RESEARCH

Evaluating trait-based analyses for microbiome relative abundance data

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Abstract

First part title: Text for this section.

Second part title: Text for this section.

Keywords: sample; article; author

Methods

Constructing the trait database

Table 1 Sample table title. This is where the description of the table should go

Name	Description	Source
ProTraits		http://protraits.irb.hr/data.html
MediaDB		https://mediadb.systemsbiology.net/
Microbe Directory		https://github.com/microbe-directory/microbe-directory
JGI GOLD		https://gold.jgi.doe.gov/index
FAPROTAX		https://pages.uoregon.edu/slouca/LoucaLab/archive/FAPROTAX/lib/php/index.php
BacDive		https://bacdive.dsmz.de/

$$E\big[Z_1(vT_x)\big] = \frac{\mu}{r}\log x \int_0^{v\wedge 1} x^{1-u} x^{(\lambda_1/r)(v-u)}\, du. \tag{1}$$

Thus we observe that this expected value is finite for all $v > 0$ (also see [1, 2, 3, 4, 5, 6]).

Appendix

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Acknowledgements
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Funding
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Abbreviations
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Availability of data and materials
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Ethics approval and consent to participate
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Competing interests
The authors declare that they have no competing interests.

Consent for publication
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Authors' contributions
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Figures

Figure 1 Sample figure title

Figure 2 Sample figure title

Tables

Table 2 Sample table title. This is where the description of the table should go

	B1	B2	B3
A1	0.1	0.2	0.3
A2	...	..	.
A3	..	.	.

Additional Files
Additional file 1 — Sample additional file title
Additional file descriptions text (including details of how to view the file, if it is in a non-standard format or the file extension). This might refer to a multi-page table or a figure.

Additional file 2 — Sample additional file title
Additional file descriptions text.