

Review of: "Regular Consumption of Lacto-fermented Vegetables has Greater Effects on the Gut Metabolome Compared with the Microbiome"

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The authors presented an interesting study investigating an under-researched aspect of diet modification with lacto-fermented vegetables for potential health benefits. While there is general recognition of this ancient tradition found in all cultures, details about mechanisms and effects of these foods on the human gut microbiome are sparse (1, 2).

Similar to the findings reported from the American Gut Project (3), in this study a substantial increase of LFV consumption to a daily amount of four servings (4 ounces) had only minor effects on the taxonomic composition or diversity of the microbiome comparing consumers and non-consumers. Authors are aware of generalisability concerns where in this small cohort (n=47) only some participants showed differences of abundances of certain taxa such as *Leuconostoc mesenteroides* and *Rhodotorula mucilaginosa*. Microbiome diversity as an indicator of health is increasingly being questioned: some studies found that healthier individuals have a greater diversity of taxa and functional pathways while others reported the opposite (4, 5). Intervention studies reported improvements of symptoms without increases of diversity or greater diversity in the unhealthy group compared to healthy controls (6, 7). This reiterates the necessity of the debate about diversity as a biomarker for health, let alone the issue that we are still yet to define a healthy microbiome (8).

Here, a shift from discussing a 'core of healthy species' to a 'core of functions' common to all human beings has changed our understanding of the influence of the microbiome. Guse et al. reported in this regard a significantly greater metabolomic diversity post intervention, particularly a potential increase of butyrate producers and a general increase of short-chain fatty acid production. SCFAs have been identified as beneficial to human health with their broad ranging effects as energy sources for enterocytes or signalling molecules for a wide range of physiological processes well beyond the intestinal tract (9). At the same time, we are alerted that some 95% of SCFAs are absorbed and amounts measured in faecal samples do not adequately represent production rates (10). Furthermore, there is no generally recognised level of adequate amounts of SCFAs. The reported increase of SCFA levels as a result of increased LFV consumption is nevertheless an intriguing fact requiring further investigation. The authors mentioned that it is not clear if the consumption of LFV itself caused the increase in SCFA levels or if it supported the existing population of producers to increase their production. One way or another it appears that an increased intake has the health benefit of increased SCFA levels for the individual. In addition, SCFAs are metabolites of fibre fermentation and higher levels found in the LFV group poses an intriguing question given that there were no differences in fibre intake between the groups. Authors hint that the lack of lactic acid consumed with the fermented vegetables may indicate its metabolization to butyrate. Elucidating the

mechanisms behind the higher abundances of valeric, butyric, and acetic acids in the LFV group would indeed be of great interest.

Potential health benefits from LFV consumption faces the well-known challenges of correlating metabolomic data with dietary data from self-administered questionnaires. Reporting bias and accuracy of study participants is one issue, the limitations of the assessment tool are another. More fundamental issues plague nutritional epidemiology where millions of possible combinations of food items and their ingredients seriously hamper classifications of a healthy food or diet. This had been hotly debated in the field for many years and is unlikely to be resolved anytime soon (11, 12). LFV consumption, its effect on the microbiome and potential health benefits could be assessed in much greater detail in controlled feeding studies, as the authors suggest. Their finding in the present study of a higher metabolite pool and higher abundances of SCFAs in the consumer group is encouraging for further research in this area.

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