

CORRECTION

Correction: Hepatic glycogen storage diseases are associated to microbial dysbiosis

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Fig 3 is incorrect. The authors have provided a corrected version here.

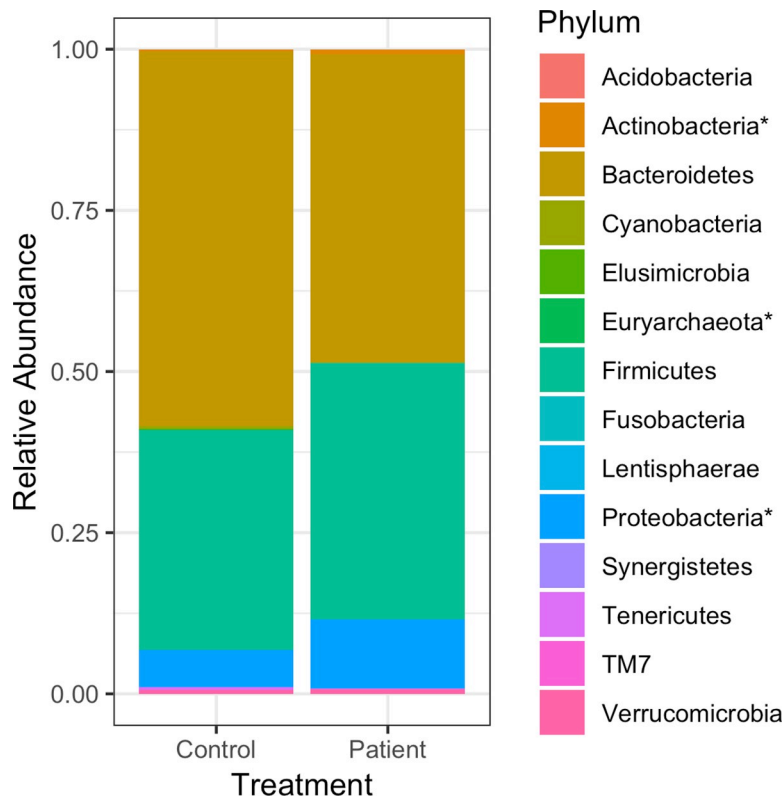


Fig 3. The average relative abundance of phyla found in GSD patients and healthy controls. Phyla followed by an asterisk (*) are different, both in terms of statistics and biological consistency, between patients and controls at p and $FDR \leq 0.05$: *Euryarchaeota* (LDA score = 1.75), *Actinobacteria* (LDA score = 3.06) and *Proteobacteria* (LDA score = 3.94). Firmicutes was marginally significantly different with $p = 0.064$, LDA score = 4.52 and $FDR = 0.112$.

<https://doi.org/10.1371/journal.pone.0218254.g001>

Fig 4 is incorrect. The authors have provided a corrected version here.



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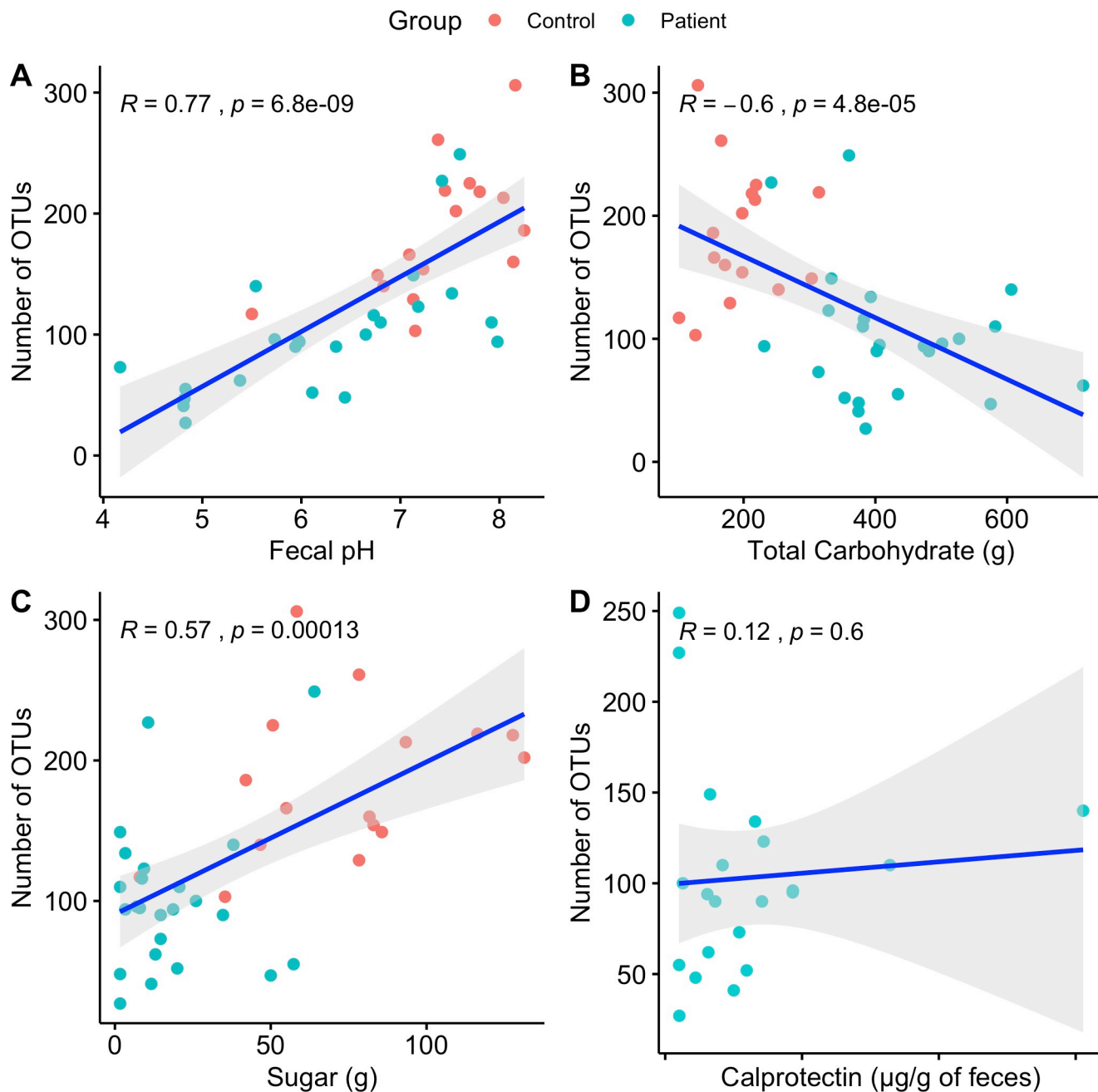


Fig 4. Correlations between the microbiota and diet, faecal pH, and gut inflammation.

<https://doi.org/10.1371/journal.pone.0218254.g002>

Reference

1. Colonetti K, Bento dos Santos B, Nalin T, Moura de Souza CF, Triplett EW, Dobbler PT, et al. (2019) Hepatic glycogen storage diseases are associated to microbial dysbiosis. *PLoS ONE* 14(4): e0214582. <https://doi.org/10.1371/journal.pone.0214582> PMID: 30939160