

### Development of high throughput miniaturized robotic platforms for metabolism investigations of microorganisms

Cécilia Bergès, Edern Cahoreau, Lindsay Peyriga, Stephanie Heux, Floriant Bellvert, Jean-Charles Portais

#### ▶ To cite this version:

Cécilia Bergès, Edern Cahoreau, Lindsay Peyriga, Stephanie Heux, Floriant Bellvert, et al.. Development of high throughput miniaturized robotic platforms for metabolism investigations of microorganisms. Biosynsys, Jul 2023, Toulouse, France. hal-04207380

#### HAL Id: hal-04207380 https://hal.science/hal-04207380

Submitted on 14 Sep 2023

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

# Development of high throughput miniaturized robotic platforms for metabolism investigations of microorganisms



## <u>Cécilia Bergès<sup>1,2</sup>, Edern Cahoreau<sup>1,2</sup>, Lindsay Peyriga<sup>1,2</sup>, Stéphanie Heux<sup>1</sup>, Floriant Bellvert<sup>1,2</sup> and Jean-Charles Portais<sup>1,2,3</sup></u>

1 : TBI, Université de Toulouse, CNRS, INRAE, INSA, Toulouse. France

2: MetaToul-MetaboHUB, National Infrastructure of Metabolomics & Fluxomics (ANR-11-INBS-0010)

3 : RESTORE, Université Toulouse III - Paul Sabatier, EFS, Institut National de la Santé et de la Recherche Médicale, Centre National de la Recherche Scientifique

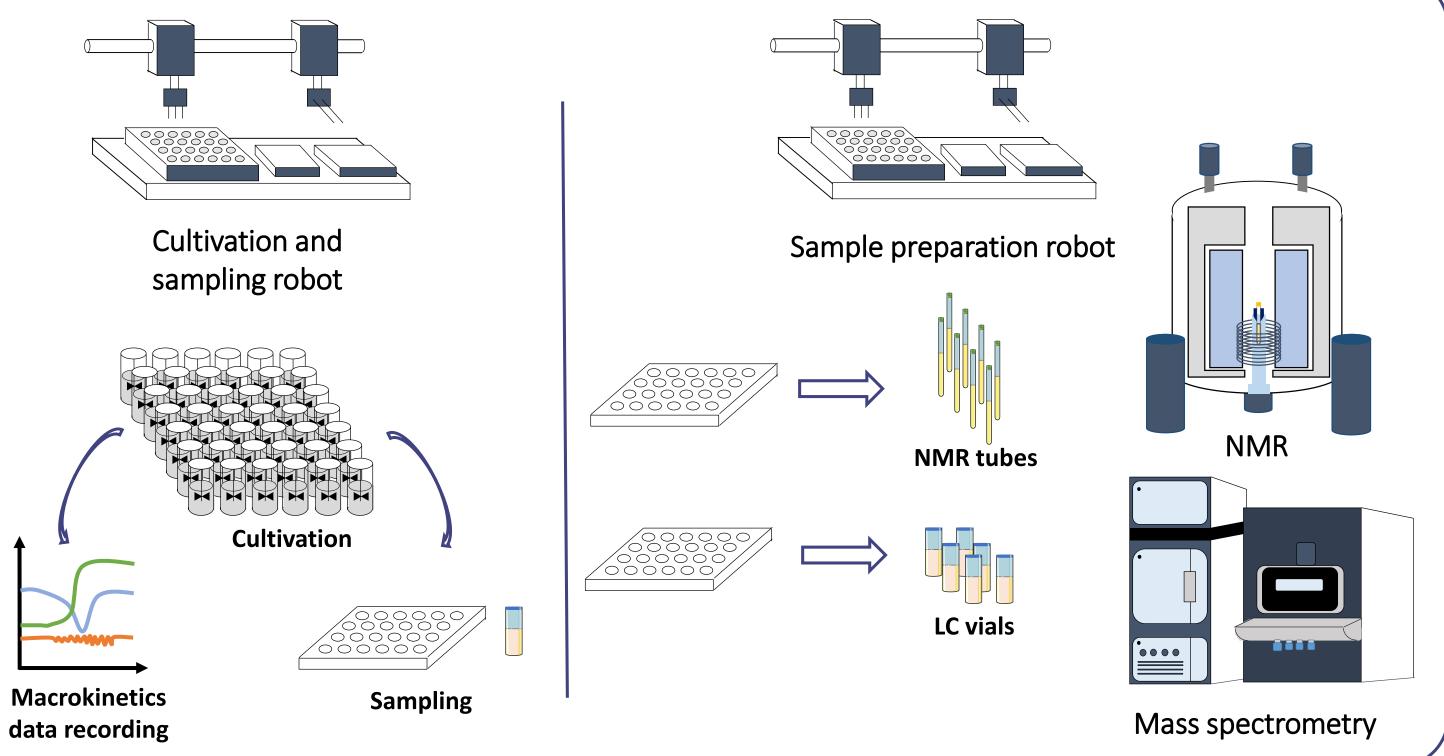
**Context : MetaboHUB MetaToul platform : Robotic facilities** 

High-thoughput (HT) omics approaches enable the large scale analysis of metabolism and its regulation. Among those, fluxomics (metabolic flux analysis) aims at measuring the actual rates of biochemical reactions in metabolic networks.

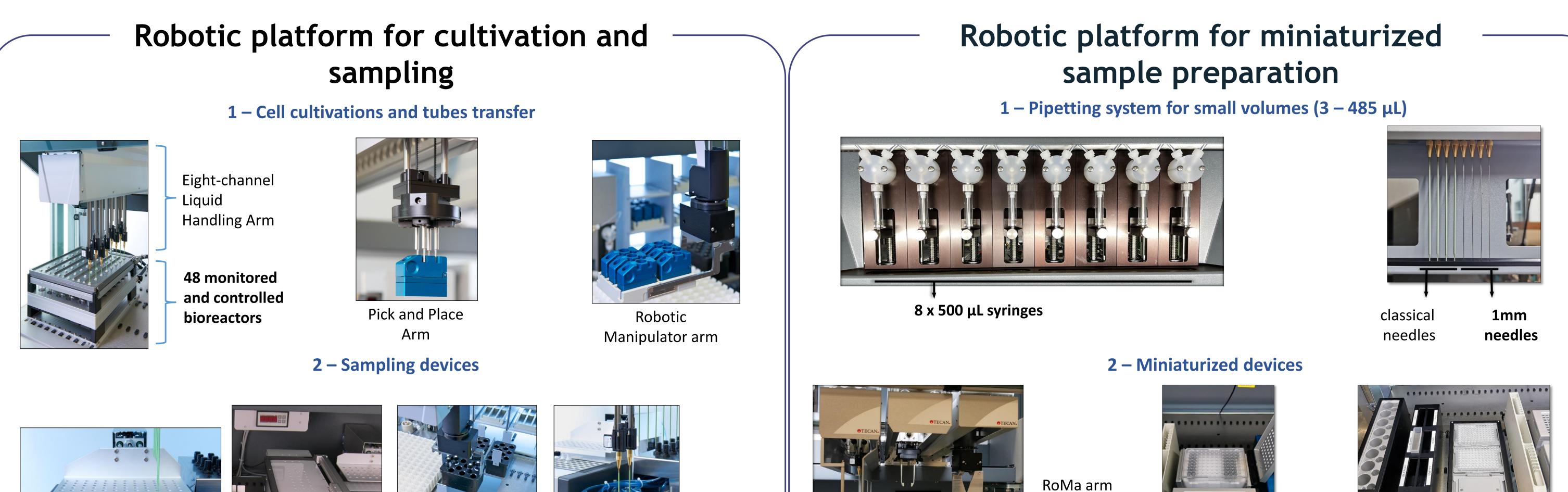
MetaToul develops and provides to the community a full methodological framework for highthroughput (HT) metabolism investigation applicable to a broad range of micro-organisms and applications. It includes experiment design and bioinformatical tools, analytical platforms (MS and NMR), and robotic platforms for cultivation and sample preparation.

In this poster, you'll find a focus on the possibilities offered by our two robotic platforms :

An automated, high-throughput cultivation and sampling workstation that allows to perform ulletrobust and HT metabolomics and fluxomics investigations in controlled physiological conditions.



An sample preparation workstation for low-volume handling or small-quantity samples in order to analyze matrices for which we have few amounts available or to concentrate classical samples and have access to minority metabolites.



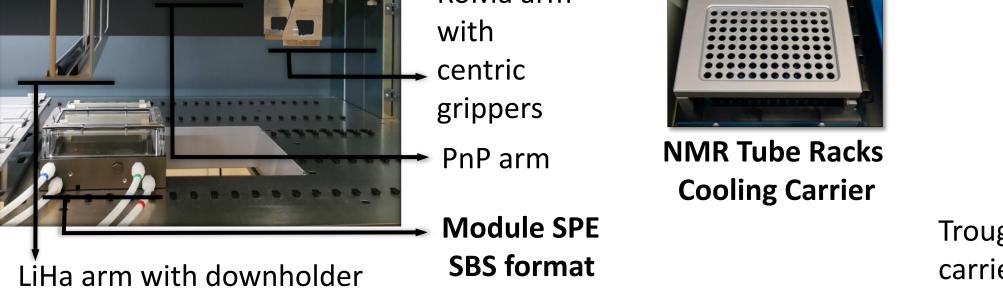


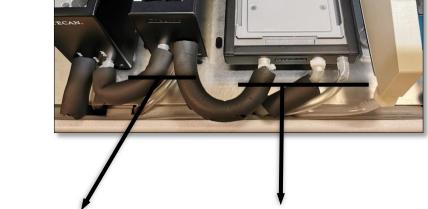
**Cooling module** 

Centrifuge

Filtration module

This robotic platform has the capability to run **48 cultivations in parallel** and **monitor pH, O2 and OD**<sub>600</sub> (optical density). This system also runs fully automated metabolite harvesting and extraction for downstream analyses.





Troughs cooling **Customized Inserts for** carrier SBS format

This robotic platform is capable of handling a complete range of sample racks (NMR tubes, vials, multiwell plates, vials, cryotubes, etc.), as well as carrying **out various types of manipulation** such as dilution, aliquoting, addition of buffer and internal standard, all while ensuring traceability of all samples.

APPLICATIONS OF AUTOMATIC HIGH THROUGHPUT WORKFLOW			
Monitoring physiology	HT Quantitative metabolomics	HT Fluxomic experiments	<b>Customized experiments</b>
<ul> <li>48 cultivations in parallel</li> </ul>	<ul> <li>48 cultivations in parallel</li> </ul>	<ul> <li>48 cultivations in parallel</li> </ul>	<ul> <li>48 cultivations in parallel</li> </ul>
<ul> <li>Monitoring of :</li> <li>OD600 every 10 min</li> <li>pH and pO2 every 1 min</li> </ul>	<ul> <li>Monitoring of :</li> <li>OD600 every 10 min</li> <li>pH and pO2 every 1 min</li> </ul>	<ul> <li>Monitoring of :</li> <li>OD600 every 10 min</li> <li>pH and pO2 every 1 min</li> </ul>	<ul> <li>Monitoring of :</li> <li>OD600 every 10 min</li> <li>pH and pO2 every 1 min</li> </ul>
<ul> <li>Collection of 48 culture medium all along the cultivation kinetic</li> </ul>	<ul> <li>Collection of 48 whole broth and filtrate when OD target is achieved</li> </ul>	<ul> <li>Collection of 48 culture medium all along the cultivation kinetic</li> </ul>	<ul> <li>Collection of supernatant or extract all along the cultivation kinetic</li> </ul>
<ul> <li>NMR analysis to determine by- products production/consumption</li> </ul>	<ul> <li>Mass spectrometry analysis to determine intracellular pools : amino acids, nucleotides, Krebs cycle, glycolysis and pentose phosphate pathway intermediates.</li> </ul>	<ul> <li>Cells harvesting at user defined OD600</li> <li>Mass spectrometry analysis to determine carbon isotopologues distributions</li> </ul>	<ul> <li>Sample preparation of your samples</li> </ul>
	L	<ul> <li>NMR analysis to determine by-products production/consumption</li> </ul>	

## Conclusions

This two systems allows to significantly accelerate the applications of metabolomics and fluxomics to investigate microbial metabolism, and to address challenges not only in metabolic engineering, synthetic biology and biotechnology but also in health, toxicology and pharmacology, biomarker and drug development.

**Customized protocols** can be designed on the two robotic platforms

Applications :

- Large-scale analysis of strains, mutants, species
- Large-scale analysis of the effects of environmental parameters, biological effectors, drugs, etc.
- **Design and metabolic optimization** of cell factories

Poster presented at Biosynsys 2023. Toulouse. France

## Acknowledgments

MetaboHUB (MetaboHUB-ANR-11-INBS-0010)

