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# Corrigendum: Altered composition of the oral microbiota in depression among cigarette smokers: A pilot study

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avoidance, activation, BADS, metagenomics, oral microbiome

### A corrigendum on

Altered composition of the oral microbiota in depression among cigarette smokers: A pilot study

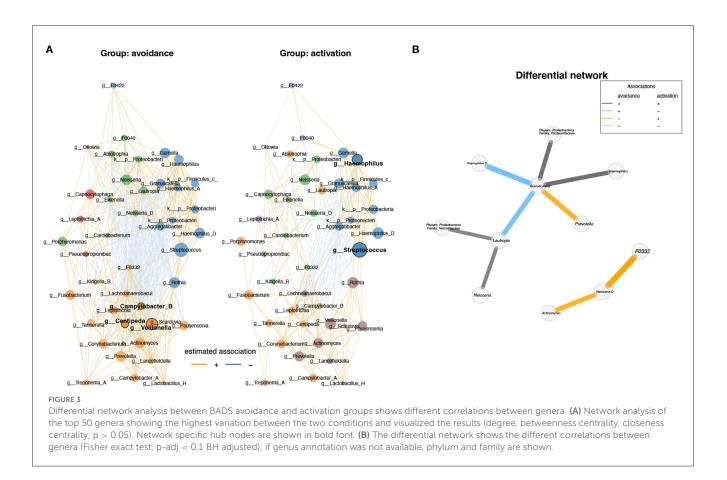
by Al Bataineh, M. T., Künstner, A., Dash, N. R., Abdulsalam, R. M., Al-Kayyali, R. Z. A., Adi, M. B., Alsafar, H. S., Busch, H., and Ibrahim, S. M. (2022) *Front. Psychiatry* 13:902433. doi: 10.3389/fpsyt.2022.902433

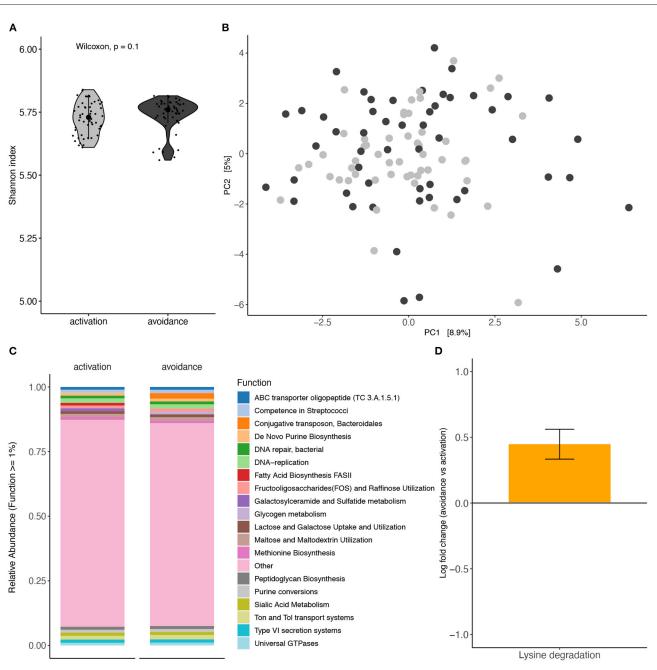
In the published article, there was an error in Figure 3 and Figure 4 as published. In the published article, the edges between nodes were not visible (Figure 3) and the figure/text was blurry (Figures 3, 4). The corrected Figure 3 and Figure 4 and their caption appear below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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## FIGURE 4

Functional profiling analysis of oral microbiota based on BADS. Functional profiling between BADS avoidance and activation groups (A) alpha diversity (Shannon index) and (B) beta diversity (Aitchison distance). (C) Average functional profiles with an abundance >1% are shown. (D) Lysine degradations different abundance between the two conditions (ANCOM-BC, q = 0.0692).