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Jérémy Monteiro, Antoine Lefèvre, Oury Adeline, Diane Dufour-Rainfray, Lydie Nadal-Desbarats, et al.. Exploration of SCFA in human multi-compartments using 3-nitrophenylhydrazine-LC-MS. Journée de la Recherche Biomédicale Tours/Angers 2023, Dec 2023, Tours (FR), France. hal-04352671

#### HAL Id: hal-04352671 https://hal.science/hal-04352671

Submitted on 19 Dec 2023

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# Exploration of SCFA in human multi-compartments using 3-nitrophenylhydrazine-LC-MS

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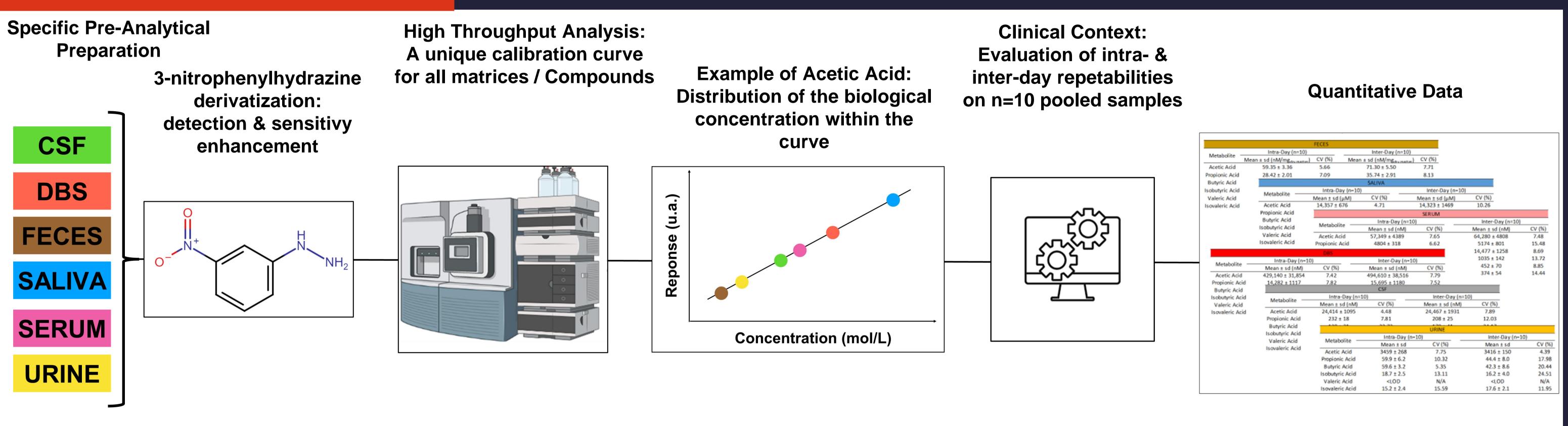
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### Introduction

Short-chain fatty acids (SCFAs) are metabolites involved in many physiological processes. A disruption of their metabolism is suspected in diseases of the digestive system (inflammatory diseases) but also in neurodegenerative diseases such as Alzheimer's disease. Currently, one of the reference methodologies for the quantitative analysis of SCFA is based on the use of GC-MS due to its sensitivity and reproducibility. However, its technical implementation is delicate, limiting in particular the throughput of the analyzes carried out. Our objective is to propose a method for absolute quantification of SCFA that is easier to implement and for future use in a clinical environment for a large number of types of biological samples (stool, urine, serum, saliva, CSF, and dried blood spot).

With this objective we selected, 3-nitrophenylhydrazine (3NPH) to promote the detection and analytical separation of these metabolites. After methodological validation (robustness, repeatability, matrix effect, carryover, LLOQ), its use makes it possible to propose reference values of SCFA concentrations for each type of sample. This quantitative methodology also allows longitudinal monitoring of SCFA concentrations.

## Material & Methods



### Results

(1) Comparison of biological concentrations of SCFAs

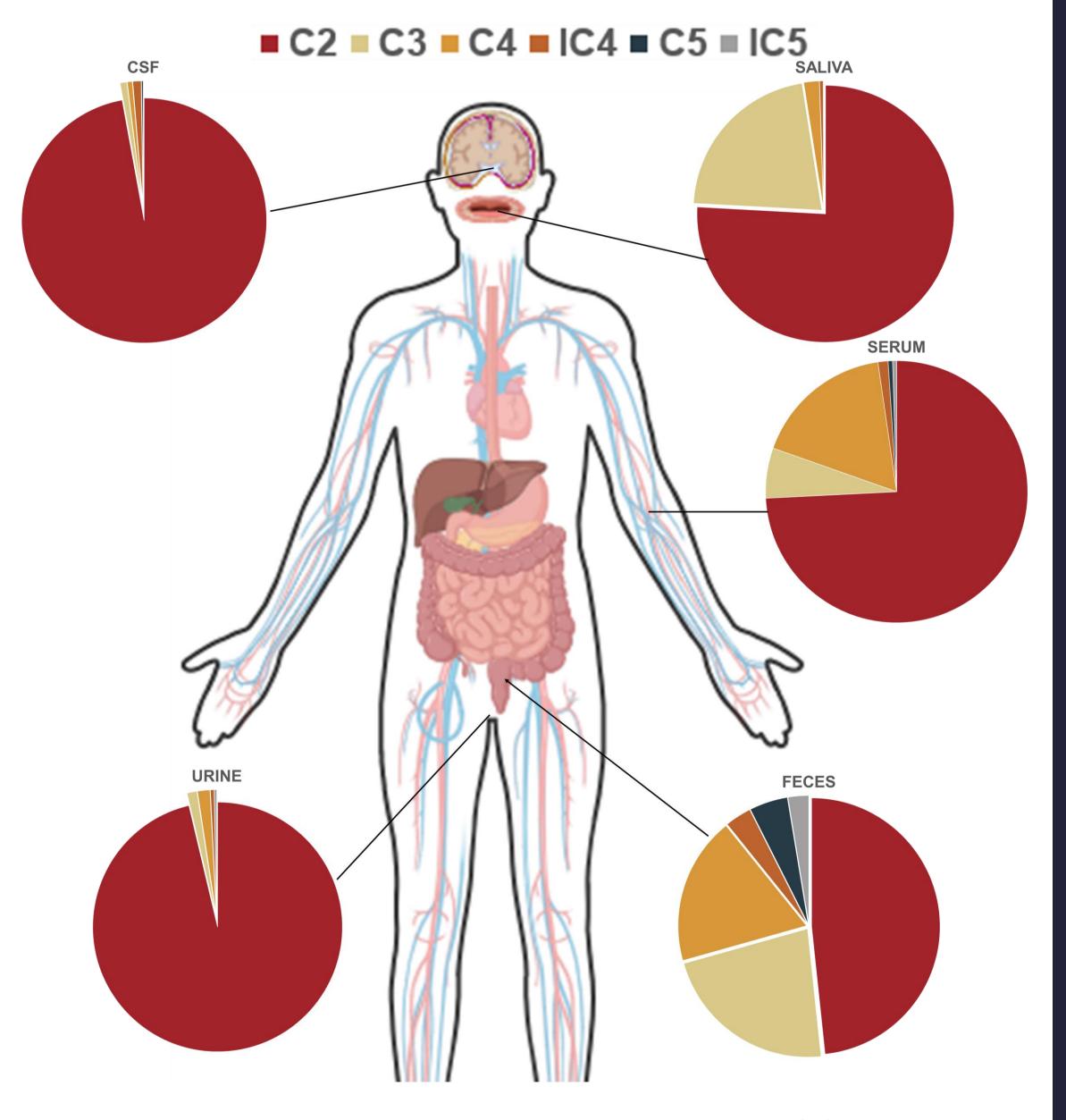
#### (2) Biological repartition of SCFAs in human

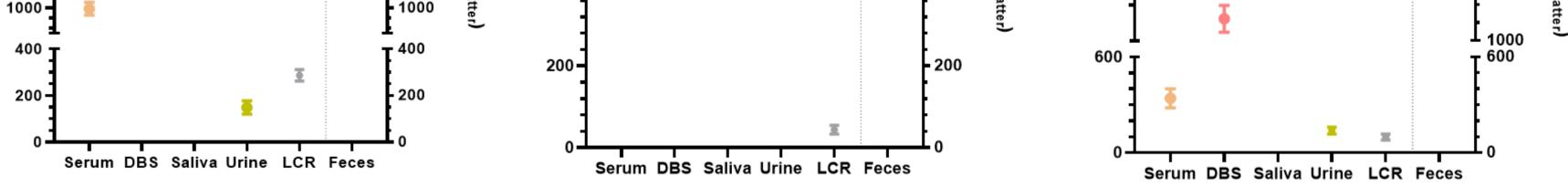
1. concentration in faces are normalized to dried matter

1: concentration in feces are normalized to dried matter 2: concentration in urine are normalized to [creatinine]

1.8×10<sup>7</sup> 1.8×10 400000 4000000 4000000 1.6×107 1.6×107 360000 360000 3000000 3000000 320000 320000-1.4×10<sup>7</sup> 1.4×107 30000 30000 1.2×107 **1.2×10**<sup>7</sup> 35000 35000 520000 520000· -25000 25000 لي 470000-25000 25000 470000 -20000 20000 420000 420000 15000 15000 80000 80000 15000 15000 Con 60000 - 5000 5000-60000 10000 10000 **-** 600 600 40000 40000 400 400 - 5000 5000 20000 20000 600 -200-200 Serum DBS Saliva Urine LCR Feces Serum DBS Saliva Urine LCR Feces Serum DBS Saliva Urine LCR Feces 10000 1000 100000 100000 80000 80000 90000 90000 8000 8000 80000 80000 70000 L 70000 6000 6000 **-** 6000 5000 6000 5000 5000 4000 الا 4000 4000-4000 4000 2000 - 2000 3000 3000 3000 2000 - 2000 - 2000 2000-- 400

(One pie chart = one matrix; the proportion is based on the concentration of compounds within the matrix studied)





Comparison of different SCFA concentrations between biological matrices, from healthy patients. The results shown gather intra-day and inter-day repeatabilities. The concentrations found in saliva are in  $\mu$ M (left axis) whereas they are expressed in nM for the other matrices.

Acetic acid is always the most concentrated SCFA, to the point of obscuring the others in the CSF and urine. In the faeces, all the SCFAs are concentrated: these are the main products of digestion. Note the high concentrations of propionic acid in saliva, produced by the enzymes it contains.

## Conclusion

SCFAs have been extensively and intensively studied by 3-nitrophenylhydrazine-LC-MS<sup>2</sup> derivatization due to their involvement in metabolisms and diseases. We propose a unique methodology to quantify SCFAs in 6 different biological matrices to generate FAIR data. Reference values are proposed for CSF and DBS. These validated approaches are now used to characterize metabolism and better investigate the inflammatory component of several diseases.