

A Scoping Review: Metabolomics Signatures Associated With Animal or Plant Protein Intake and Their Potential Relation to Cardiometabolic Risk

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A scoping review: metabolomics signatures associated with animal or plant protein intake and their potential relation to cardiometabolic risk

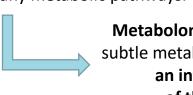
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Context:

• A dietary shift from animal protein (AP) to plant protein (PP) sources have been associated with lower cardiovascular and diabetes risk

Mechanisms are still poorly characterized and might involve many metabolic pathways.



Metabolomics is a relevant tool to explore subtle metabolic reorientations as it provides an integrative characterization of the phenotype expressed

DNA - Genome mRNA - Transcriptome Proteins - Proteome **Metabolites - Metabolome**

Objective:

This scoping review is aimed at gathering and analyzing the available metabolomics data associated with PP and AP-rich diets, and discussing the metabolic effects underlying these metabolomics signatures and their potential implication for cardiometabolic health

Results:

24 studies selected

Study design	Dietary comparison	n [ref]
Observational studies	AP and PP intake	1 [1]
	Omnivores vs vegetarians or vegans	6 [2-7]
	Adherence levels to defined dietary patterns	6 [8-13]
	Adherence to a posteriori (data-driven) dietary patterns	3 [14-16]
Interventional studies	Omnivores vs vegetarians or vegans	3 [17-19]
	Adherence levels to defined dietary patterns	5 [20-24]

Table 1: Classification of the studies selected

References

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- 8: Schmidt et al, 2015, DOI: 10.3945/ajcn.115.111989
- 4. Wang et al, 2019, DOI: 10.1002/mnfr.201900612
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- 6. Lindqvist et al, 2021, DOI: 10.1093/jn/nxaa019
- 7: Schmidt et al. 2021. DOI: 10.3390/metabo11020121 8. Playdon et al, 2017, DOI: 10.3945/ajcn.116.144501
- 9. McCullough et al, 2019, DOI: 10.1093/ajcn/ngy385
- 10. Tong et al, 2020, DOI: <u>10.1093/jn/nxz263</u>
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- 14: O'Sullivan et al, 2011, DOI: 10.3945/ajcn.110.000950
- 15: Bhupathiraju et al, 2018, DOI: 10.1093/jn/nxy074
- 16: Wei et al, 2018, DOI: 10.1002/mnfr.201800583
- 17: Stella et al, 2006, DOI: 10.1021/pr060265y
- 18: Rådjursöga et al, 2018, DOI: 10.3390/nu10081063
- 19: Djekic et al, 2020, DOI: 10.1161/JAHA.120.016518
- 20: Andersen et al, 2014, DOI: 10.1021/pr400964s
- 21: Khakimov et al, 2016, DOI:
- 10.1021/acs.jproteome.6b00109
- 22: Acar et al, 2019, DOI: 10.1002/mnfr.201800215
- 23. Vázquez-Fresno et al, 2015, DOI: 10.1021/pr5007894
- 24. Meslier et al, 2020, DOI: 10.1136/gutjnl-2019-320438
- 439 metabolites reported across studies (mean $S_M = 0.89$)
- 46 metabolites associated to AP or PP-rich diet with a high level of evidence $(S_M > 1,51)$
- **High heterogeneity** among the reported metabolites
- Metabolites related to amino acid and protein metabolism have the highest contribution
- proteinogenic amino acids (30% of the 46 metabolites)
- Biomarkers of animal products intake
- TCA cycle intermediates
- Very few lipids

Method:

1. Criteria for study selection

Metabolomics signatures

- Metabolomics analyses only
- Urine, plasma and serum
- Human only
- Observational and interventional studies

AP and PP intake

- Direct comparison of AP and PP intake
- Comparison of diets with contrasted AP and **PP** intake:
- o omnivores with vegetarians or vegans
- o adherence levels to defined dietary patterns (Mediterranean diet, HEI, DASH)
- o a posteriori (data-driven) dietary patterns

Protein package

Plant proteir

Dietary pattern

Protein package

Environment

INRAO

: 6:

U. ...

2. Scoring system for study contribution

Dietary contribution score S_P to account for the expected AP and PP intake

Type of dietary comparison	S _P
Comparison of a posteriori (data-driven) dietary patterns	0.25 (+0.125)
Comparison based on adherence to defined dietary patterns	0.5 (+0.125)
Comparison of omnivore, vegetarian and vegan diets	0.75 (+0.125)
Comparison based on AP and PP intake	1

differences across studies:

Type of dietary comparison	S _P
Comparison of a posteriori (data-driven) dietary patterns	0.25 (+0.125)
Comparison based on adherence to defined dietary patterns	0.5 (+0.125)
Comparison of omnivore, vegetarian and vegan diets	0.75 (+0.125)
Comparison based on	1

+ 0,125 for studies providing detailed dietary data confirming intake differences in AP and PP-rich foods across groups

3. Metabolite selection

- Compilation of all metabolites reported as discriminant between groups in each study and cross-analysis between studies
- Attribution of a metabolite score **S_M** to each metabolite reported:

 S_{M} (metabolite A) =

 $\sum S_c$ (studies reporting metabolite A)

Selection of the metabolites with the highest S_M

4. Metabolite review

Literature review on the links between the selected metabolites and cardiometabolic risk

Assigning an overall contribution score S_c to each study included: $S_c = (S_p + S_p) / 2$

Literature review (not specific to

metabolomics)

Discussion: Main metabolites discriminant between AP and PP-rich diets and their potential implication in cardiometabolic risk

As identified across selected metabolomics studies

Lifestyle Human urine, plasma and serum metabolome Glycine Glutamine Isoleucine, Leucine, Valine ■ Tyrosine, Phenylalanine

- Short-chain acylcarnitines
- Glutamate

PNCA

Trimethylamine-N-oxide

Cardiometabolic 1 Increased level associated to AP or PP Possible association

Conclusions:

- AP and PP-rich diets are associated with distinct metabolites which are plausible candidates to explain some of the differential associations between diet type and cardiometabolic risk
- Very little data is available regarding the metabolomics signatures specifically associated to **AP** and **PP** intake
- More studies with a specific focus on protein type, detailed dietary data and tight intake control (see poster **P28-011-21**) are needed to better characterize the metabolomics signatures associated to AP and PP intake

Want to know more?

Article in press for publication in Advances in Nutrition: Lépine et al, A scoping review: metabolomics signatures associated with animal and plant protein intake and their potential relation to cardiometabolic risk

Study design contribution score S_D to

account for the difference in evidence

level providing by the study design:

0.2

0.4

0.8

1

Study design

Mechanistically based

reasoning

Case-control studies

Non-randomized

controlled cohort

Randomized trials

Systematic review of

randomized trials