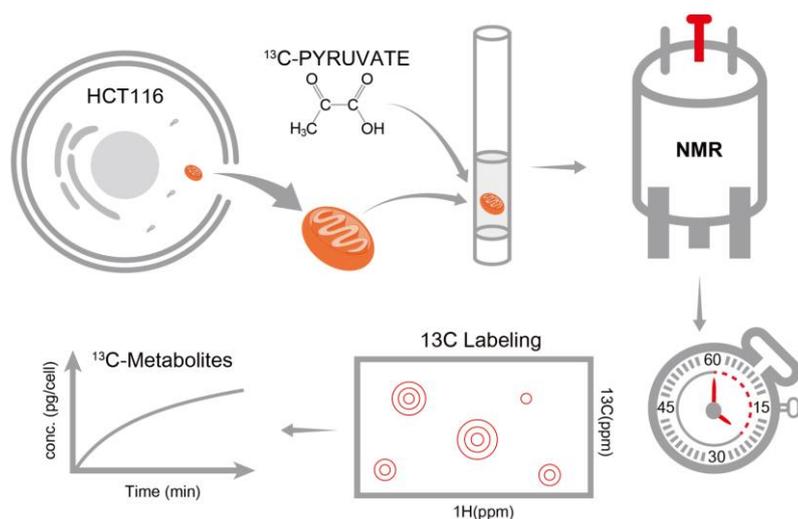


Real-time Live Metabolomics with 2D Heteronuclear NMR in Hierarchical Biological Systems

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Recent studies point out the link between altered metabolism and various diseases, but metabolic monitoring of biological systems have mostly been performed at fixed time points on extracted material. As NMR is a non-destructive technique, we applied heteronuclear 2D NMR to living or functional biological systems. At the protein level, we present a triple resonance NMR-based approach for specific detection of glutaminase activity using stable-isotope labeled glutamine in live cells. Compared to conventional methods involving coupled enzyme assays, the proposed approach is direct because it detects the presence of the H–N–CO amide spin system. The approach was applied to investigating the effects of a glutaminase inhibitor and the inhibitory effects of glucose on glutamine metabolism in live cells. The approach can be extended to other enzymes targeting carbon bonds using HCACO. At the organelle level, we monitored the mitochondrial metabolism in real time. We used ¹³C pyruvate as a tracer to monitor its metabolism in isolated and functional human mitochondria. We found acetyl phosphate synthesis from pyruvate in live human mitochondria which has been largely forgotten since its first description about 70 years ago in bacteria. The kinetic profile of acetyl phosphate formation was biphasic, and its transient nature suggested its role as a metabolic intermediate. The method also allowed for the estimation of pyruvate dehydrogenase (PDH) enzyme activity through monitoring of the acetylCoA formation, independent of competing cytosolic metabolism. We also detected lactate production from pyruvate in live mitochondria which is modulated by p53, a well-known tumor suppressor. Although lactate production has been known only in cytosol, the mitochondrial formation was confirmed with the recovery of lactate formation upon treatment of specific inhibitor of mitochondrial pyruvate carrier followed by a membrane permeabilizer. At the cell level, the live metabolomic method allowed for metabolomic differentiation between cancer and normal cells on the basis of time dependent changes in metabolite concentrations. Cancer cells were found to have large in- and out-flux of pyruvate as well as increased net production of alanine and acetate. The method also enabled evaluation of the metabolic effects of galloflavin whose anticancer effects have been attributed to its specific inhibition of lactate dehydrogenase. Our approach revealed previously unknown functional targets of galloflavin, which were further confirmed at the protein levels. How the live metabolomic method can be expanded to other hierarchical biological systems will be also presented.



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