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## CHU AMBROISE PARÉ

## Background

Breast cancer (BC) is the most common diagnosed cancer and the leading cause of cancer death in women worldwide, accounting for 25% of all cancer cases and 15% of all cancer deaths among females. There is an obvious need for a better understanding of BC biology.

The shift in cellular metabolism, also called Warburg effect, is an emerging hallmark of cancer cells (CC) orchestrated by oncogenic proteins. MIF, a pleiotropic cytokine, is involved in BC cells proliferation and multiple aspects of carcinogenesis but little is known about its possible role in cellular metabolic activities.

Metabolomics, the global qualitative and quantitative evaluation of metabolites in a biological system, by NMR spectroscopy, mass spectrometry or combined techniques, has emerged as a unique tool to investigate the modification of metabolites of cancer cells or in bio fluids and tissues of cancer patients.

Using MIF knockdown (KD) technique and <sup>1</sup>H-RMN spectroscopy, we evaluated the influence of MIF on BC triple receptor negative MDA-MB-231 and hormones receptors positive MCF-7 metabolism cells lines in the hope of identifying therapeutic targets.



## **Metabolites extraction**

After a quick rinsing, cells were scrapped and quenched using cold methanol.

The solution containing the quenched cells was pipetted for intracellular metabolites extraction (methanol, chloroform and water extraction procedure).

In this study, only the aqueous phase containing water soluble low-molecularweight endogenous metabolites was used. Prior to <sup>1</sup>H-NMR analysis, solvents were completely removed using a vacuum concentrator. Before acquisition on a Bruker Avance spectrometer at 500 MHz, each sample was reconstituted in phosphate buffer mixed with TSP.



# Macrophage migration inhibitory factor (MIF) involvement in breast cancer cell metabolism: A<sup>1</sup>H-NMR spectroscopy evaluation

V. Richard<sup>1-4\*</sup>, R. Conotte<sup>2,4</sup>, N. Kindt<sup>3</sup>, S. Saussez<sup>3</sup>, J-M. Colet<sup>2,4</sup> \*e-mail: vincent.richard@hap.be

<sup>1</sup> CHU Ambroise Paré, Mons Belgium ; <sup>2</sup> Department of Human Biology and Toxicology, Faculty of Medicine and Pharmacy, UMONS, Mons, Belgium; <sup>3</sup> Department of Anatomy and Cell Biology, Faculty of Medicine and Pharmacy, UMONS, Mons, Belgium ; <sup>4</sup> Biomedical Profiling Unit, UMHAP, Mons, Belgium







<sup>1</sup>H-NMR spectroscopy of intracellular MDA-MB-231 and MCF-7 cells metabolomes shows significant different profiles suggesting that MDA-MB-231 cells have higher levels of glucose and glutamine consumptions than MCF-7, producing more lactate. This suggests that MDA-MB-231 cells are more dependent on glutamine but less on lactate and, possibly, different responses to targeted metabolic drugs. Moreover the decrease levels of glutamine observed in both cells lines when MIF is under-expressed suggests that this cytokine could be involved in the regulation of this amino acid metabolism.

## Results

## Conclusion