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Miniaturisation of metabolomic and fluxomic workflow on spheroids cultures applied to the treatment of gastric cancer





Nina Lager-Lachaud^{1,2}, George Alzeeb⁴, Noémie Butin^{1,2,3}, Thibaud Lanté^{1,2}, Hanna Kulyk^{1,2}, Cécilia Bergès^{1,2}, Catherine Corcos⁴, Laurent Corcos⁴, Floriant Bellvert^{1,2}

¹ Toulouse Biotechnology Institute (TBI), Université de Toulouse, CNRS, INRAE, INSA, 31077 Toulouse, France

² MetaboHUB-MetaToul, National Infrastructure of Metabolomics and Fluxomics, Toulouse, France

³ RESTORE, Université de Toulouse, CNRS ERL5311, EFS, ENVT, Inserm U1031, UPS, Toulouse, France

⁴ INSERM, Faculté de Médecine, 29200 Brest, France

Introduction



Spheroid (3D) culture models are becoming essential to mimic what happens in tissues *in vivo*. They are more representative of the cell-cell contact, dynamic interactions and reduce the impact of plastic compared to traditional 2D cultures. According to the models of cultures, metabolic differences were observed (in glycolytic activity, ATP production and sugar nucleotides synthesis). Expression of drug resistance genes may be also different and spheroids are often more resistant to the drug than 2D cultures. Culture of single spheroids are used to analyse with high throughput a large number of conditions. The challenge of this project was to be able to perform a study of metabolism for single spheroid (metabolomics and ¹³C metabolomics) of gastric cancer cell line.

-> Objective : Development of automated workfow (sample preparation and mass spectrometry) to analyse a single spheroid

Material and methods



Results and discussion

Spheroids metabolism analysis

Spheroids metabolome coverage

Comparison with 2D cultures





 \rightarrow Validation of a high throughput sample preparation protocol applied to single spheroid.

-> Ability to analyze a single spheroid (400 cells injected) for absolute quantification and isotopic profiling by mass spectrometry.

-> Metabolic differences between 2D and 3D cultures : Higher level of most central metabolites in spheroids / Stronger impact of drugs on 2D than on 3D cultures.

 \rightarrow Non targeted analysis in process...

References

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