







Metabolomic fingerprinting of phenolic compounds in blood serum from rats treated with chestnut shells extract

Diana Pinto, Andreia Almeida, Anallely López-Yerena, Bruno Sarmento, Rosa Lamuela-Raventós, Anna Vallverdú-Queralt, Cristina Delerue-Matos, Francisca Rodrigues*

*francisca.rodrigues@graq.isep.ipp.pt





State of art



✓ 1/3 of food production is wasted





Principal causes



- √ 88 M tonnes are generated annually
- ✓ Associated costs estimated at 143 billion euros







Sustainability Challenge















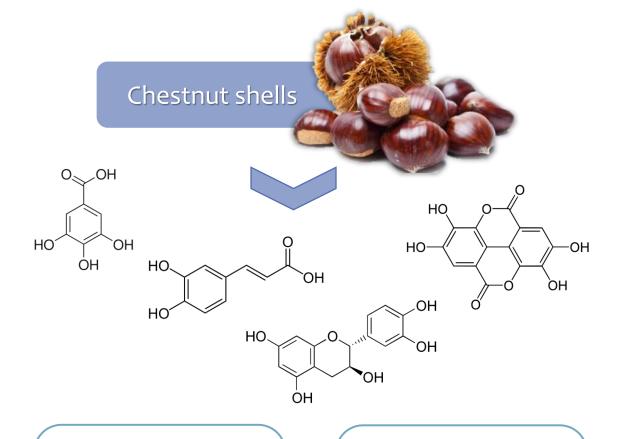
State of art



Macronutrients & Micronutrients



Vitamins Mineral **Polyphenols**



In vitro

Antioxidant
Anti-inflammatory
Hypoglycemic
Hypolipidemic

In vivo

Anti-obesity Anti-inflammatory Hepatoprotective

enols in blood

Objectives

Investigate the targeted metabolomic profile of polyphenols in blood serum from rats orally treated with an eco-friendly chestnut shells extract

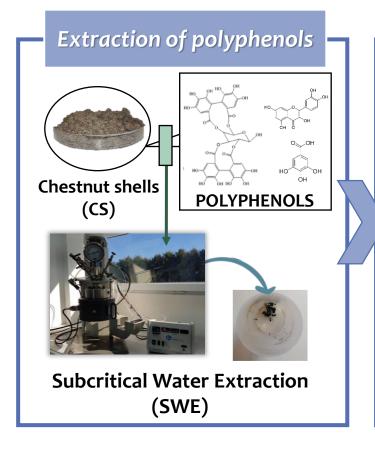


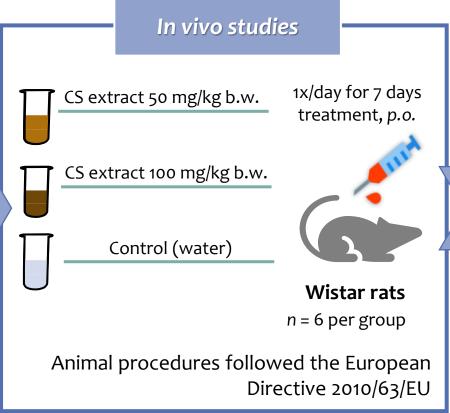


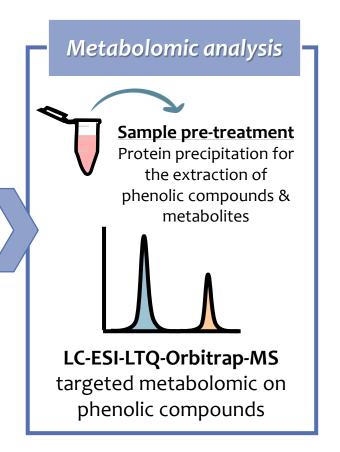
LC-ESI-LTQ-Orbitrap-MS

This is the **first study** that proposes a comprehensive analysis of the **metabolomic fingerprinting of phenolics-enriched chestnut shells** extract in **blood serum** after oral treatment of rats.

Experimental design



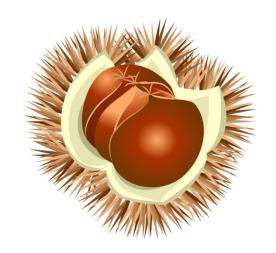




Phenolic acids



Hydroxybenzoic acids & Hydroxycinnamic acids



Flavonoids

Lignans

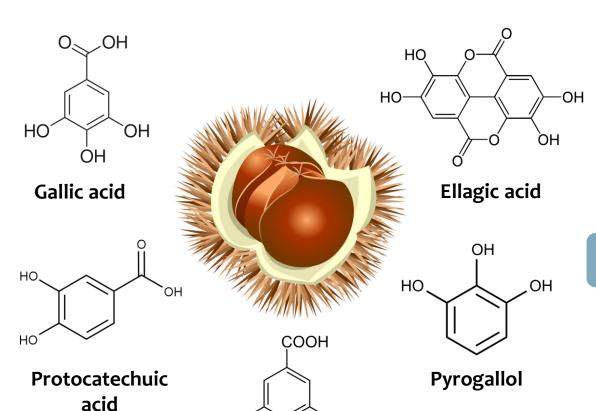
Other polyphenols

Phase I

Hydrogenation

Phase II

- Methylation
- Sulfation
- Methylation + Sulfation
 - Unmetabolized form



Phase II & gut microbiota

- Dimethylation
 - Unmetabolized form &

Urolithin A metabolites from methylation & sulfation

Phase II

- Sulfation
- Glucuronidation
- (Di)methylation + Sulfation or Glucuronidation

Syringic acid

H₃CO



Unmetabolized form

Phase II

Methylation

OCH₃

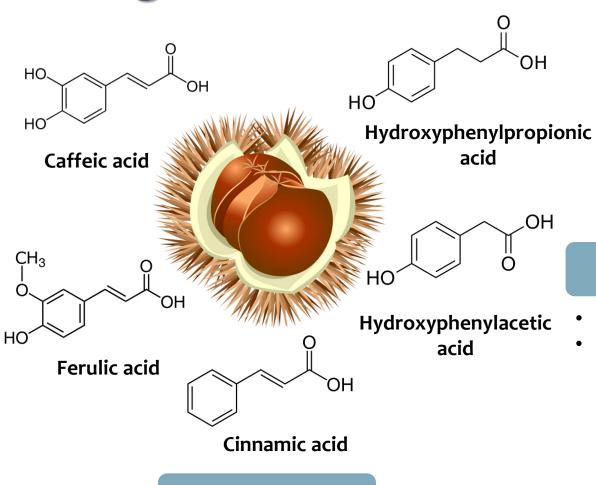
• (Di)methylation + Sulfation

Phase I & II

- Hydrogenation
- Hydrogenation + Sulfation or Glucuronidation
- Sulfation

Phase I & II

- Hydroxylation
- Hydrogenation
- Hydrogenation + Sulfation or Glucuronidation
- Methylation + Sulfation or Glucuronidation
- Hydrogenation + Methylation + Sulfation



Phase II

- Sulfation
- Dimethylation + Sulfation

Unmetabolized form

Phase II

- Sulfation
- Glucuronidation

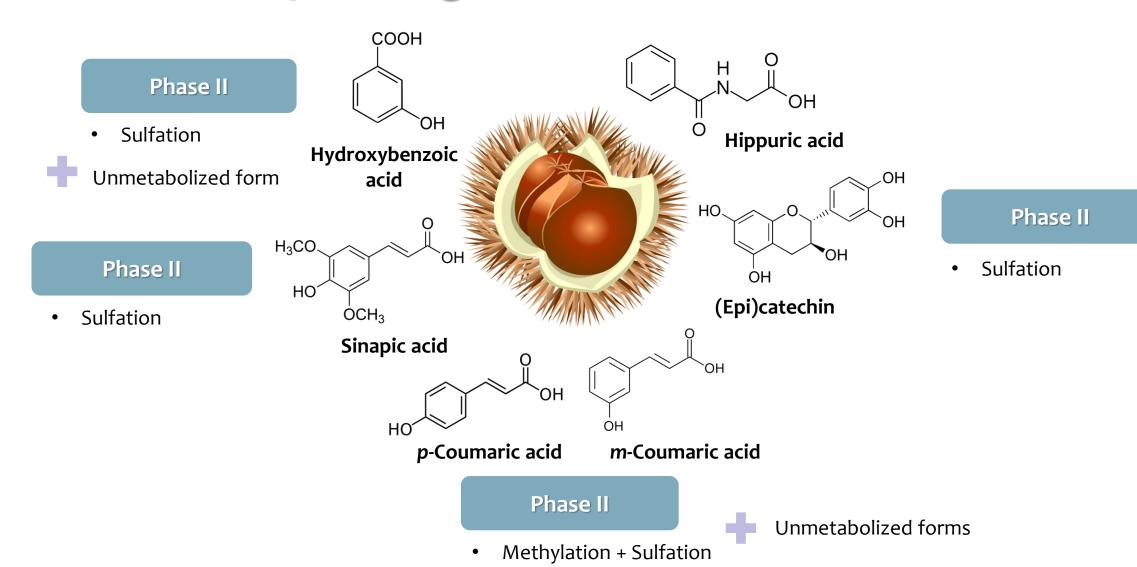
Phase II

Glucuronidation



Unmetabolized form

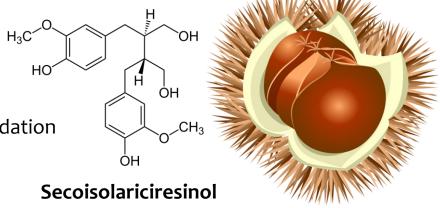
.OH



Phase II & gut microbiota

- Enterodiol + Disulfation
- Enterolactone + Glucuronidation

Unmetabolized form



ОН

Catechol

Phase II

- Methylation
- Glucuronidation
- Sulfation
- Methylation + Sulfation

- ✓ A total of 52 compounds were identified, mostly phenolic acids & metabolites.
- ✓ 80% of the metabolites resulted from **phase II** reactions; the remaining 20% derived from **phase I**.
 - ✓ 11 compounds correspond to **parent compounds**; the remaining represent their metabolites.

Conclusion



The **detection** of parent compounds in serum **attested their absorption** in **unmetabolized** form.



Phase II metabolites were secreted into circulating blood due to their high polarity and molecular weight.



Identical metabolomic profile for both CS extract groups.



This work validate for the first time a new nutraceutical ingredient extracted from chestnut shells.



Acknowledgments















- PTDC/ASP-AGR/29277/2017
- UIDB/50006/2020 & UIDP/50006/2020
- SFRH/BD/144534/2019
- CEECIND/01886/2020
- RYC-2016-19355













Thank you for the attention!