



Predictive Analysis of Allergens in Novel Foods by Using Advanced Proteomic and Bioinformatic Tools

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BACKGROUND & OBJECTIVES

One of the most significant challenges in bringing novel foods to market is ensuring food safety. New food scenarios drive the need to detect novel allergens that must be identified and quantified for proper labeling. Allergenic reactions are primarily caused by proteins that are abundant in foods, typically of low molecular mass, glycosylated, water-soluble, and highly stable to proteolysis. The most relevant plant and animal food allergens, such as lipid transfer proteins, profilins, seed storage proteins, lactoglobulins, caseins, tropomyosins, and parvalbumins from fruits, vegetables, nuts, milk, eggs, shellfish, and fish, have been studied. As a massive preliminary screening, proteomic methods should be employed to search for potential allergens. This work focuses on proteomic and bioinformatic tools for food researchers to identify allergens in novel foods.

ALLERGENS IN TRADITIONAL FOOD

Table 1. Several protein allergens in food. For the WHO/IUIS nomenclature, the allergens are named according to the species source of food.			
Food	Protein Name	Species	Allergen
Milk	Caseins	<i>Bos taurus</i>	
	α S1-casein (23.6 kDa)		Bos d 9
	α S2-casein (25.2 kDa)		Bos d 10
	β -casein (24 kDa)		Bos d 11
	κ-casein (19 kDa)		Bos d 12
	β-lactoglobulin (18.3 kDa)		Bos d 5
	α-lactalbumin (14.2 kDa)		Bos d 4
Eggs	Serum albumin (66.3 kDa)	<i>Gallus domesticus</i>	Bos d 6
	YGP42 (35 kDa)		Bos d 7
	Ovomucoid (28 kDa)		Gal d 1
	Ovalbumin (44 kDa)		Gal d 2
	Ovotransferrin (78 kDa)		Gal d 3
Fish	Lysozyme (14 kDa)	<i>Gadus callarias</i> (Baltic cod)	Gal d 3
	α-livetin (69 kDa)		Gal d 5
	YP42 (35 kDa)		Gal d 6
	Parvalbumin		Gad p 2
Shellfish	α-parvalbumin (13 kDa)	<i>Gadus callarias</i> (Baltic cod)	Gad p 1
	β-parvalbumin (11.6 kDa)		Gad p 1
Peanuts/tree nuts	Tropomyosin (34 kDa)	<i>Metapenaeus ensis</i> (Shrimp)	Met e 1
	7 S seed storage globulin, vicilins (64 kDa)		Ara h 1
	2 S albumin (17 kDa)		Ara h 2, Ara h 6, Ara h 7
	Nonspecific lipid transfer proteins		Ara h 9, Ara h 16, Ara h 17
Soy	Oleosins	<i>Arachis hypogaea</i>	Ara h 10, Ara h 11, Ara h 14, Ara h 15
	Defensins		Ara h 12, Ara h 13
	Profilins		Ara h 5
	Plant pathogenesis-related proteins PR-10		Ara h 8
Wheat	7 S seed storage globulin, β-conglycinin	<i>Glycine max</i>	Gly m 5
	11 S seed storage globulin, glycinin		Gly m 6
Sesame	α-amylase inhibitor (13 kDa)	<i>Triticum aestivum</i>	Tri a 28
	Gamma gliadin (88 kDa)		Tri a 20
Soy	Elongation factor 1	<i>Glycine max</i>	Tri a 45
	2 S albumins		Ses i 1, Ses i 2
Sesame	7 S vicilin-type globulin (45 kDa)	<i>Sesamum indicum</i>	Ses i 3
	Oleosins		Ses i 4, Ses i 5
Soy	11 S globulin, legumins	<i>Glycine max</i>	Ses i 6, Ses i 7
	Profilin		Ses i 8

ALLERGENS IN NOVEL FOODS

Figure 1. Novel protein sources of plant, algal, fungal and insect origins are being researched by the food industry.

Food	Protein Name	Species
Microalgae	C-phycoerythrin	<i>Microalgae spirulina (A. platensis)</i>
	Thioredoxins	
	Superoxide dismutase	
	Glyceraldehyde-3-phosphate dehydrogenase	
Microalgae	Triosephosphate isomerase	<i>Microalgae chlorella (C. vulgaris)</i>
	viz. calmodulin	
Insects	Fructose-bisphosphate aldolase	<i>Microalgae chlorella (C. vulgaris)</i>
	Fructose-bisphosphate aldolase	
Insects	Tropomyosin, myosin, actin, troponin C (muscle proteins)	<i>Microalgae chlorella (C. vulgaris)</i>
	Tubulin (cellular proteins)	
Insects	Hemocyanin, defensin (circulating proteins)	<i>Microalgae chlorella (C. vulgaris)</i>
	Arginine kinase, glyceraldehyde 3-phosphate dehydrogenase (GAPDH), triosephosphate isomerase, α-amylase, trypsin, phospholipase A, hyaluronidase (enzymes)	

PROTEOMIC APPROACHES TO IDENTIFY ALLERGENS IN NOVEL FOODS

Novel Food	Bioinformatic Tool	Goal/Main Achievements
Bread wheat spelt and rye	Database of Allergen Families-AllFam AllergenOnline Allergome	Comparison of allergenicity in cereal products
Cashews	BLASTP Search against AllergenOnline sequence	Analysis of allergen stability under heat treatment
Goji berries	AlgPred software hybrid approach	Identification of 11 IgE-binding proteins
Macadamia nut	AllergenOnline Immune Epitope Database Analysis Resource (IEDB)	Analysis of homology and linear epitope similarities to known allergens
Medicago sativa	COMPARE allergen database	Identification of three allergenic protein families
Lentil (Lens culinaris)	Blast2GO—Functional Annotation and Genomics	Quantification of major allergen proteins
White- and red-fleshed pitaya seeds	AllermatchTM webtool AlgPred 2.0 AllerCatPro web server	Identification of five potential allergens
Spirulina and chlorella microalgae	AllergenOnline	Six proteins exhibit significant homology with food allergens
Cricket	Allermatch TM webtool AlgPred 2.0 ABCPred Bepipred	Description of the impact of processing on allergenic reactivity of insect proteins.
Cricket Acheta domestica	Database of Allergen Families-AllFam Allergen nomenclatura (WHO/IUIS) CLC Genomics Workbench 20.0.4. AllerCatPro web server	Identification of 20 putative allergens
Lesser mealworms, black soldier flies and their protein hydrolysate	AllermatchTM webtool	Identification of potential allergens by similarity to known allergens
Anisakis simplex, Pseudoterranova decipiens, and Contracaecum osculatum	Blast2GO—Functional Annotation and Genomics AllergenOnline AllerTOP web server ver. 2.0 PREAL web server	Prediction of 53 probable allergens in three species

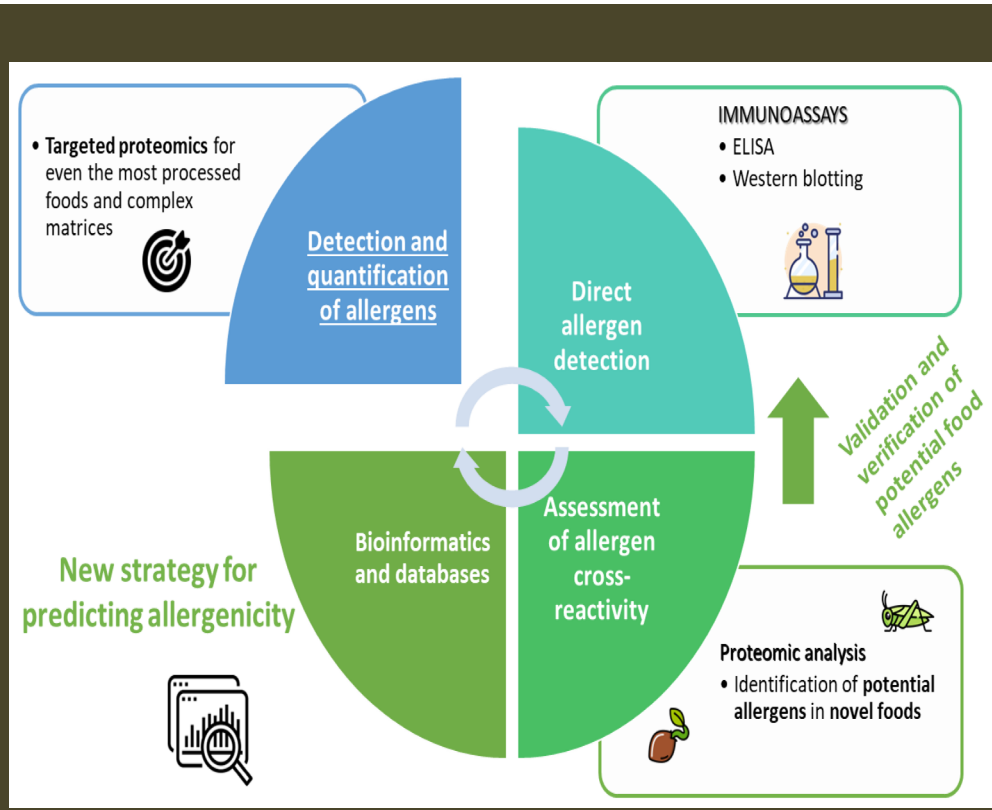
BIOINFORMATIC TOOLS

Table 2. Bioinformatic software tools most used for allergen analysis.

Name	Link (Website)	Description	Name	Link (Website)	Description
Allergen nomenclature	http://www.allergen.org (accessed on 12 February 2023)	Official site for the systematic allergen nomenclature provided by the World Health Organization and International Union of Immunological Societies (WHO/IUIS)	Allergome	http://www.allergome.org (accessed on 12 February 2023)	A website with detailed information on Allergenic Molecules (Allergens) causing an IgE-mediated (allergic, atopic) disease (anaphylaxis, asthma, atopic dermatitis, conjunctivitis, rhinitis, urticaria).
AllerBase	http://bioinfo.unipune.ac.in/AllerBase/Home.html (accessed on 12 February 2023)	Database of allergens detected as IgE-binding epitopes, IgE antibodies and cross reactivity. Allergen data such as experimental information on its allergenic activity and food source is compiled, resulting in a curated database.	Comprehensive protein allergen resource (COMPARE allergen database)	https://comparedatabase.org/ (accessed on 12 February 2023)	A database comprised of protein sequences of known allergens
AllerCatPro	https://allercatpro.bii.a-star.edu.sg/ (accessed on 12 February 2023)	Provides protein allergenicity potential prediction based on the similarity of amino acid sequence and 3D protein structure	Database of Allergen Families-AllFam	http://www.meduniwien.ac.at/allfam/ (accessed on 12 February 2023)	Comprises a resource for classifying allergens into protein families as well as biochemical properties and allergology significance
AllergenOnline	http://www.allergenonline.org (accessed on 12 February 2023)	Provides sequence database of allergens to identify proteins and assess the potential risk of allergenic cross-reactivity. This database offers 2233 peer-reviewed sequences from 912 taxonomic protein groups (February 2021)	Immune Epitope Database and analysis resource (IEDB)	https://www.iedb.org (accessed on 12 February 2023)	Provides experimental data on antibody and T-cell epitopes to identify allergens and to assist in the prediction and analysis of allergenicity
			Structural Database of Allergenic Proteins (SDAP)	https://femi.utmb.edu (accessed on 12 February 2023)	Tool for testing the FAO/WHO allergenicity rules in new proteins and investigating cross reactivity, also offering information about protein sequence and structure

TAKE-HOME MESSAGE

- Proteomic approaches using advanced MS will continue providing relevant information in food safety.
- Detection, identification and quantification of known allergens in complex matrices and highly processed food have already been developed, and targeted MS allows monitoring of them during food processing.
- Identification of novel protein allergens in insects, seaweeds, microalgae or other non-common vegetable foods is one of the most important challenges over the next few years.
- Bioinformatic tools and curated allergen databases will enable the prediction of potential allergens, which should be validated subsequently.
- This information could be used to improve the design and safety of food products by novel devices.
- Advanced technologies, including biosensors, could identify specific interactions between receptors and allergens, enabling us to address the challenges of food safety monitoring.



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