

Plasma Metabolomic Effects of 8-Week Red Wine Consumption in CHD Patients

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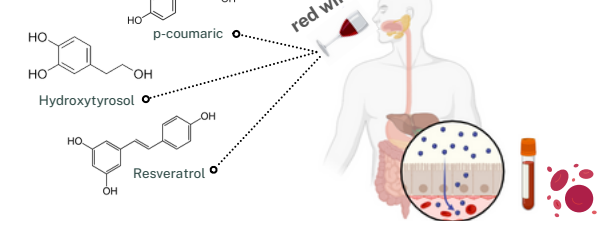
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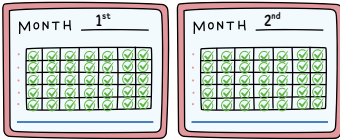
INTRODUCTION

Objective

To investigate the metabolic effects of **red wine consumption over 8 weeks**, focusing on the identification of key metabolites and their temporal changes, while **addressing inter-individual variability using multilevel analysis**.



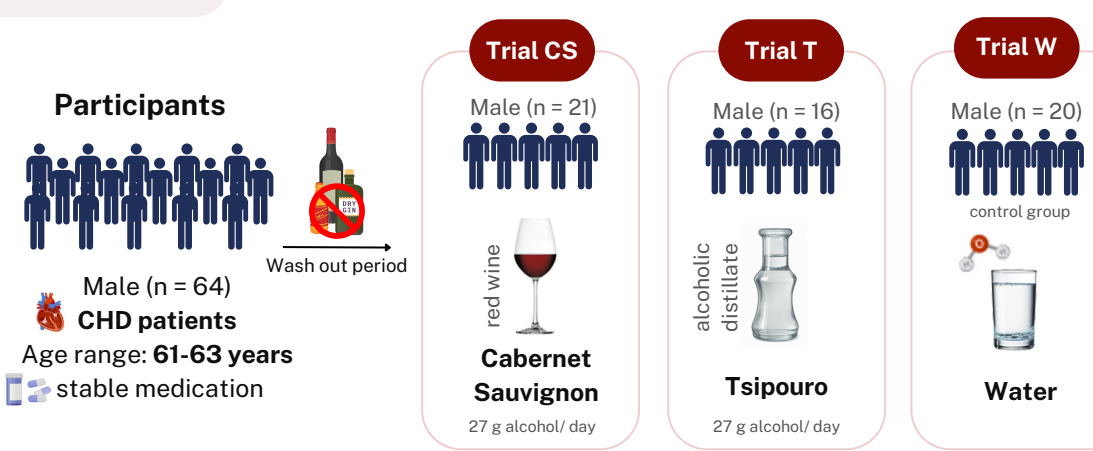
8 weeks of daily wine consumption



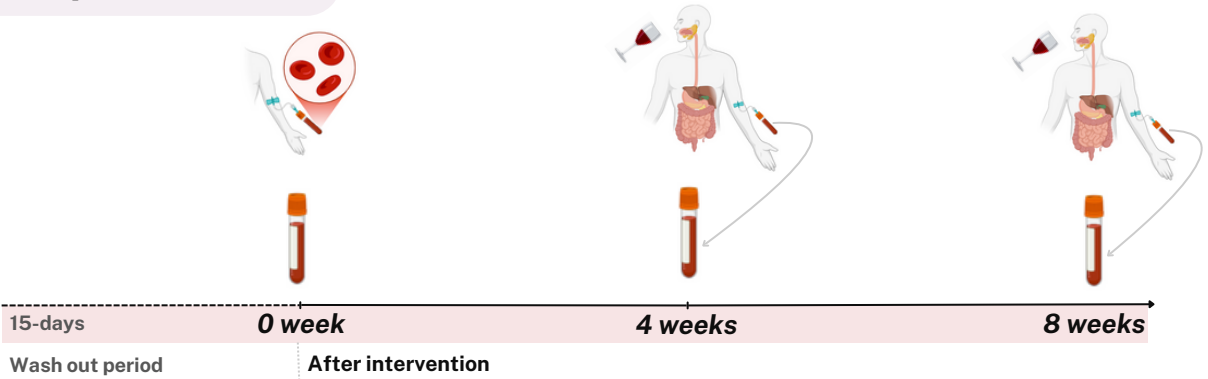
Moderate alcohol consumption, particularly **red wine**, has long been **associated with cardioprotective effects**, largely attributed to its polyphenolic content. However, the **long-term metabolic impact** of such consumption **in patients with coronary heart disease (CHD) remains poorly characterized**. To date, most studies have either focused on short-term interventions or observational associations without directly comparing ethanol versus polyphenol-rich alcoholic beverages. **This study addresses this gap by conducting the first 8-week randomized clinical trial using untargeted metabolomics**.

EXPERIMENTAL FLOW CHART

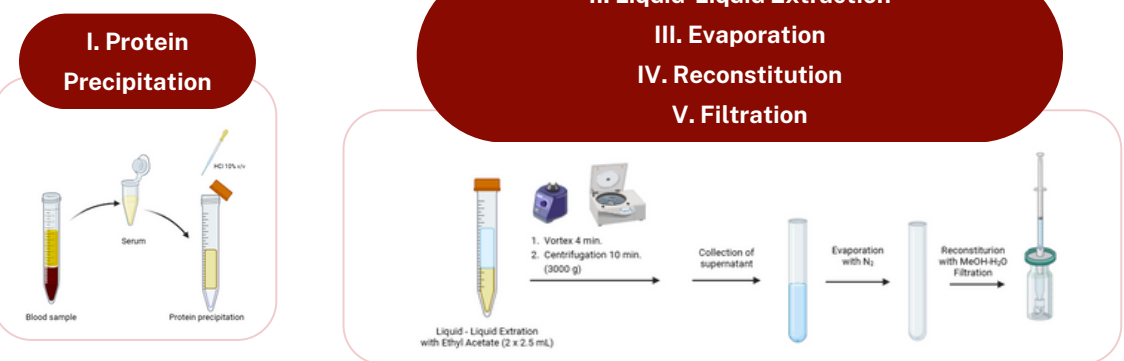
1. Clinical Trial



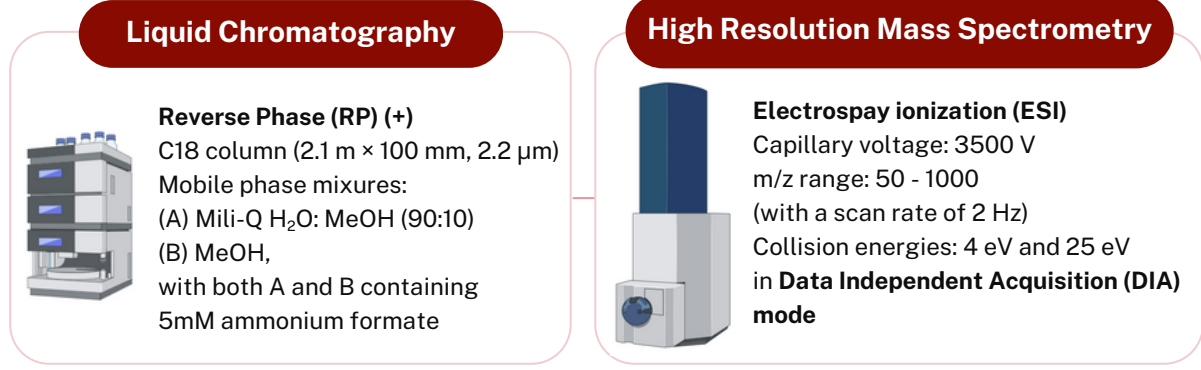
2. Sample Collection



3. Sample Preparation



4. Analysis



Data Processing

Non-targeted metabolomics data were acquired in AIF mode and processed in **MS-DIAL**. The dataset was **log-transformed** and **Pareto-scaled** in **RStudio** using **MetaboAnalystR**. Features with **RSD > 30%** were removed using **QC samples**. Multilevel analysis was performed with **mixOmics** and **MetaboAnalystR** to account for intra-subject variability.

RESULTS

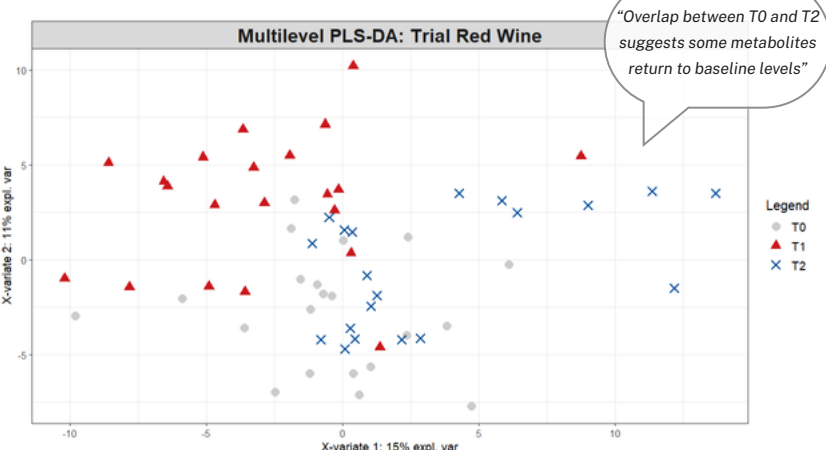


Figure A1. Partial least squares discriminant analysis (PLS-DA) of the red wine trial, with adjustment for inter-individual variability (multilevel).

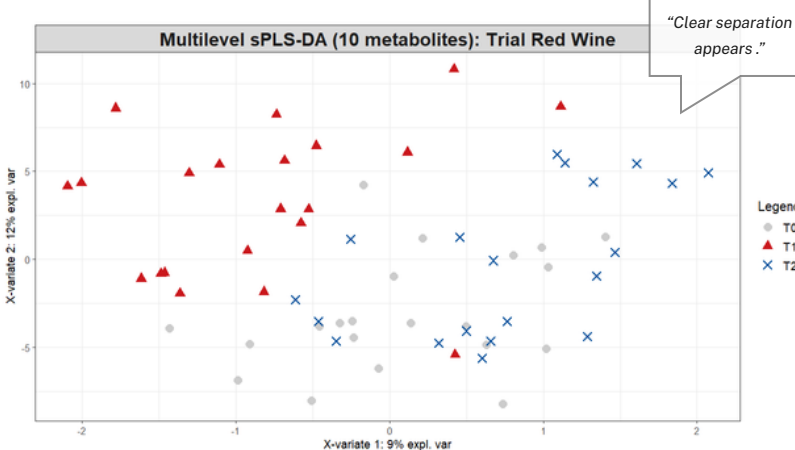


Figure A2. Multilevel sparse PLS-DA (sPLS-DA) of the red wine trial, with top 10 metabolites.

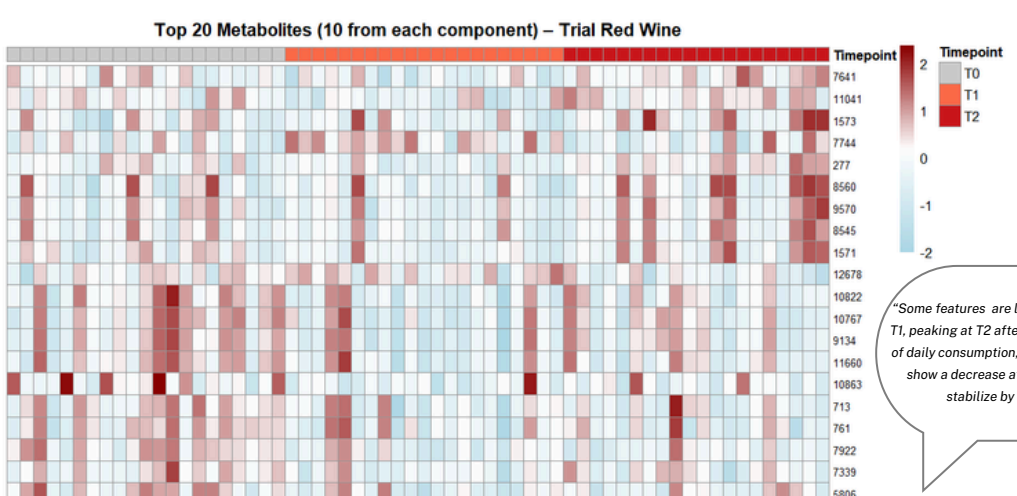


Figure B. Scaled heatmap of the top 20 discriminating metabolites identified by multilevel sPLS-DA (10 from each component), illustrating inter-individual and temporal variation across T0, T1, and T2.

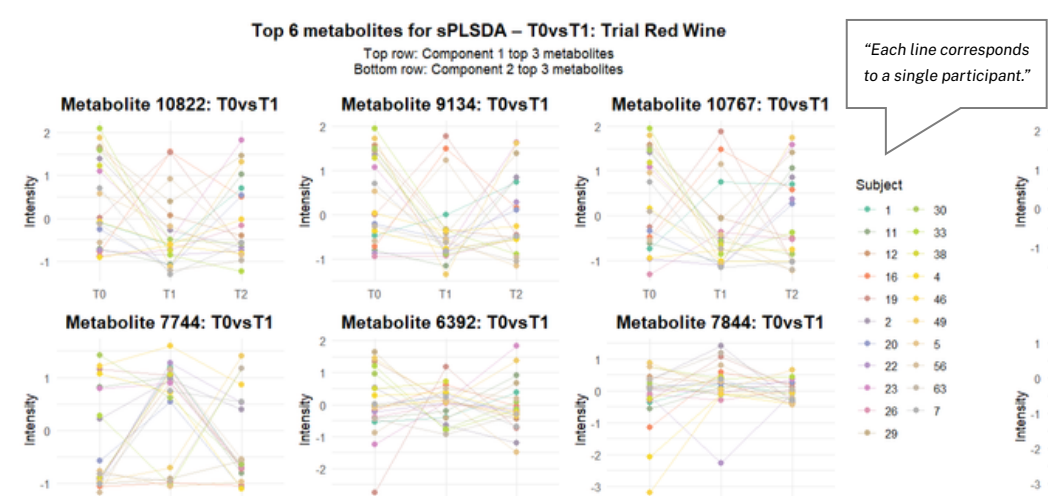


Figure C1. Longitudinal trajectories for the top 6 discriminating scaled features from the sPLS-DA comparison between T0 and T1.

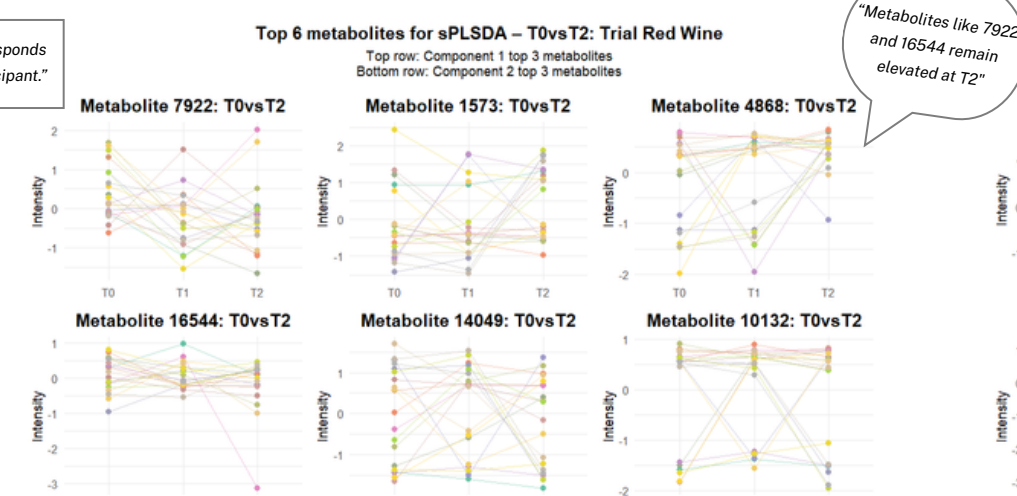


Figure C2. Temporal evolution of the top 6 discriminating features between T0 and T2 in the red wine group.

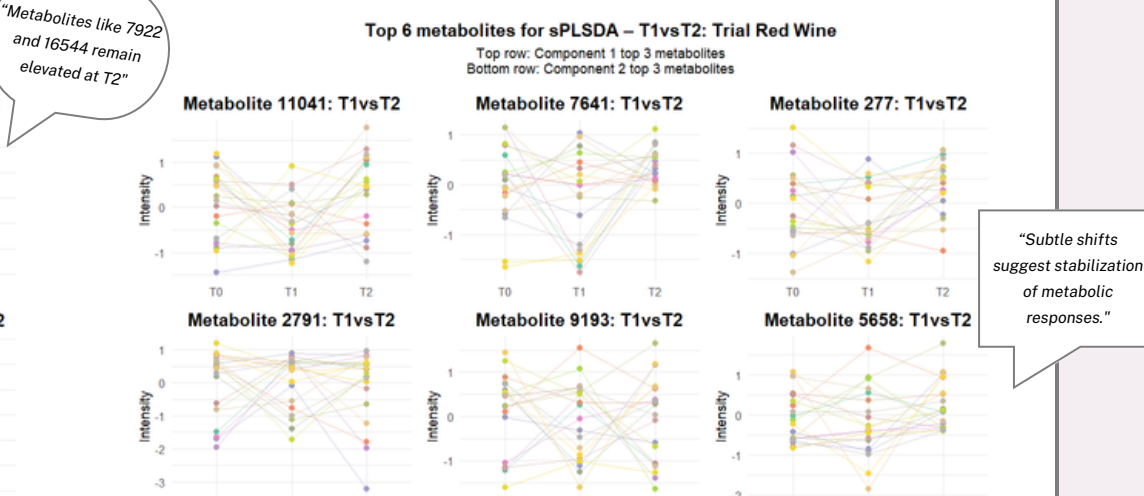


Figure C3. Inter-subject trajectories of the top 6 features distinguishing the 1st and 2nd month timepoints (T1 vs T2).

CONCLUSIONS / FUTURE PERSPECTIVES

- Metabolic Changes:** Red wine consumption induces significant metabolic shifts, with key metabolites increasing at T2 and others decrease after T1.
- Inter-Individual Variability:** Substantial variability across subjects highlights the need for multilevel modeling to account for baseline differences.
- Metabolic Stabilization:** Some metabolites return to baseline (T0) by T2, indicating metabolic adaptation to red wine consumption.

"Future work will focus on annotating metabolites and integrating them into pathway analyses to uncover underlying metabolic mechanisms."

Related Literature

[1] J. M. Guilford and J. M. Pezzuto, "Wine and Health: A Review," **2011**
[2] M. Lombardo, A. Feraco, E. Camajani, M. Caprio, and A. Armani, "Health Effects of Red Wine Consumption: A Narrative Review of an Issue That Still Deserves Debate," **2023**
[4] E. Fragopoulou, M. Choleva, S. Antonopoulou, and C. A. Demopoulos, "Wine and its metabolic effects. A comprehensive review of clinical trials," **2018**
[5] P. Lekka, E. Fragopoulou, A. Terpou, and M. Dasenaki, "Exploring Human Metabolome after Wine Intake—A Review," **2023**

