Summer Meeting hosted by the Irish Section, 16-19 July 2012, Translational nutrition: integrating research, practice and policy

Development of a biomarker for sucrose intake based on metabolomics

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Carbohydrates are the major source of energy in the diet and their intake is an important determinant of health. The intake of extrinsic sugars, such as sucrose, is of increasing public health concern as evidence suggests high intakes are associated with metabolic abnormalities leading to hyperlipidaemia, insulin resistance and increased risk of degenerative disease⁽¹⁾. It is important, therefore, to have an accurate determination of dietary sucrose intake and metabolomics has recently been shown to be a useful tool for measuring dietary exposure⁽²⁾. The aim of this study was to develop biomarkers of sucrose intake using metabolomics.

We performed a randomised, parallel, single-dose feeding study on 90 healthy females (aged 29.9 ± 4.7 yr, BMI 23.3 ± 2.5 kg/m²) consuming either 0 g, 50 g or 100 g sucrose in 500 ml water (30 subjects per group). Blood and urine samples were taken before and for 24 h post sucrose intake. Standard low carbohydrate meals were provided during the study period. Blood was analysed for markers of glucose control and lipid status, urine was measured for electrolytes and sucrose excretion. Urine and plasma samples underwent FIE-MS fingerprinting analysis using established protocols⁽³⁾, followed by GCMS and LCMS analysis.

FIE-MS fingerprinting analysis showed that the 3 h time point was the most informative in urine and plasma (PC-LDA T_w values>2) and out of 120 explanatory signals, highlighted 12 metabolite signals in urine and 23 metabolite signals in plasma above a statistical significance threshold (p<1 × 10 $^{-4}$) that were discriminatory and either positively or negatively correlated with sucrose intake over time. In urine, the main signals positively correlated with sucrose intake were attributed to sucrose at 3, 6 and 9 h, and fructose and erythronic acid at 3 h. In plasma erythronic acid was also discriminatory and several unknowns were also positively correlated with sucrose intake. In urine and/or plasma samples several discriminatory signals attributed to keto- and hydroxy acids, medium to long chain non-esterified fatty acids and derivatives (acyl-carnitines), indicative of a fasting response, were highly abundant following consumption of water and were generally negatively correlated with sucrose intake, certain species increasing with time in the 0 g group (e.g acetylcarnitine, 3-OH-butanoic acid). GCMS and LCMS profiling analysis of urine and/or plasma samples confirmed fingerprinting data in that sucrose, fructose and erythronic acid were the strongest dose responsive discriminators for sucrose intake and metabolites associated with a fasting response being negatively correlated with sucrose intake.

In conclusion, we have found sucrose, fructose, and erythronic acid to be positively correlated with sucrose intake suggesting that these could be developed further into quantitative biomarkers of sucrose intake. A fasting-to-fed metabolic switch was also evident over time indicating that total energy intake during the study was low.

This work was supported by the Food Standards Agency (N05084) and the Department of Health (024/0044). Any views expressed are the author's own.

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