

source: https://cen.acs.org/physical-chemistry/protein-folding/Software-predicts-complex-protein-structures/99/web/2021/11

The importance of plant proteomics

Plants:

- make up more than 80% of the planet's biomass
- are foundational for global ecological and economic systems

Article | Published: 04 February 2019

The global burden of pathogens and pests on major food crops

<u>Serge Savary</u>, <u>Laetitia Willocquet</u>, <u>Sarah Jane Pethybridge</u>, <u>Paul Esker</u>, <u>Neil McRoberts</u> & <u>Andy Nelson</u> □

Nature Ecology & Evolution 3, 430–439 (2019) Cite this article

Climate change: Key crops face major shifts as world warms

source: BBC news

By Matt McGrath
Environment correspondent

(1) 26 January

Our food system isn't ready for the climate crisis

The world's farms produce only a handful of varieties of bananas, avocados, coffee and other foods - leaving them more vulnerable to the climate breakdown

by Nina Lakhani, Alvin Chang, Rita Liu, and Andrew Witherspoon

source: The Guardian

A Pan-plant Protein Complex Map Reveals Deep Conservation and Novel Assemblies

Claire D. McWhite, Ophelia Papoulas, Kevin Drew, Rachael M. Cox, Viviana June, Oliver XiaoouDong, Taejoon Kwon, Cuihong Wan, Mari L. Salmi, Stanley J. Roux, Karen S. Browning, Z. Jeffrey Chen, Pamela C. Ronald, and Edward D. Marcotte

in Cell 181, 460-474, April 16, 2020



presented by Annika Bamberger & Ann-Kathrin Otto

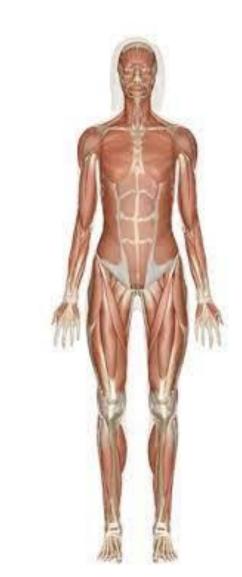
Why did so many plant proteins remain uncharacterized for such a long time?

- plant genomes are complex:
 - polyploidity
 - whole-genome duplications



source: https://agricsite.com/wheat-farming/

- hexaploid
- > 100,000 genes
- 42 chromosomes



source: https://www.innerbody.com/image/musfov.html

- diploid
- ca. 30,000 genes
- 46 chromosomes
- → Plants have mulitple near-identical proteins

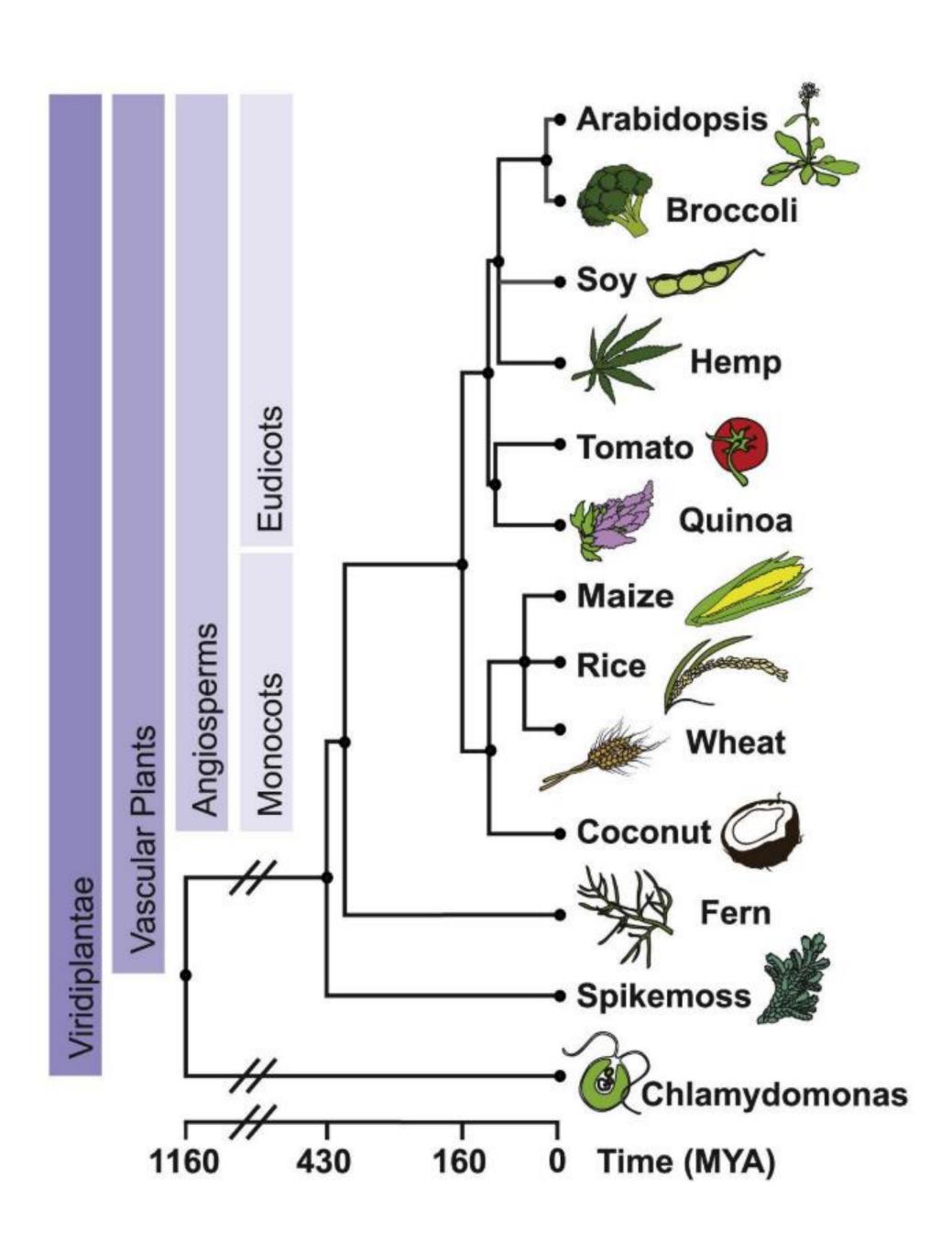
An evolution-informed protein grouping approach

Orthogroups (OGs): "Sets of genes in modern organisms that derive from the same original gene in those organisms' last common ancestor." (McWhite et al., 2020)

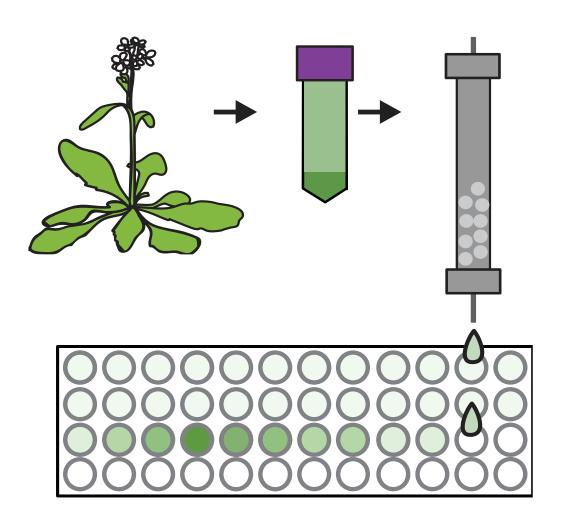


- eggNOG-mapper
- eggNOG database
- highly related proteins clustered into same OG
- overcome problem of polyploidity and redundant proteins

The dataset

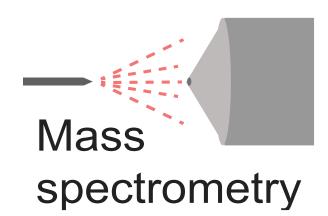


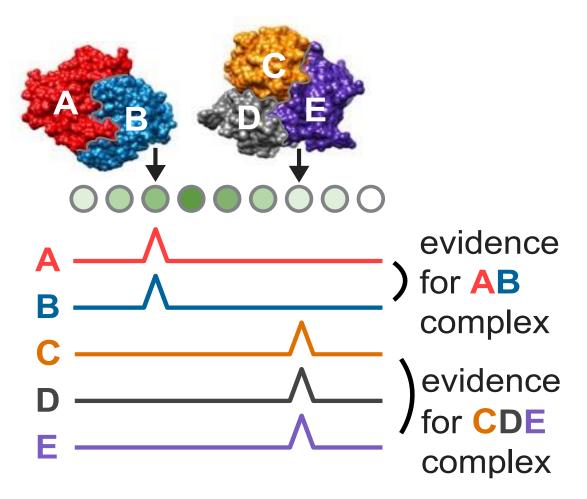
Co-fractionation mass spectrometry (CF-MS)



1. Chromatography/Fractionation

 separate protein complexes into fractions

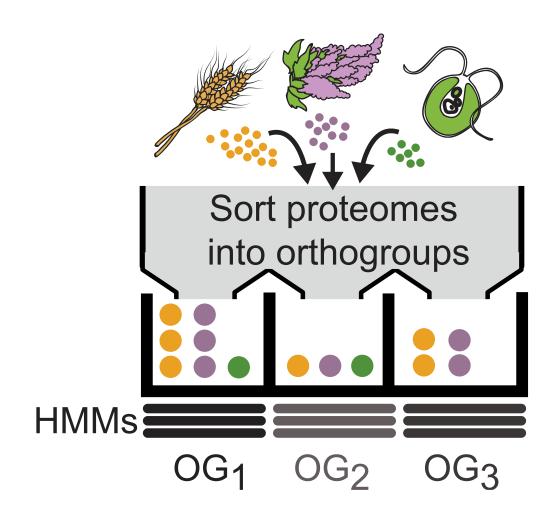


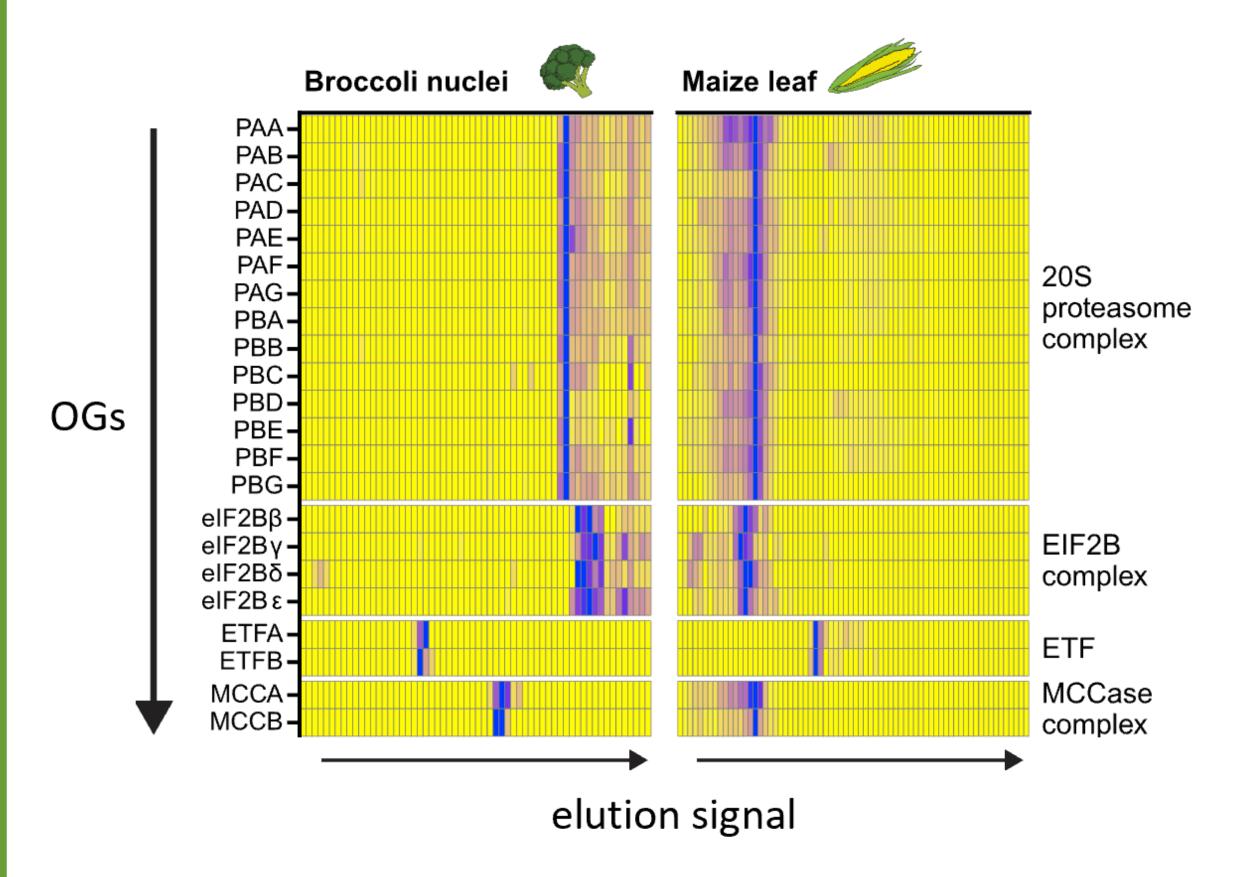


2. Mass Spectrometry (MS)

- identify proteins within fractions
- elution-profile
- co-elution

A cross-species OG detection profile



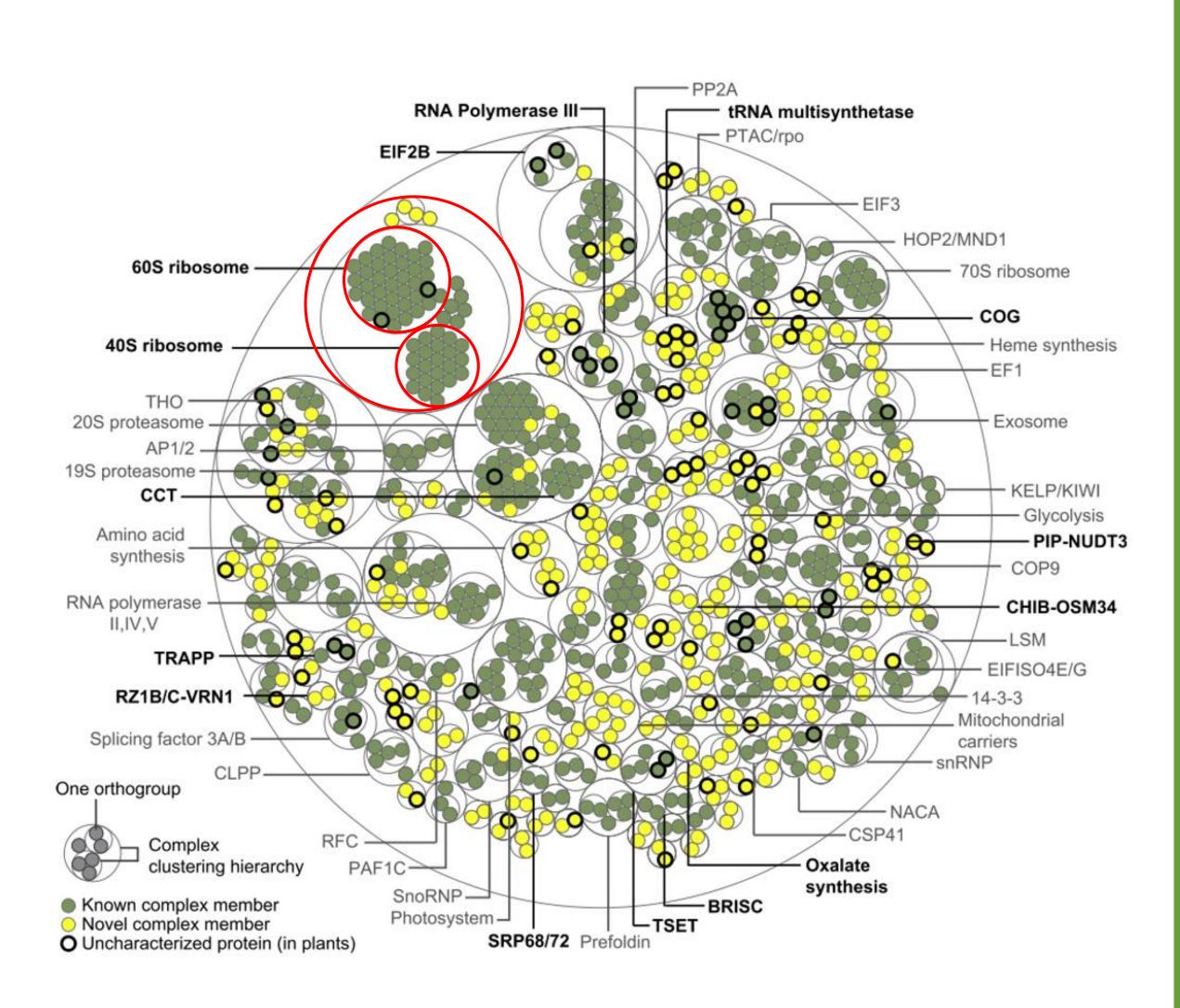


Computational framework

- for the identification of co-eluting proteins
- supervised Machine Learning approach
- → probabilistic CF-MS score: likelihood of a physical association

Identification of many new complex members

unsupervised hierarchical clustering



Differences in protein interactions in plants and humans

- arrangements of interacting proteins
 specific subunits
- analog assemblies
 - similar functions
 - different molecular interactions
- → Sequence analysis not always suffices to predict the function and structure of a protein complex.

Findings that could help solve food crop problems

- linking protein interactions to phenotypes
 - identification of complexes involved in pathogen defense
 - example: CHIB-OSM34, involved in combatting gray mold infections





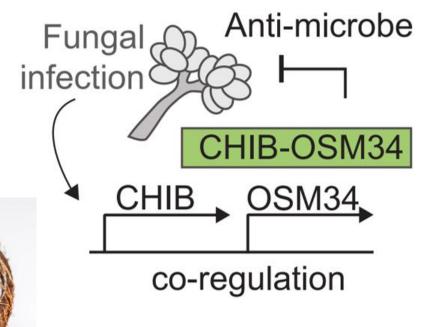




image sources:

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Thank you for your attention.

List of additional literature:

Martinez-Perez et al. (2003). Chromosomes form into seven groups in hexaploid and tetraploid wheat as a prelude to meiosis. *The Plant Journal* **36**, Issue 1, pp. 21-29.

National Human Genome Research Institute, (2020). Human Genome Project FAQ. [online] https://www.genome.gov/human-genome-project/Completion-FAQ

World Economic Forum, (2021). Visualizing the total biomass of every animal on earth. [online] What is the total weight of all the animals on Earth? | World Economic Forum (weforum.org)

Link to the quiz: https://forms.gle/g6ygUuGeyK4icJBL6