Modulation of Gut Microbiota and Short-Chain Fatty Acid Production by Gac Fruit Juice and Its Fermentation in In Vitro Colonic Fermentation

Marisa Marnpae,^{a,b} Vernabelle Balmori,^{a,c} Kritmongkhon Kamonsuwan,^a Uarna Nungarlee,^b Suvimol Charoensiddhi,^d Thavaree Thilavech,^e Tanyawan Suantawee,^a Pavaret Sivapornnukul,^f Prangwalai Chanchaem,^f Sunchai Payungporn,^f Winai Dahlan,^b Nazimah Hamid,^g Thumnoon Nhujak,^h Sirichai Adisakwattana^{a,b*}

^aCenter of Excellence in Phytochemical and Functional Food for Clinical Nutrition, Department of Nutrition and Dietetics, Faculty of Allied Health Sciences, Chulalongkorn University, Bangkok 10330, Thailand. Email: Sirichai.a@chula.ac.th

^b The Halal Science Center, Chulalongkorn University, Bangkok 10330, Thailand

^c Department of Food Science and Technology, Southern Leyte State University, Southern Leyte 6606, Philippines

^d Department of Food Science and Technology, Faculty of Agro-Industry, Kasetsart University, Bangkok 10900, Thailand.

^e Department of Food Chemistry, Faculty of Pharmacy, Mahidol University, Bangkok 10400, Thailand

^f Center of Excellence in Systems Microbiology, Department of Biochemistry, Faculty of Medicine, Chulalongkorn University, Bangkok, 10330, Thailand

^g Department of Food Science, Faculty of Health and Environment Sciences, Auckland University of Technology, Auckland 1010, New Zealand

^h Department of Chemistry, Faculty of Science, Chulalongkorn University, Bangkok 10330, Thailand

Supplementary File

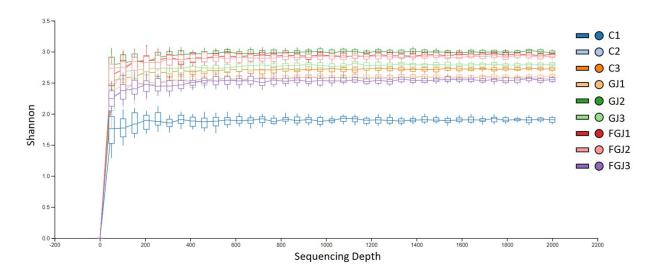


Fig. S1 Alpha rarefaction curve at 24h. Shannon indices of the nine groups kept steady indicating that the sequencing depth of the groups was adequate to characterize the composition of the microbial communities.

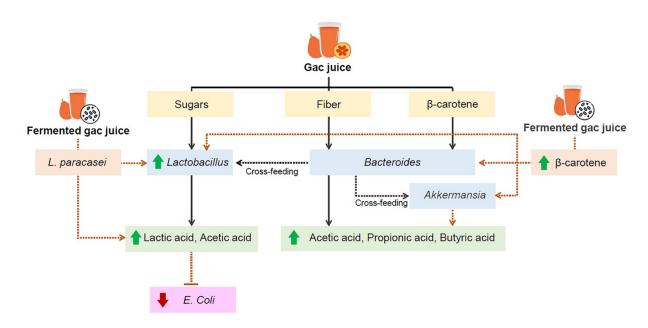


Fig. S2 Proposed Relationship between the composition of gac Juice (GJ) and fermented gac juice (FGJ) and mainly changes in gut microbiota and short-chain fatty acids (SCFAs).