Allergy (2020) 10:13 https://doi.org/10.1186/s13601-020-00318-x Clinical and Translational Allergy

REVIEW Open Access

COST Action 'ImpARAS': what have we learnt to improve food allergy risk assessment. A summary of a 4 year networking consortium

Kitty Verhoeckx^{1*}, Katrine Lindholm Bøgh², Anne Constable³, Michelle M. Epstein⁴, Karin Hoffmann Sommergruber⁵, Thomas Holzhauser⁶, Geert Houben¹, Annette Kuehn⁷, Erwin Roggen⁸, Liam O'Mahony⁹, Ben Remington¹ and René Crevel¹⁰



- Introduction of new foods should not add to the burden of food allergy
- Need of a reliable, harmonized, evidence-based and validated allergenicity RA strategy
- To make use of novel methods and knowledge
- To focus on future research

EFSA activities





SCIENTIFIC OPINION

ADOPTED: 18 May 2017 doi: 10.2903/j.efsa.2017.4862

Guidance on allergenicity assessment of genetically modified plants

EFSA Panel on Genetically Modified Organisms (GMO),
Hanspeter Naegeli, Andrew Nicholas Birch, Josep Casacuberta, Adinda De Schrijver,
Mikolaj Antoni Gralak, Philippe Guerche, Huw Jones, Barbara Manachini, Antoine Messéan,
Elsa Ebbesen Nielsen, Fabien Nogué, Christophe Robaglia, Nils Rostoks, Jeremy Sweet,
Christoph Tebbe, Francesco Visioli, Jean-Michel Wal, Philippe Eigenmann, Michelle Epstein,
Karin Hoffmann-Sommergruber, Frits Koning, Martinus Lovik, Clare Mills,
Francisco Javier Moreno, Henk van Loveren, Regina Selb and Antonio Fernandez Dumont

Abstract

- Non-IgE-mediated adverse immune reactions
- In vitro protein digestibility
- Endogenous allergenicity

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Keywords: guidance, allergenicity assessment, newly expressed proteins, endogenous allergenicity, GMO

EFSA activities focused on:

- A statement on the usefulness of the pepsin resistance test
- A workshop organised 15-16
 June 2021
- A scientific opinion on future development needs

Looking closer at the future needs



More robust bioinformatic approaches:

- Current methodologies date back to 2001
- Databases development
- EFSA procurement on peptide modeling ongoing
- More refined in vitro protein digestion protocols
 - Classical pepsin resistance test in place but questioned
 - EFSA procurement on in vitro digestion finalised but more work needed
- More predictive cell-based approaches/ HLA-phenotyping

EFSA procurement on in vitro digestion



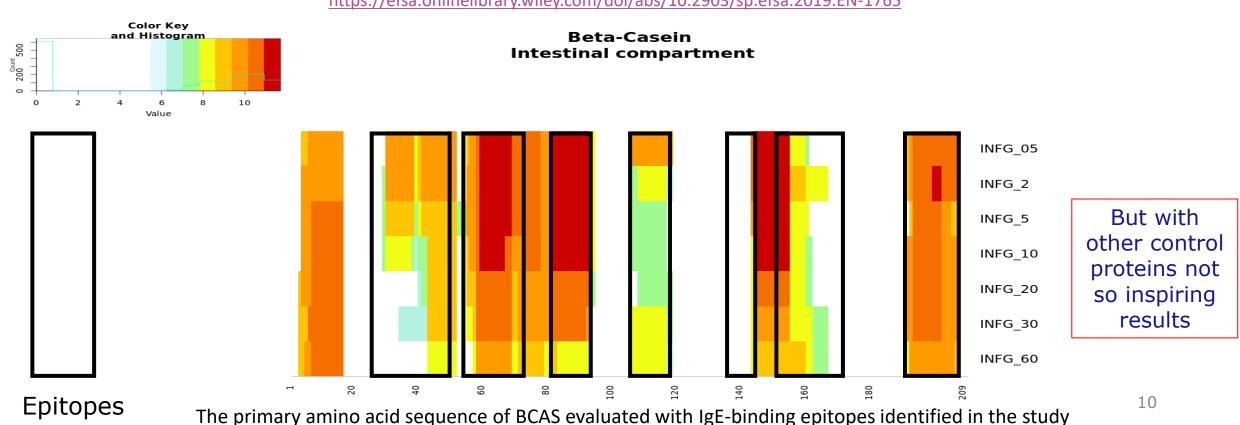
Report on EFSA project OC/EFSA/GMO/2017/01

"In vitro protein digestibility" (Allergestion)

Alan Mackie¹, Didier Dupont², Amelia Torcello-Gómez¹, Julien Jardin², Amélie Deglaire²

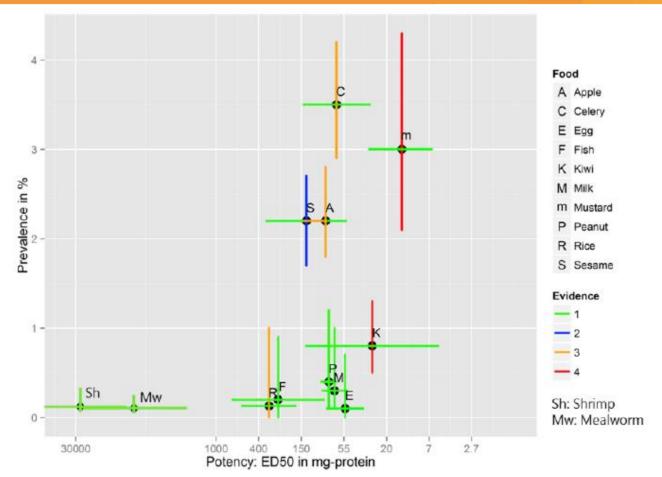
- 1. School of Food Science and Nutrition, University of Leeds, Leeds, LS2 9JT, UK
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https://efsa.onlinelibrary.wiley.com/doi/abs/10.2903/sp.efsa.2019.EN-1765



Example: Scaling allergy risks of foods relatively





Strategy for ranking the allergic potential of known proteins as a way forward

(FAO/WHO, 2001; EFSA GMO Panel, 2017; Remington et al 2018; Verhoeckx et al 2020; Fernandez et al 2020)

Houben et al 2019. Food and chemical toxicology, 127, pp 61-69

Whole foods

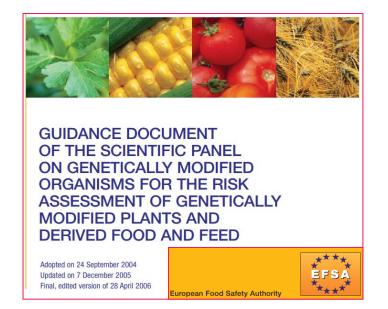


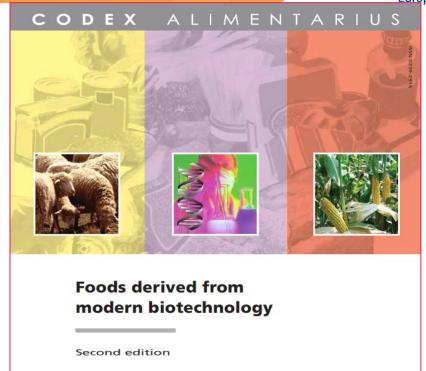
Endogenous allergenicity

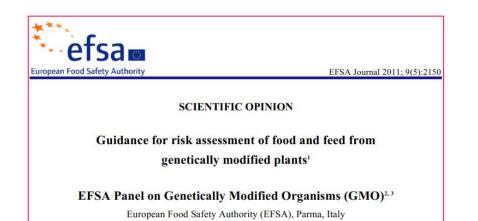


Comparative approach





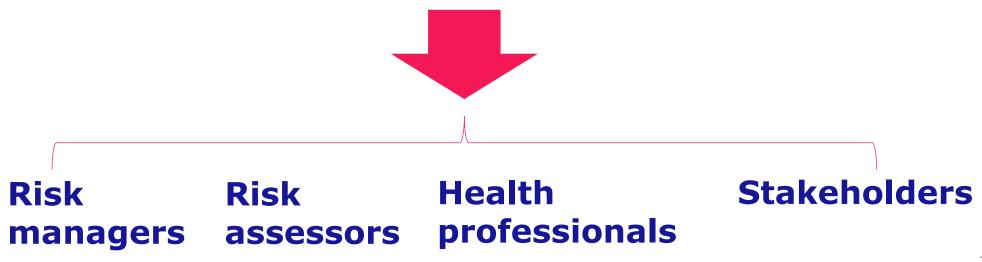




Relevant plants for analysis



- Analysis performed on a case-by-case basis
- For «common allergenic foods» (EU Reg. 2003, 2011)
- To date: soybean is the main crop analysed
- Other GM plants than soybean: whenever considered necessary



Relevant allergens for quantification



EU Regulation 503/2013



OECD consensus soy 2012 Soy allergen list:

'potential soybean allergens'

Table 30. Potential soybean allergens

IgE-binding provides	Allerges nomenclotuse	Molecular weight (kDs)	Semily
Bytroptobic governs	Opn fi	7925	Lipid transfer provin
Delenia	Oy = 7	8.9	Swage protein
hollin	Oym 5	34	Profile
90EE	Olym 4	166	Petropaccia related pretein PS-10
D4	00y to 880.59 K	34	Project
Takanwa Asa-liaked ghosponeia	60 a 50 30 K	36	Usborn
(Conglycials (vicilia, 75 photos)	Gym 5	140-178	Stocop protein (with submits)
Gycinia (legonia, 115 globulia)	Clym f	330-169	Storage product (with robusts)
25 eBonin	Notanipad	- 11	Probasis
Lettin	Nationiped	130	Lectio
Ljonygase	Notoniped	161	Encyme
Energy regular statistics	Not integer?	- 31	Protocor solution
Eslaces	Notseaped	36	Unknown
Eslaces	Youniped	90	Homology to delongity if A-B teaching protein
R0-25	Notossigned	23-25	Unicore

Source stopped from L. Rocine and Boys, (2007); updated with information from WHO LTS (2011)

Evidence check



Evaluation of literature for all single allergens

and

Comparison and complementation with databases (EFSA, 2010)

and/or

Systematic Reviews



Clinical relevance shown



Relevance for GMO risk assessment

EFSA Guidance 2017

¹ WHO IUS (341) Albegos associates excipated by VBO and IUS

Methodology



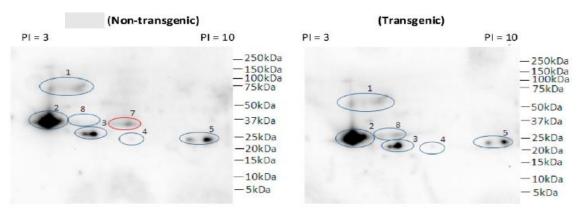
- Quantitative ELISA
- Quantitative mass spectrometry

Future development of an allergen database (<u>natural variability</u>)

comparative approach



Historically: human sera (IgE-binding)



Goodman et al. 2013, J. Agric. Food Chem. 2013, 61, 8317-8332

Data interpretation



Natural variability of allergens





- On case-by-case basis
 - Magnitude and number of changes
 - Clinical relevance of the allergen(s) involved
 - Exposure considerations
 - Clinical evaluation

Few overall remarks



Scientific developments

Translated into

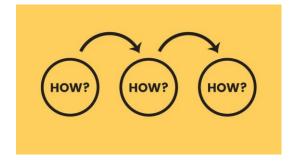


Risk assessment requirements

A reliable, harmonised, and evidence-based allergenicity RA strategy (e.g., Codex, EFSA, Cost actions, others)



Consensus permeable to evolution



EFSA Workshop – 15/16 June 2021



Session on clinical relevance and risk assessment

How to better use available human data on allergic reactions to known allergens?

Session on in silico approaches

What is the usefulness of current FASTA algorithms and of alternative approaches? How can the current allergen database be improved?

Session on in vitro/in vivo approaches

What are the most relevant test materials to predict allergenicity?
When and how should human sera or alternatives be used to address challenges?

Session on outstanding questions

What are the most effective methods to predict allergenicity? How should they be integrated? How best to validate *in silico* and *in vitro* methods (e.g., animal models)? How can PMM tools be developed and how to determine acceptable levels and/or thresholds of protein?



Thank you very much

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