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Author Correction: Gut microbiome composition in the Hispanic Community Health Study/Study of Latinos is shaped by geographic relocation, environmental factors, and obesity

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AUTHOR CORRECTION

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Author Correction: Gut microbiome composition in the Hispanic Community Health Study/Study of Latinos is shaped by geographic relocation, environmental factors, and obesity

Robert C. Kaplan^{1,2*}, Zheng Wang¹, Mykhaylo Usyk³, Daniela Sotres-Alvarez⁴, Martha L. Daviglius⁵, Neil Schneiderman⁶, Gregory A. Talavera⁷, Marc D. Gellman⁶, Bharat Thyagarajan⁸, Jee-Young Moon¹, Yoshiki Vázquez-Baeza^{9,10}, Daniel McDonald¹¹, Jessica S. Williams-Nguyen², Michael C. Wu², Kari E. North¹², Justin Shaffer¹¹, Christopher C. Sollecito³, Qibin Qi¹, Carmen R. Isasi¹, Tao Wang¹, Rob Knight^{10,11,13,14} and Robert D. Burk^{1,3,15}

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Following publication of the original paper [1], an error was reported in the third paragraph in the section “**Analysis of GMB composition and its correlates**” (page 3 of the PDF). The first sentence of the text should refer to Table 2, but mistakenly refers to Table 1.

The sentence should read:

“Fungal (ITS1) GMB populations were dominated by *Aspergillus proliferans* and *Saccharomyces cerevisiae* in both mainland US-born and Latin American-born groups (Table 2).”

As a result Table 2 appears further in the article, after the second reference to Table 2.

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