University of New Hampshire

University of New Hampshire Scholars' Repository

Faculty Publications

8-29-2023

Reply to: Contribution of carbon inputs to soil carbon accumulation cannot be neglected

Feng Tao Tsinghua University

Benjamin Z. Houlton *Cornell University*

Serita D. Frey University of New Hampshire, Durham

Johannes Lehmann Cornell University

Stefano Manzoni Stellowhethis University on al works at: https://scholars.unh.edu/faculty_pubs

Part of the Biogeochemistry Commons

See next page for additional authors

This is a preprint posted to bioRxiv, available online: https://dx.doi.org/10.1101/2023.08.20.552557

Recommended Citation

Feng Tao, Benjamin Z. Houlton, Serita D. Frey, Johannes Lehmann, Stefano Manzoni, Yuanyuan Huang, Lifen Jiang, Umakant Mishra, Bruce A. Hungate, Michael W. I. Schmidt, Markus Reichstein, Nuno Carvalhais, Philippe Ciais, Ying-Ping Wang, Bernhard Ahrens, Gustaf Hugelius, Toby D. Hocking, Xingjie Lu, Zheng Shi, Kostiantyn Viatkin, Ronald Vargas, Yusuf Yigini, Christian Omuto, Ashish A. Malik, Guillermo Peralta, Rosa Cuevas-Corona, Luciano E. Di Paolo, Isabel Luotto, Cuijuan Liao, Yi-Shuang Liang, Vinisa S. Saynes, Xiaomeng Huang, Yiqi Luo. Reply to: Contribution of carbon inputs to soil carbon accumulation cannot be neglected, bioRxiv 2023.08.20.552557; doi: https://doi.org/10.1101/2023.08.20.552557

This Article is brought to you for free and open access by University of New Hampshire Scholars' Repository. It has been accepted for inclusion in Faculty Publications by an authorized administrator of University of New Hampshire Scholars' Repository. For more information, please contact Scholarly.Communication@unh.edu.

Authors

Feng Tao, Benjamin Z. Houlton, Serita D. Frey, Johannes Lehmann, Stefano Manzoni, Yuanyuan Huang, Lifen Jiang, Umakant Mishra, Bruce A. Hungate, Michael W. I. Schmidt, Markus Reichstein, Nuno Carvalhais, Philippe Ciais, Ying-Ping Wang, Bernhard Ahrens, Gustaf Hugelius, Toby D. Hocking, Xingjie Lu, Zheng Shi, Kostiantyn Viatkin, Ronald Vargas, Yusuf Yigini, Christian Omuto, Ashish A. Malik, Guillermo Peralta, Rosa Cuevas-Corona, Luciano E. Di Paolo, Isabel Luotto, Cuijuan Liao, Yi-Shuang Liang, Vinisa S. Saynes, Xiaomeng Huang, and Yiqi Luo

Reply to: Contribution of carbon inputs to soil carbon accumulation cannot be neglected

Feng Tao^{1, 2, 3}, Benjamin Z. Houlton⁴, Serita D. Frey⁵, Johannes Lehmann⁶, Stefano Manzoni⁷, Yuanyuan Huang⁸, Lifen Jiang⁹, Umakant Mishra^{10, 11}, Bruce A. Hungate^{12, 13}, Michael W. I. Schmidt¹⁴, Markus Reichstein², Nuno Carvalhais^{2, 15}, Philippe Ciais¹⁶, Ying-Ping Wang¹⁷, Bernhard Ahrens², Gustaf Hugelius⁷, Toby D. Hocking¹³, Xingjie Lu¹⁸, Zheng Shi¹⁹, Kostiantyn Viatkin^{6, 3}, Ronald Vargas³, Yusuf Yigini³, Christian Omuto³, Ashish A. Malik²⁰, Guillermo Peralta³, Rosa Cuevas-Corona³, Luciano E. Di Paolo³, Isabel Luotto³, Cuijuan Liao¹, Yi-Shuang Liang¹, Vinisa S. Saynes³, Xiaomeng Huang^{1, *}, and Yiqi Luo^{9, *}

¹Department of Earth System Science, Ministry of Education Key Laboratory for Earth System Modelling, Institute for Global Change Studies, Tsinghua University, Beijing, China ²Max Planck Institute for Biogeochemistry, Jena, Germany

³Food and Agricultural Organization of the United Nations, Rome, Italy

⁴Department of Ecology and Evolutionary Biology and Department of Global Development, Cornell University, Ithaca, NY, USA

⁵Center for Soil Biogeochemistry and Microbial Ecology, Department of Natural Resources and the Environment, University of New Hampshire, Durham, NH, USA ⁶Soil and Crop Sciences Section, School of Integrative Plant Science, Cornell University, Ithaca NY, USA

⁷Department of Physical Geography and Bolin Centre for Climate Research, Stockholm University, Stockholm, Sweden

⁸Key Laboratory of Ecosystem Network Observation and Modeling, Institute of Geographic Sciences and Natural Resources Research, Chinese Academy of Sciences, Beijing, China.

⁹School of Integrative Plant Science, Cornell University, Ithaca NY, USA

¹⁰Computational Biology & Biophysics, Sandia National Laboratories, Livermore, CA, USA

¹¹Joint BioEnergy Institute, Lawrence Berkeley National Laboratory, Emeryville, CA,

USA

¹²Center for Ecosystem Science and Society, Department of Biological Sciences, Northern

Arizona University, Flagstaff, AZ, USA.

¹³School of Informatics, Computing and Cyber Systems, Northern Arizona University,

Flagstaff, AZ, USA

¹⁴Department of Geography, University of Zurich, Zurich, Switzerland

¹⁵Departamento de Ciências e Engenharia do Ambiente, DCEA, Faculdade de Ciências e

Tecnologia, FCT, Universidade Nova de Lisboa, Caparica, Portugal

¹⁶Laboratoire des Sciences du Climat et de l'Environnement, LSCE/IPSL, CEA-CNRS-

UVSQ, Université Paris-Saclay, Gif-sur-Yvette, France

¹⁷CSIRO Environment, Aspendale, Victoria, Australia

¹⁸School of Atmospheric Sciences, Sun Yat-sen University, Guangzhou, China

¹⁹Institute for Environmental Genomics and Department of Microbiology and Plant Biology,

University of Oklahoma, Norman, OK, USA

²⁰School of Biological Sciences, University of Aberdeen, Aberdeen, UK

*Corresponding Authors: Yiqi Luo: yiqi.luo@cornell.edu; Xiaomeng Huang: hxm@tsinghua.edu.cn

Abstract

In the accompanying Comment¹, He et al. argue that the determinant role of microbial carbon use efficiency in global soil organic carbon (SOC) storage shown in Tao et al. (2023)² was overestimated because carbon inputs were neglected in our data analysis while they suggest that our model-based analysis could be biased and model-dependent. Their argument is based on a different choice of independent variables in the data analysis and a sensitivity analysis of two process-based models other than that used in our study. We agree that both carbon inputs and outputs (as mediated by microbial processes) matter when predicting SOC storage – the question is their relative contributions. While we encourage further studies to examine how the evaluation of the relative importance of CUE to global SOC storage may vary with different model structures, He et al.'s claims about Tao et al. (2023) need to be taken as an alternative, unproven hypothesis until empirical data support their specific parameterization. Here we show that an additional literature assessment of global data does not support He et al.'s argument, in contrast to our study, and that further study on this topic is essential.

Main

The higher explanatory power of carbon input than microbial CUE to SOC storage envisaged by He et al. does not hold at the global scale when more data are considered (Table 1). He et al. proposed that carbon input is potentially more important than microbial CUE by using the net primary productivity (NPP) as carbon input to explain the spatial variation of SOC in 132 datasets in a mixed-effects model. However, the statistical models in Tao et al. (2023) were not applied to evaluate the relative importance of either CUE or NPP to SOC, but to determine whether microbial CUE is positively or negatively correlated with SOC. This issue raised by He et al. might become relevant if NPP obscures the CUE-SOC relationship to the extent of changing its direction. In Supplementary Table 3, Tao et al. (2023) showed that the

inclusion of NPP in a mixed-effects model does not influence the positive CUE-SOC correlation. Moreover, NPP may have high explanatory power for SOC across these 132 local sites, but not necessarily at the global scale. We extracted NPP information from a MODIS-based product³ and the PRODA-retrieved CUE from 57,267 soil profiles (Extended Data Fig. 1b of Tao et al., 2023) and found that CUE explains more spatial variation of SOC (37%) than NPP (9%) at the global scale (Table 1). Indeed, the notion that NPP is a minor factor in explaining SOC dynamics and spatial variation at both regional and global scales has been well documented in the literature⁴⁻⁶.

Process-based models can theoretically generate an array of patterns and predictions, as correctly argued by He et al. and as well documented in the literature. However, models yield realistic predictions only after they are constrained by observations. He et al. used two models to examine varying sensitivities of SOC storage in response to changes in a parameter, β , that represents the density dependence of microbial mortality⁷, arguing that SOC storage could be more sensitive to changes in NPP than in microbial CUE under certain parameterizations. Similarly, the microbial model used in Tao et al. (2023) generates a nested set of sensitivities for SOC storage in response to NPP, but assuming that $\beta = 1 - i.e.$, assuming that mortality is not density-dependent (Fig. 1). SOC storage shows no response to doubled NPP when the turnover time (τ) of both enzyme (ENZ) and microbial biomass (MIC) is very short (e.g., $\tau_{ENZ,decay} = 0.05$ years and $\tau_{MIC} = 0.2$ years, Fig 1a). Alternatively, SOC storage dramatically increases with doubled NPP when the turnover time of either of these two pools is higher (e.g., $\tau_{ENZ,decay}$ increases from 0.05 years to 0.5 years, Fig. 1a-c, or τ_{MIC} increases from 0.2 years to 1.5 years, Fig. 1d-f). Nevertheless, microbial CUE emerged from our SOC data assimilation into the model to be more important than NPP to global SOC storage. It is the Bayesian framework used in our study via data assimilation

that identified the most probable mechanisms among these diverse mechanisms. We look forward to an analysis where first data assimilation is conducted to estimate the β value in order to then demonstrate that the estimated parameter could disprove the fundamental importance of microbial CUE to SOC storage.

Estimates of NPP by different process-based models and data products indeed remain uncertain, as pointed out by He et al. While the uncertainty in NPP might influence CUE-SOC relationships, our analysis showed that variation in NPP values from -10% to +10% of the CLM5 simulated values had much less effect on SOC than microbial CUE (Fig. 4b of Tao et al., 2023). Nevertheless, we greatly appreciate the suggestion to include carbon input as a parameter for optimization. We thus encourage the scientific community to conduct data assimilation studies to constrain all parameters, including mortality- and NPP-related ones, that may influence the CUE-SOC relationships.

Methods

All the data, statistical methods, and the microbial model have been described in Tao et al. (2023) and can be publicly accessed via <u>https://www.nature.com/articles/s41586-023-06042-</u><u>3</u>.

Competing interests:

The authors declare no competing interests.

Author contributions:

F. T. and Y. L. drafted the reply. All authors contributed to the text and approved the final version.

References:

- 1 He, X. *et al.* Contribution of carbon inputs to soil carbon accumulation cannot be neglected. *bioRxiv*, 2023.2007. 2017.549330 (2023).
- Tao, F. *et al.* Microbial carbon use efficiency promotes global soil carbon storage.
 Nature, 1-5 (2023).
- 3 Zhao, M. & Running, S. W. Drought-induced reduction in global terrestrial net primary production from 2000 through 2009. *science* **329**, 940-943 (2010).
- 4 Mishra, U., Drewniak, B., Jastrow, J. D. & Matamala, R. M. Spatial representation of organic carbon and active-layer thickness of high latitude soils in CMIP5 earth system models. *Geoderma* **300**, 55-63 (2017).
- Minasny, B. *et al.* Current NPP cannot predict future soil organic carbon sequestration potential. Comment on "Photosynthetic limits on carbon sequestration in croplands".
 (2022).
- 6 Crowther, T. W. *et al.* The global soil community and its influence on biogeochemistry. *Science* **365**, eaav0550 (2019).
- 7 Georgiou, K., Abramoff, R. Z., Harte, J., Riley, W. J. & Torn, M. S. Microbial community-level regulation explains soil carbon responses to long-term litter manipulations. *Nature Communications* 8, 1-10 (2017).

Tables and Figures

Table 1 | Microbial CUE explains more spatial variation of SOC storage than NPP at the

global scale. Statistics shown in the table are unstandardized coefficients of relationships between CUE (from data assimilation) or NPP (from remote sensing data) and SOC content in a mixed-effects model. CUE or NPP was set as the fixed effect to predict SOC content. Climate types that soil profiles belong to were set as the random effect. We assumed random intercepts in all regressions. The total observation size $n_{obs} = 56,270$, the random effect size $n_{climate} = 12$.

		Intercept	CUE or NPP
log10(SOC)~CUE + (1 Climate Type)			
variance explained by mixed model: 37%			
Fixed Effects	Estimates	0.89	1.18
	Std. Error	0.055	0.0096
	t value	16.07	122.44
	Р	< 0.0001	< 0.0001
Random Effects	Standard Deviation	0.19	NA
log10(SOC)~NPP + (1 Climate Type)			
variance explained by mixed model: 9%			
Fixed Effects	Estimates	1.13	2.13×10 ⁻⁴
	Std. Error	0.065	4.61×10 ⁻⁶
	t value	17.54	46.32
	Р	< 0.0001	< 0.0001
Random Effects	Standard Deviation	0.05	NA



Fig. 1 | **Varying sensitivity of SOC storage to doubled NPP under different combinations of parameter values with the microbial model used in Tao et al. (2023).** We chose different values for the turnover time (in years) for enzyme and microbial biomass carbon pools of the microbial model used in Tao et al. (2023) at one site and doubled the magnitude of NPP in model simulation for each set of parameterizations. Different panels in this figure present how the relation of CUE and SOC storage changed with increased NPP. The SOC values were standardized using the Z-score method to be comparable with the results shown in He et al.