

Rhizodéposition et séquestration/stockage de C dans les sols

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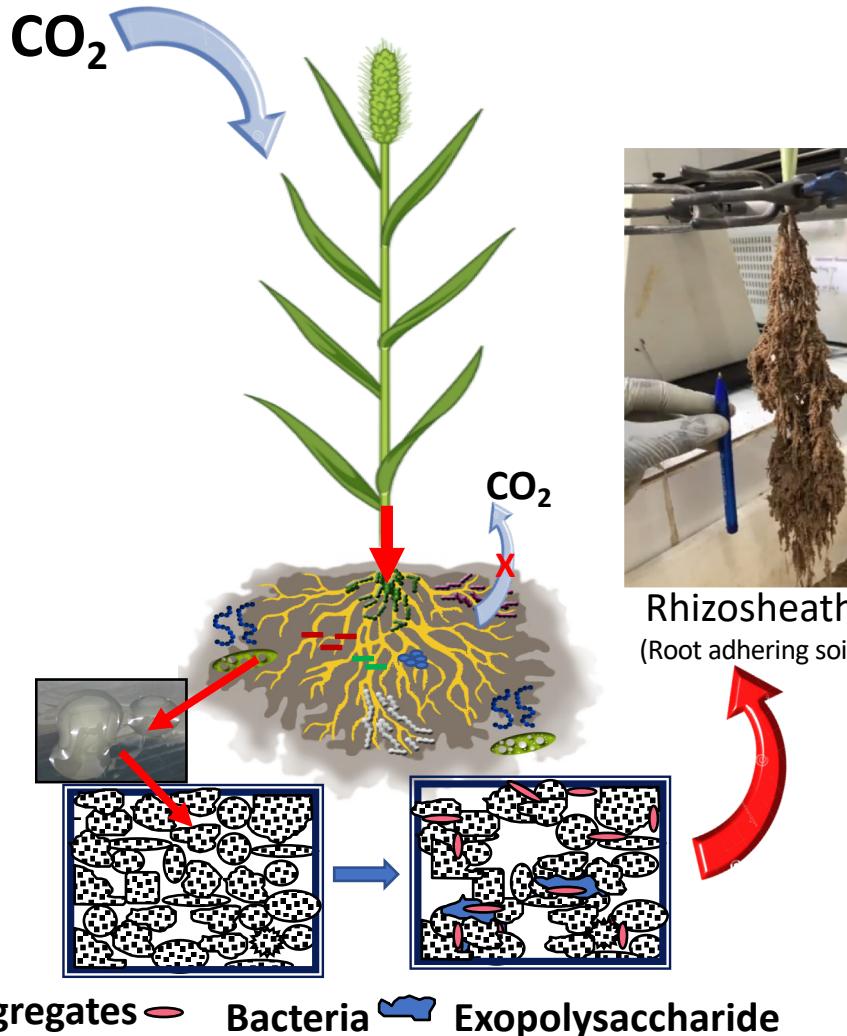
Christine HATTÉ, LSCE

Thierry HEULIN, LEMiRE-BIAM

The limits of "classic" approaches to restoring organic matter to soils :

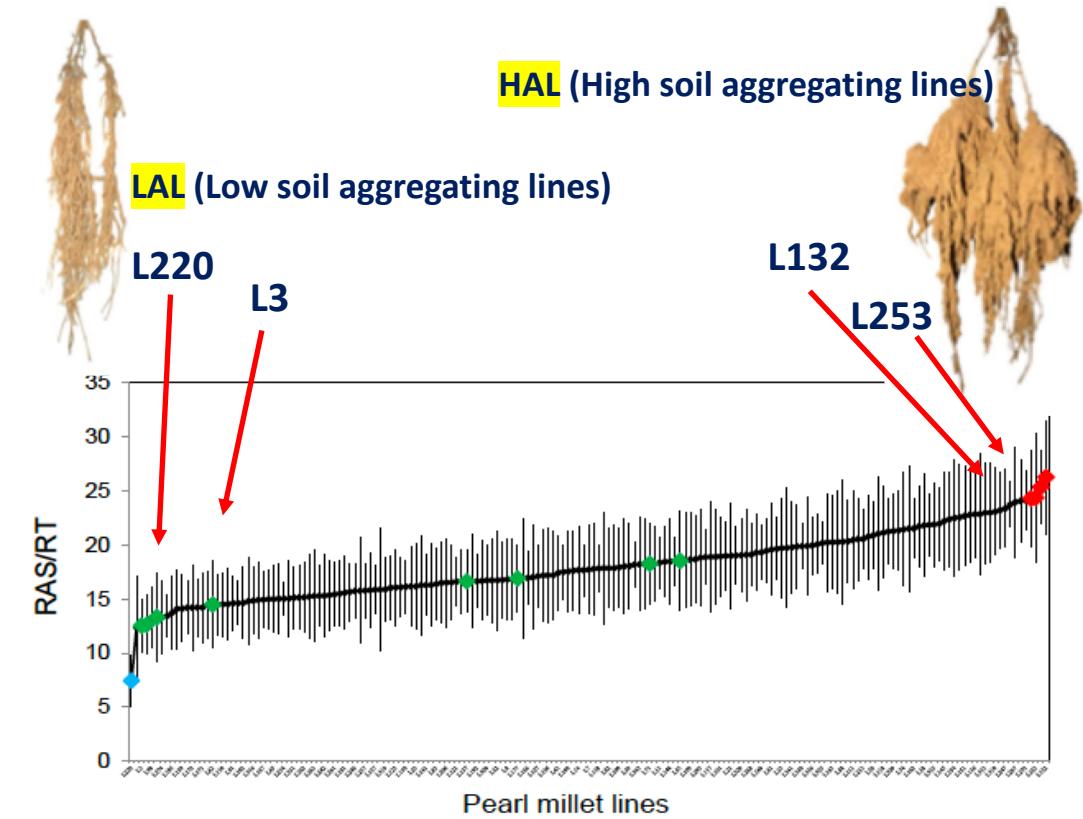
- Agroforestry could be a solution, albeit with certain constraints
- Spreading crop residues (straw), effluents or composts only enriches the soil surface and can sometimes lead to mineralization of existing C.

Can varietal selection be an alternative to conventional practices for returning C to the soil?

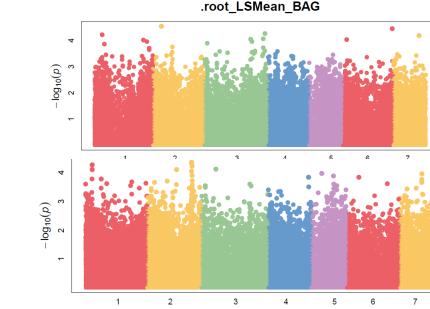
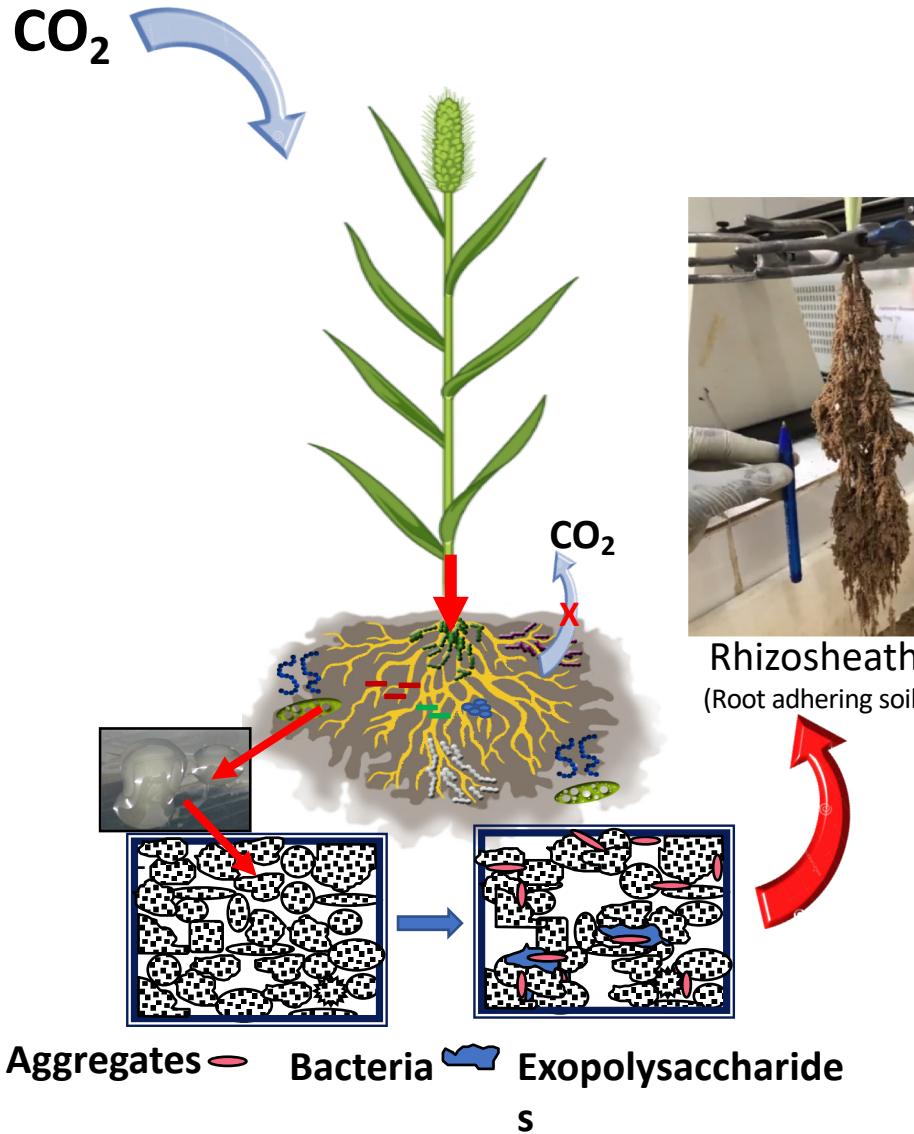


AGENCE NATIONALE DE LA RECHERCHE
ANR
RootAdapt

>180 pearl millet inbred lines have been screened:
Root Architecture/Soil structuration

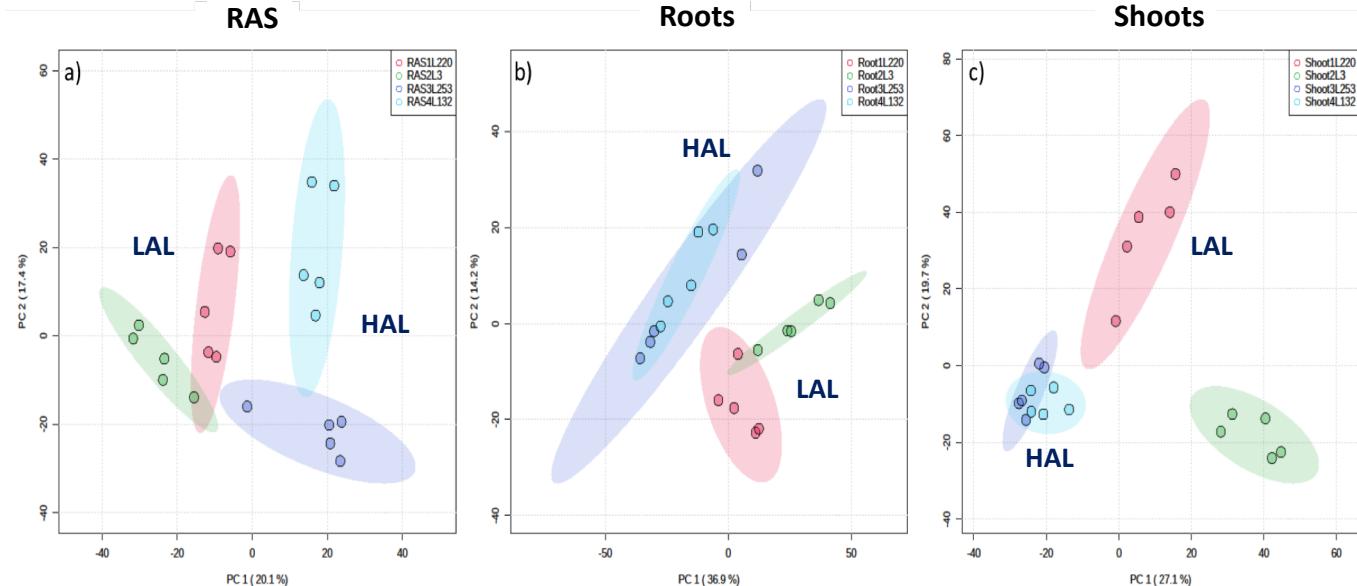


Objectives: Is there any correlation between rhizosheath size, rhizodeposition, root exudates composition, microbial community assemblage and carbon storage in soil ?

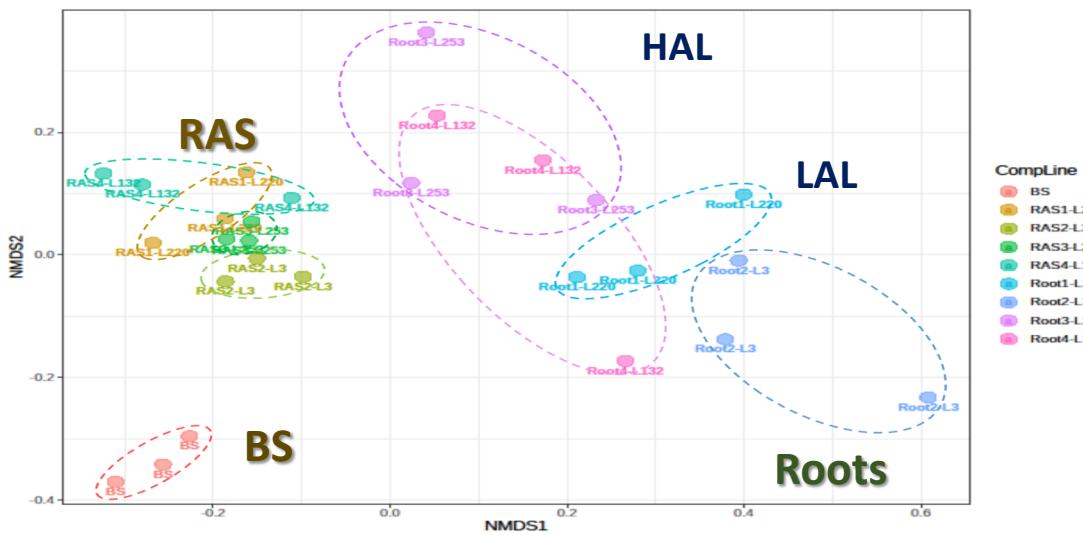


The root exudation of millet evaluated by a simple phenotype to measure (**root-adhering soil, SAR**) is genetically controlled, thus paving the way for varietal selection. (De la Fuente et al., Sc. Rep. 2022)

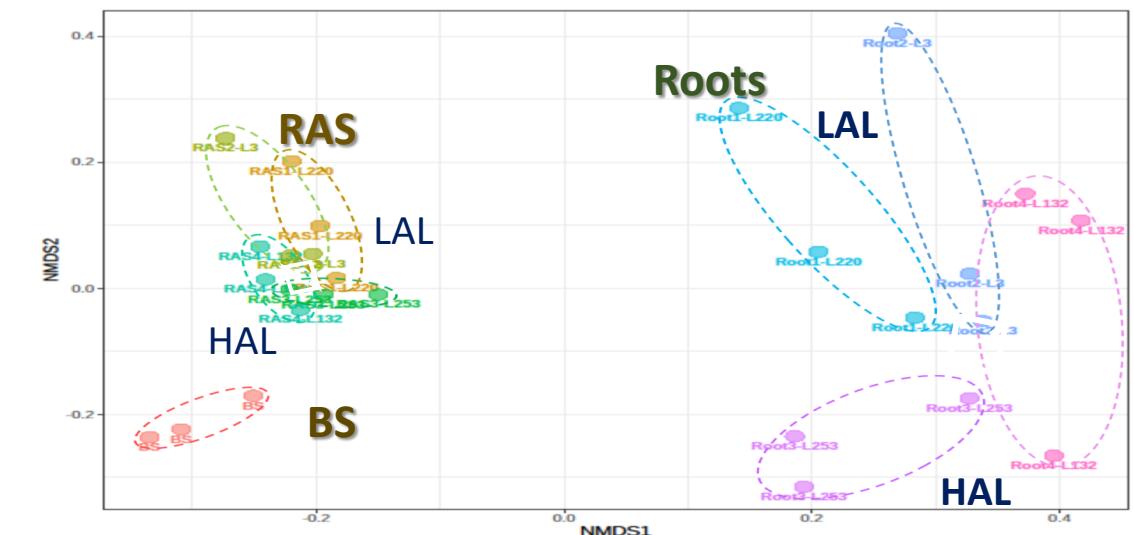
Alahmad *et al.*, Microbiome, 2024



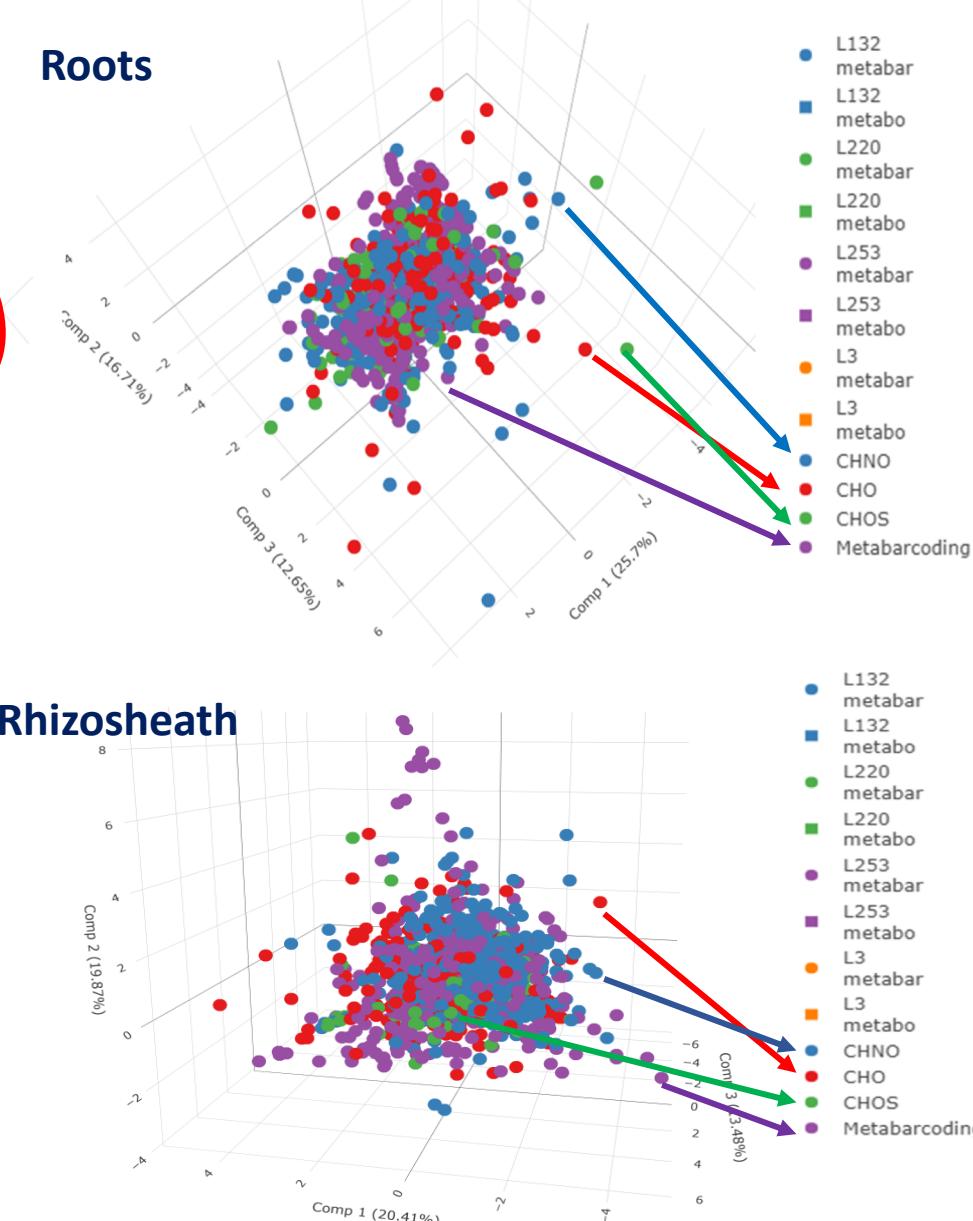
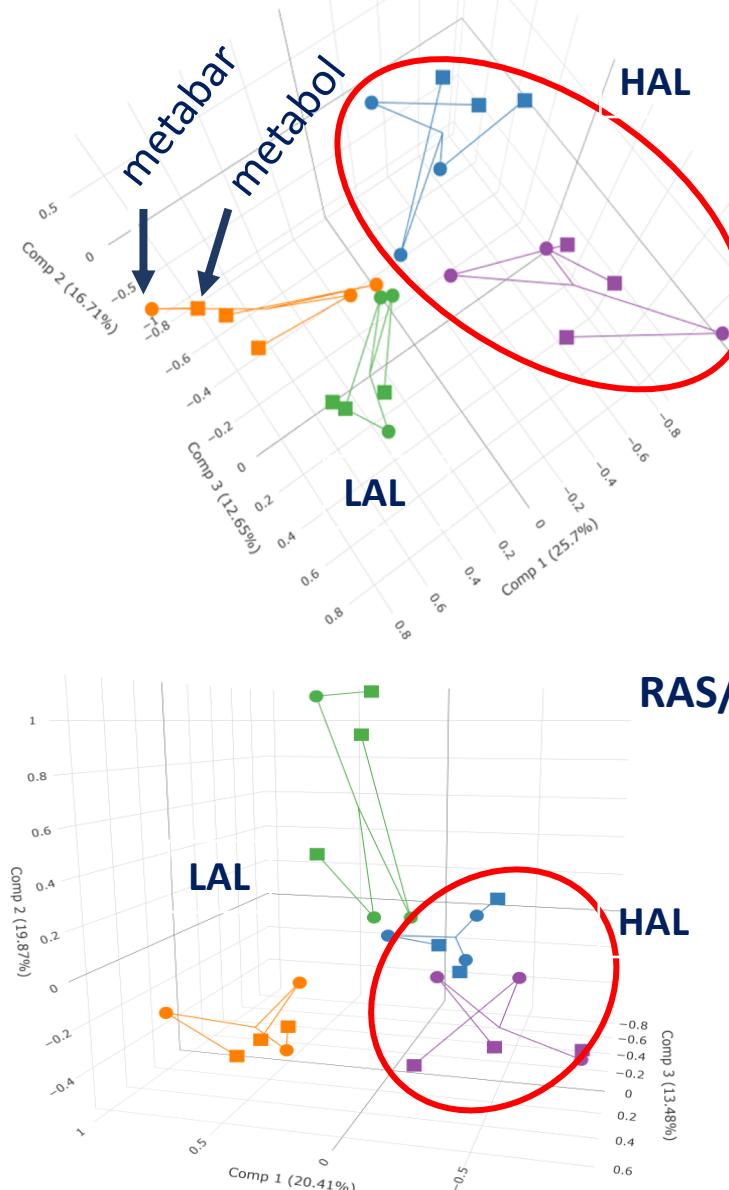
PCA 2D scores plots illustrating the distribution of assigned compounds in different compartments of each PM line.



NMDS plots of the beta-diversity of bacterial communities



NMDS plots of the beta-diversity of fungal communities



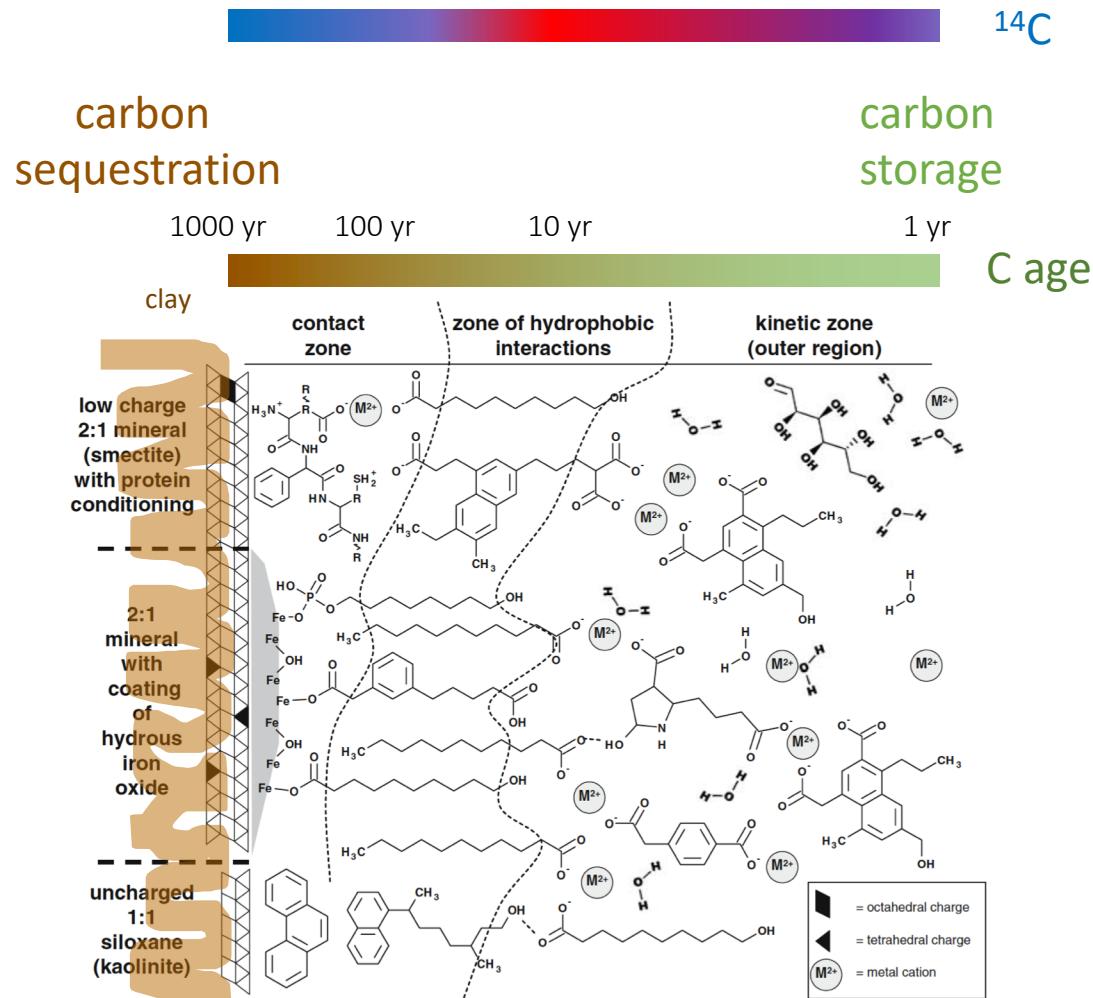
3D plot of Co-inertia analysis (CIA) of metabarcoding (circular) and metabolomics (square) data sets in the root & RAS of the four PM lines where the lines relates the position of the samples in the metabarcoding dataset in relation to the metabolomics one.

- The rhizosheath formation is under complex genetic control in pearl millet and suggests that it is mainly regulated by root exudation/rhizodeposition (de La Fuente *et al.*, 2022).
- The root-adhering soil mass is a heritable trait in pearl millet that correlates with changes in rhizosphere microbiota structure and functions, and root exudates composition (Alahmad *et al.*, 2024).
- The priming effect amplitude (C_{lost}/C_{new} ratio) was higher for the LAL than for the HAL, indicating a better C sequestration potential of the latter (Ndour *et al.*, 2022).

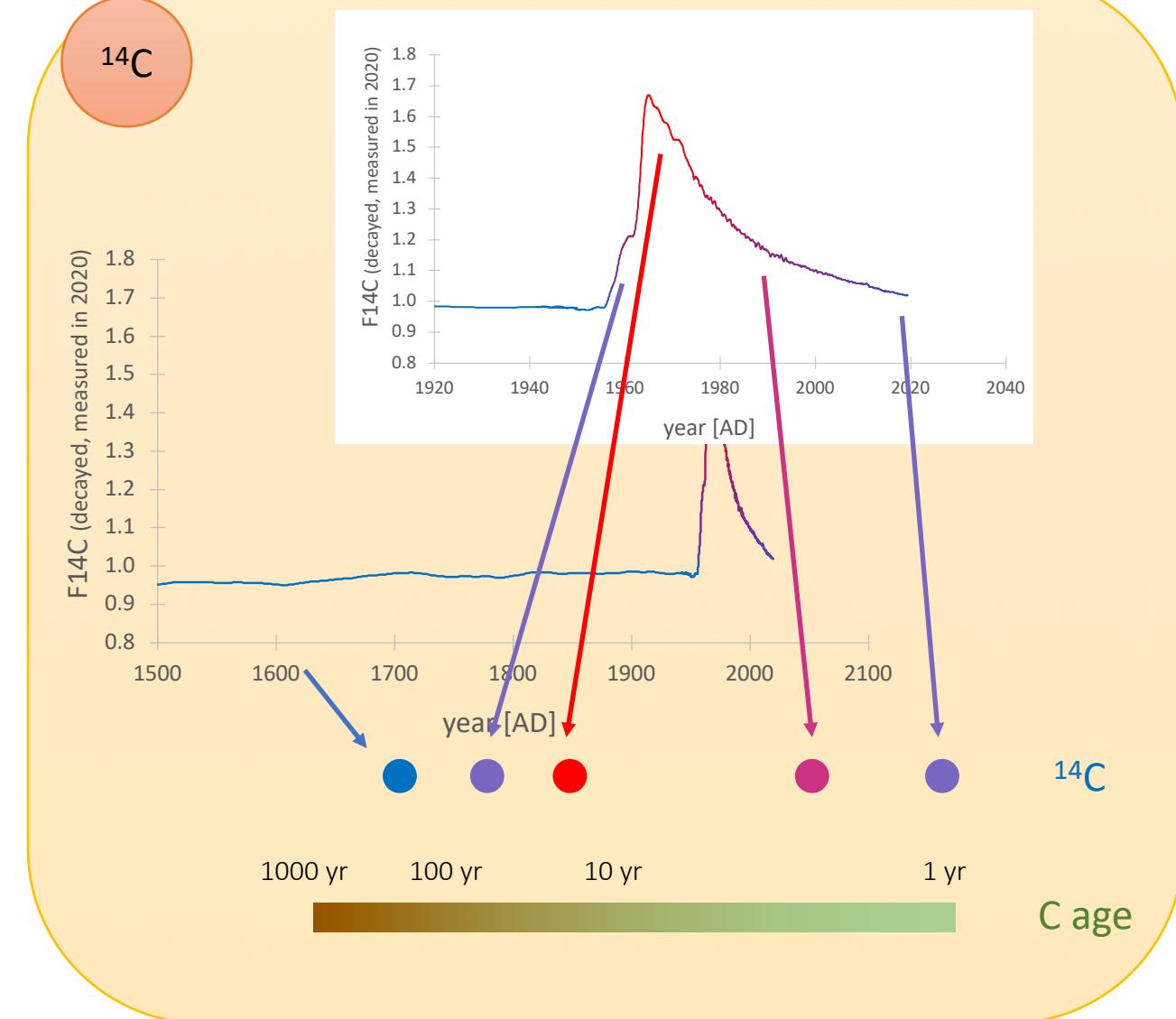
Evaluation of agronomical practices

contribution of isotopic measurements

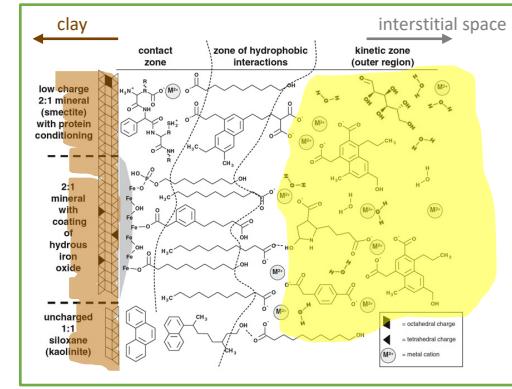
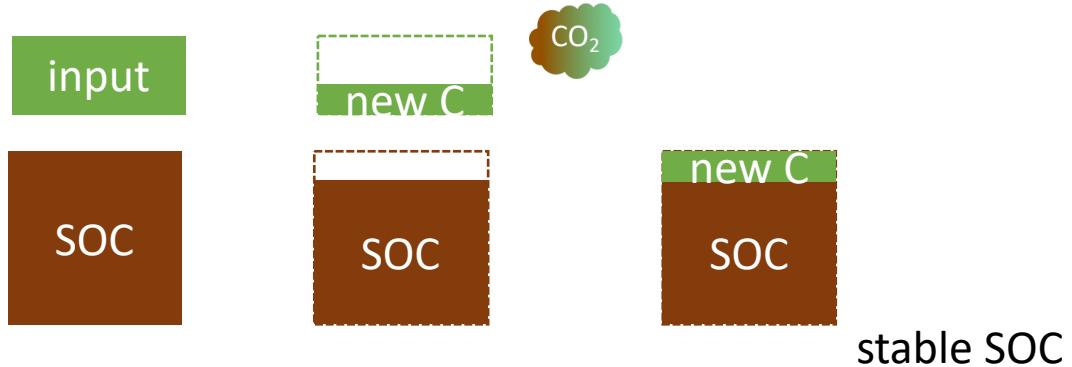
carbon storage – carbon sequestration



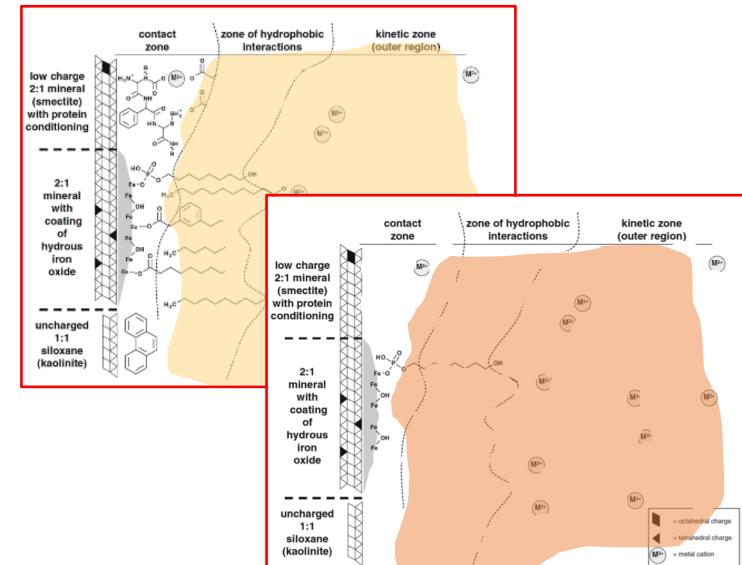
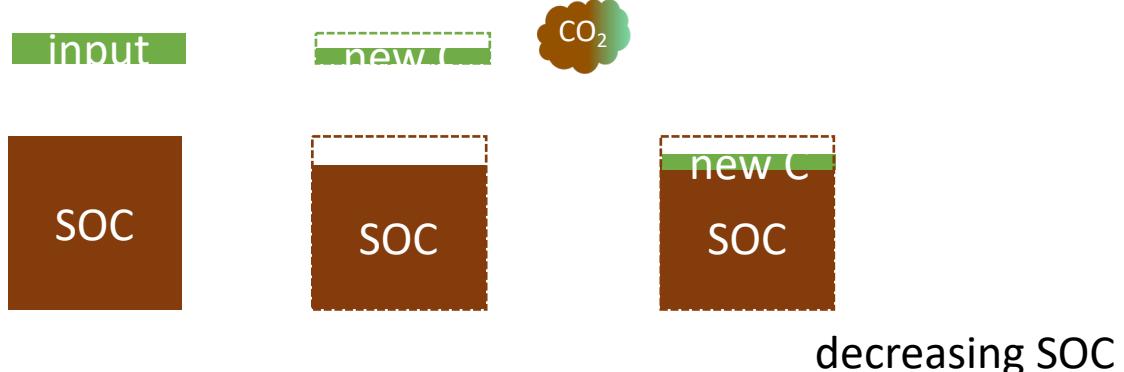
Kleber et al. 2007



heterotrophic respiration - priming effect

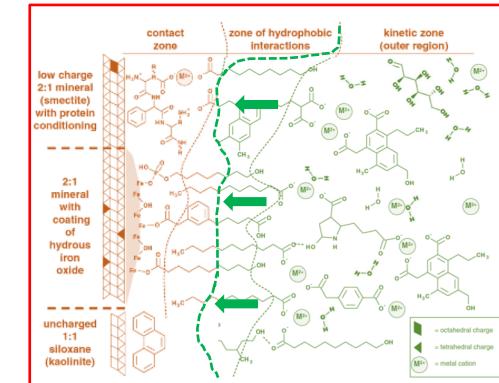
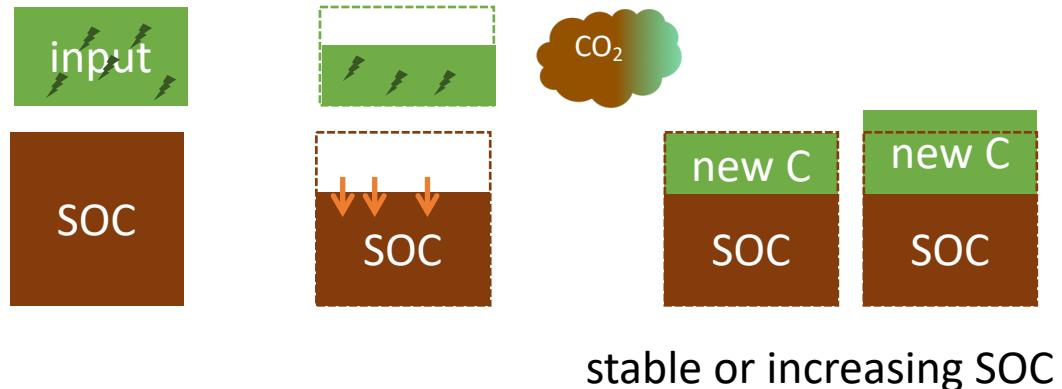


- steady-state**
- exchanges in the kinetic zone



- deleterious practices**
- hydrophobic/contact zones affected

heterotrophic respiration - priming effect



good (?) idea

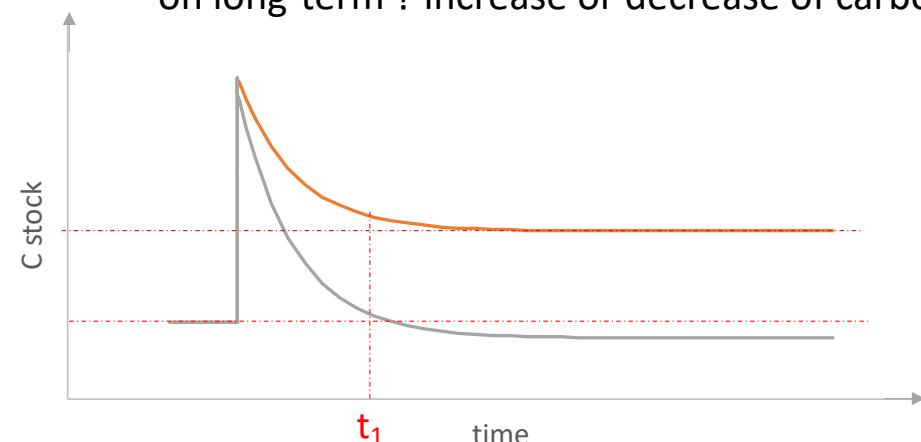
enhancement of priming effect
change in molecular cortege

- shrinking of hydrophobic zone
- enlargement of the kinetic zone

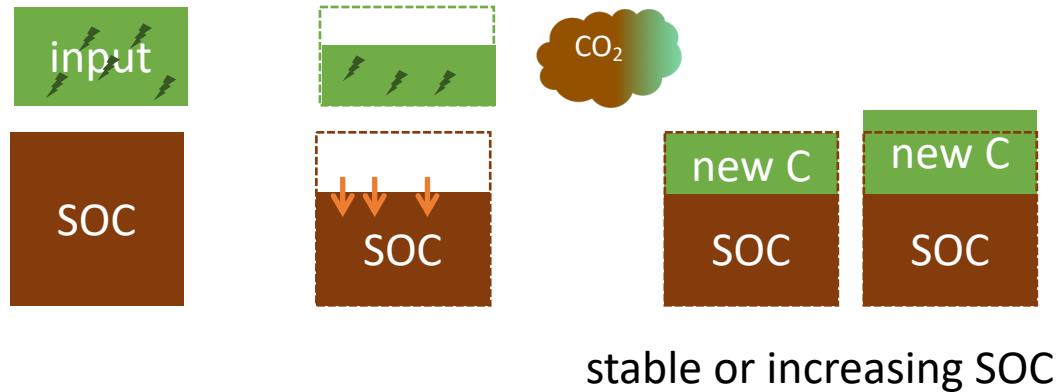
=> less sequestered carbon

=> more molecule with short turnover time

on long-term ? increase or decrease of carbon stock?



priming effect – how to assess it ?

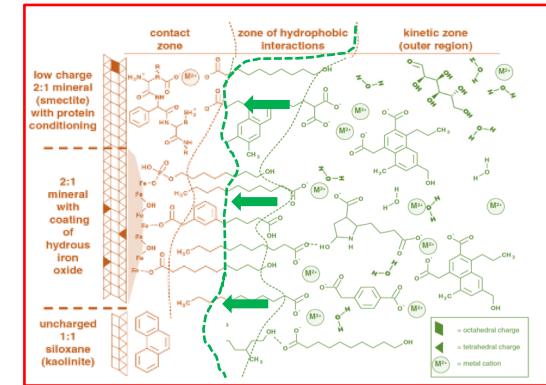


good (?) idea

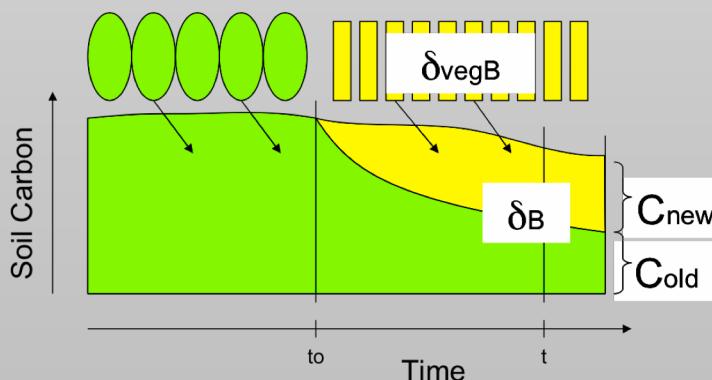
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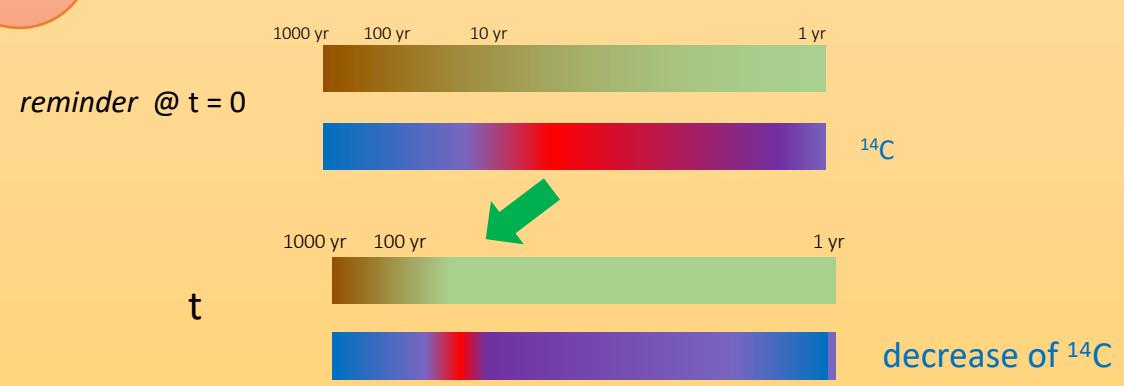
=> less sequestrated carbon
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stable isotopes - natural labelling (¹³C & ¹⁵N)
(Balesdent's approach)

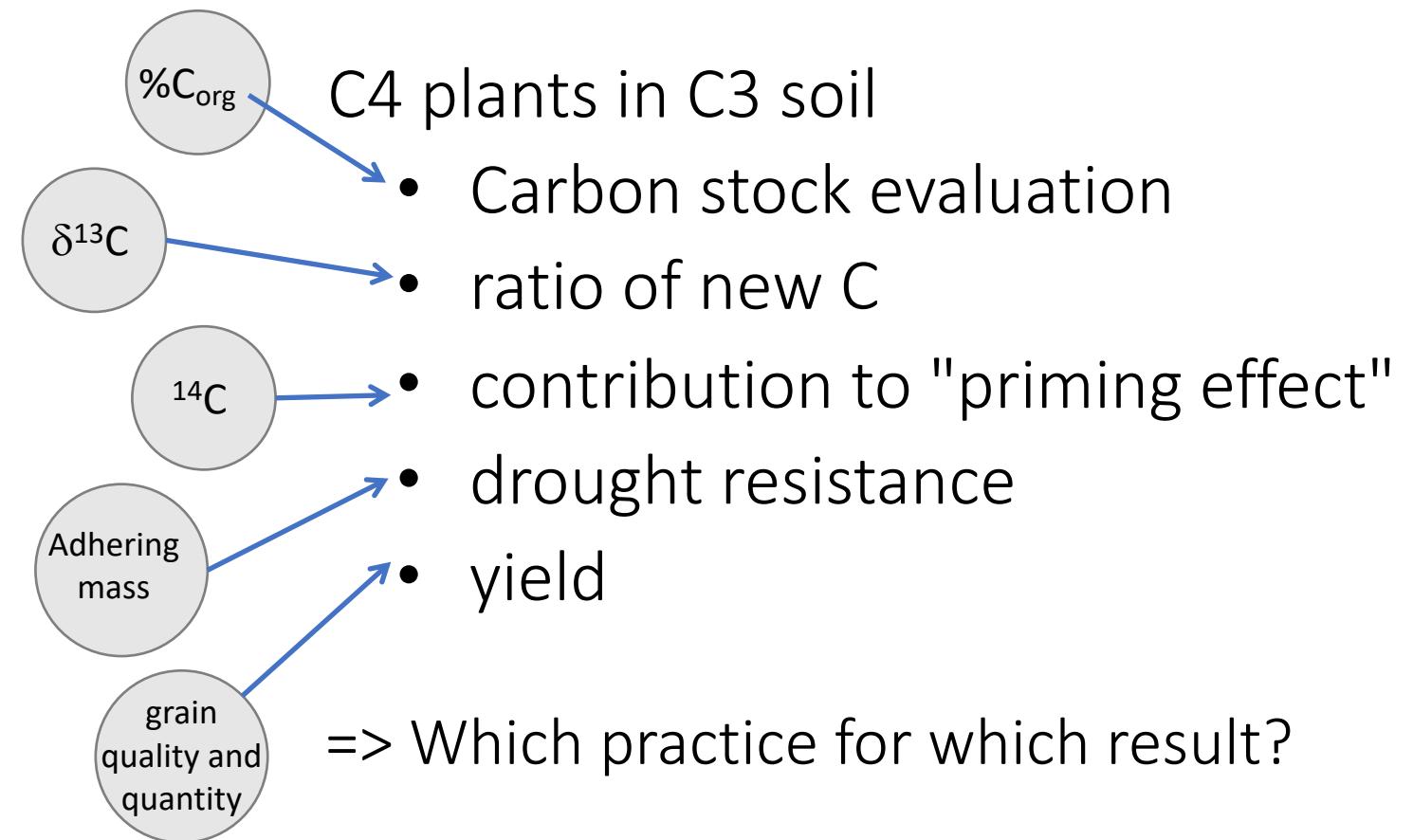
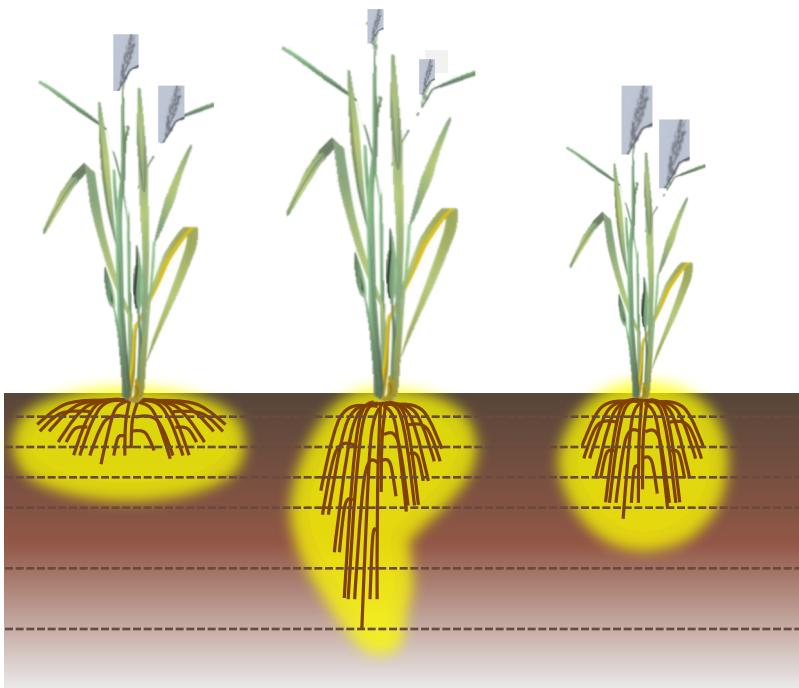


¹⁴C informs on nature of the lost carbon



Assessment – a case study

An evaluation of new agronomical practices



Assessment – a case study

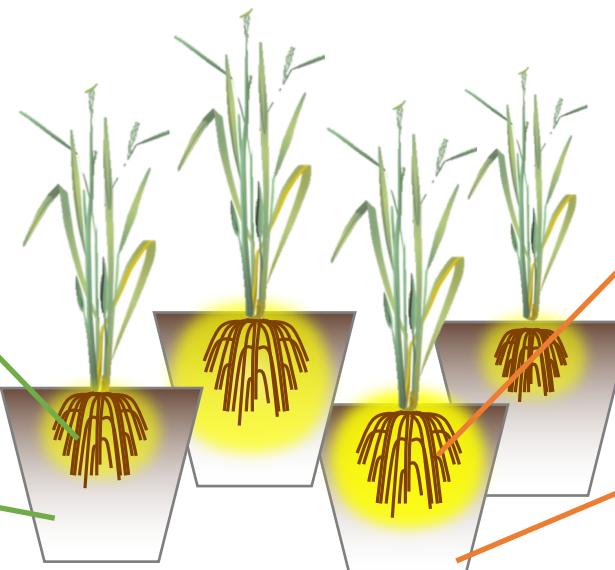
An evaluation of new agronomical practices



L3 (intermédiaire)

+ $C_{new} = 0.019 \text{ \%wt}$
-- $C_{lost} = 0.066 \text{ \%wt}$
 $\int F^{14}C_{lost} = 1.21$

$$C_{new} = -0.018 \text{ \%wt}$$



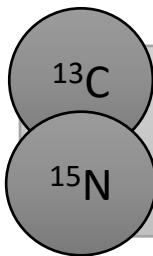
L132 (hyper)

- $C_{new} = 0.013 \text{ \%wt}$
+ $C_{lost} = 0.038 \text{ \%wt}$
 $\int F^{14}C_{lost} = 1.26$

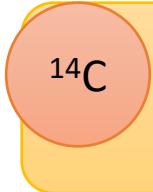
$$C_{new} = -0.010 \text{ \%wt}$$

=> L132 is a good "4%" candidate : it contributes to C input while keeping reduced loss of C
=> ATTENTION to *priming effect* ! : L3 destores more C than it brings in

priming effect @ LSCE



EA/GC – IRMS (Delta+ XP)
bulk and specific compound



ECHoMICADAS
(few µg of C)

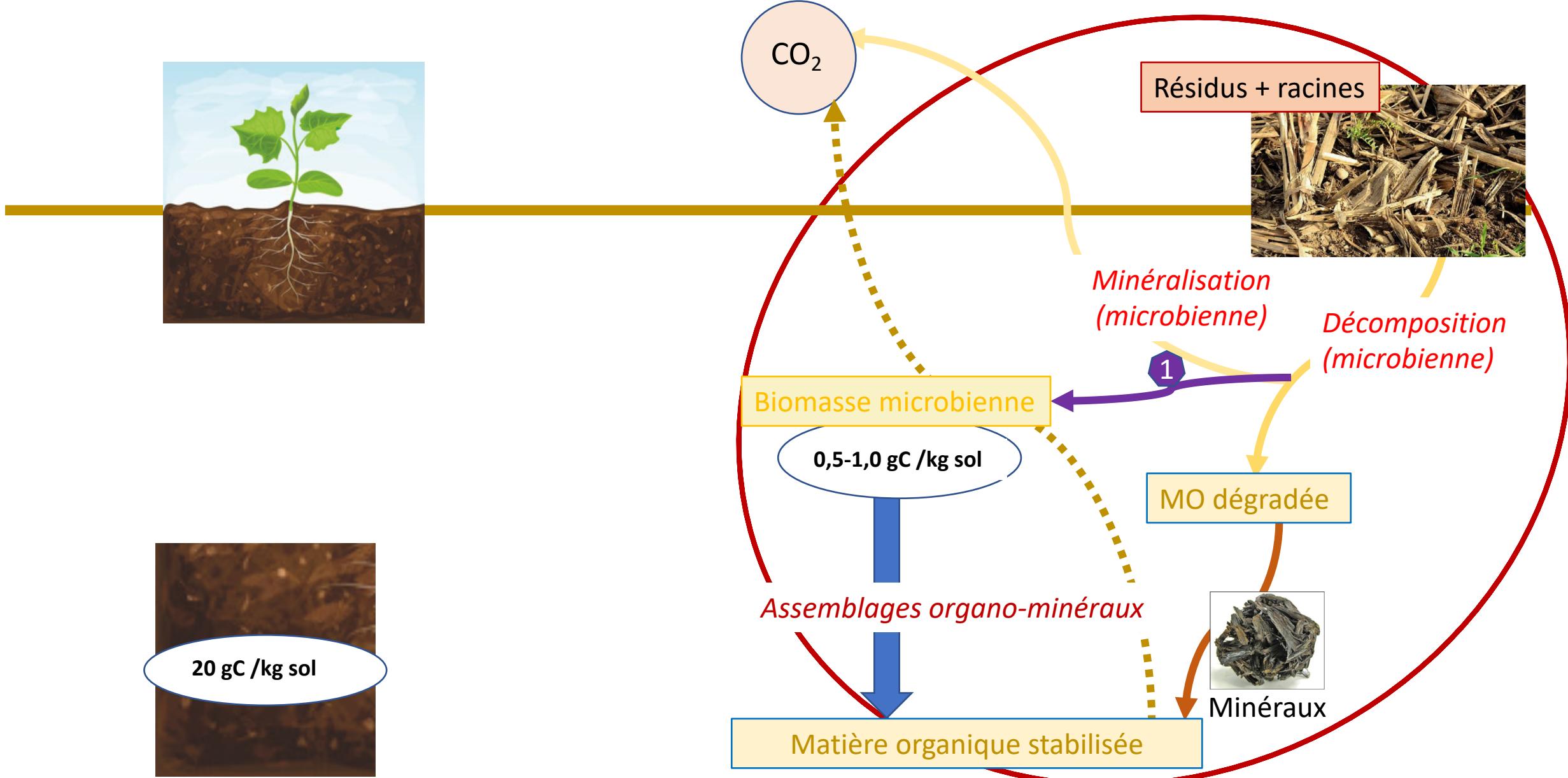
Balesdent et al. 2017, 2018
Camino-Serrano et al. 2019
Jreich et al. 2018, 2021
Kheirbeik et al. 2016
Lawrence et al. 2020
Mathieu et al. 2015

Mendez-Millan et al. 2013
Ndour et al. 2022
Paul et al. 2016, 2019
Quéro et al. 2022
Rubino et al . 2014
Tifafi et al. 2018a, 2018b

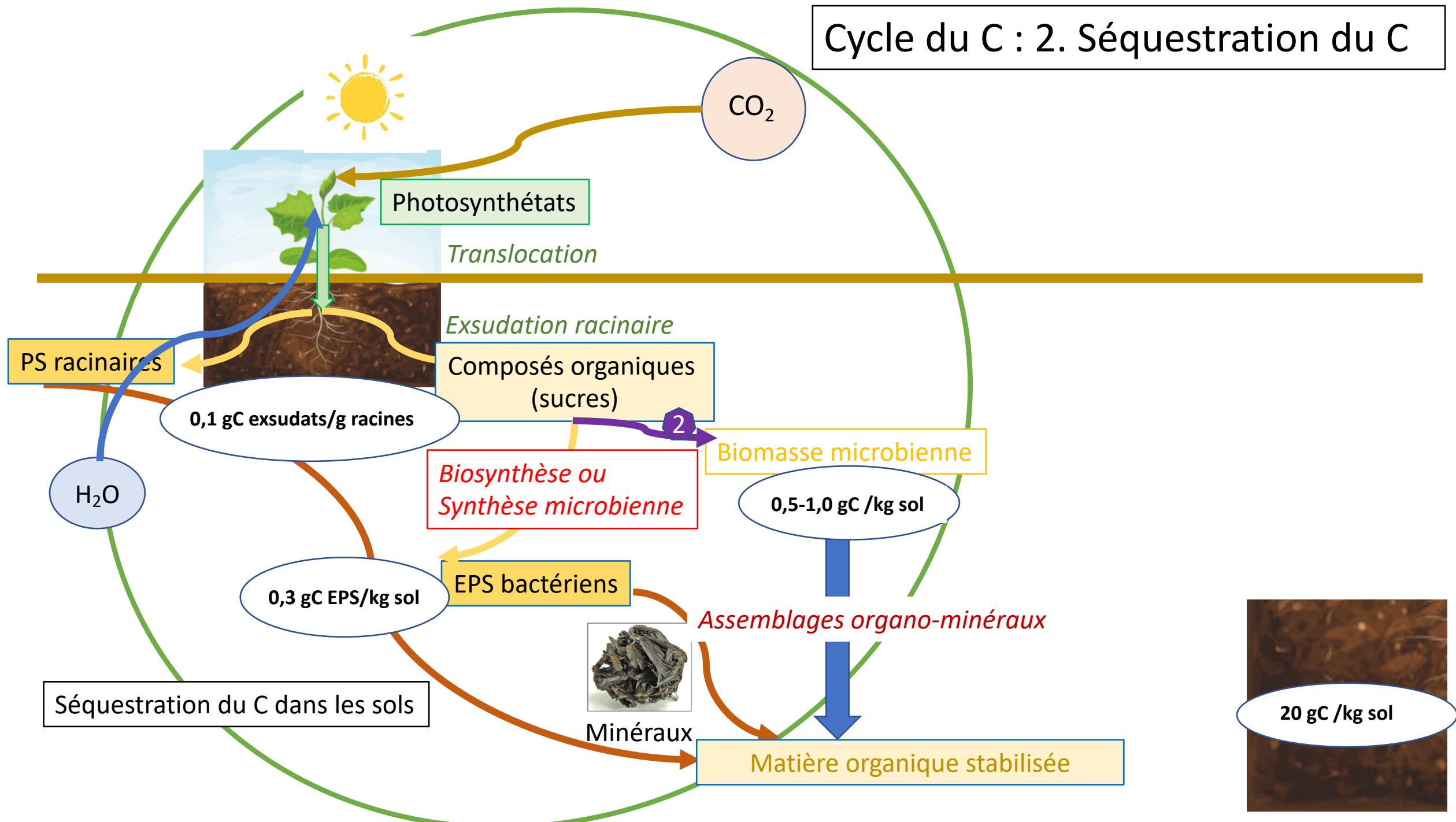


Conclusion-Perspectives

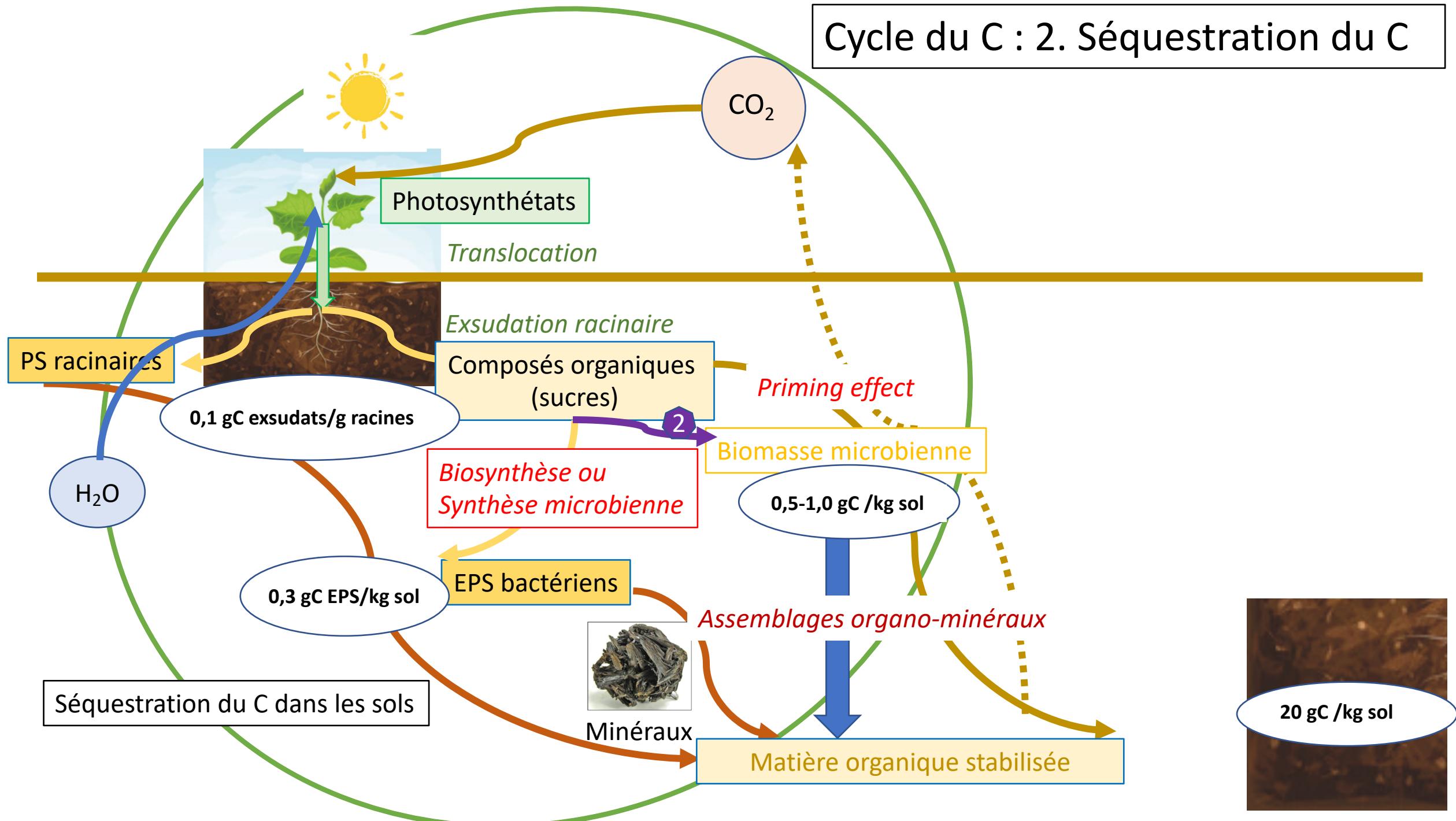
Cycle du C : 1. Stockage du C



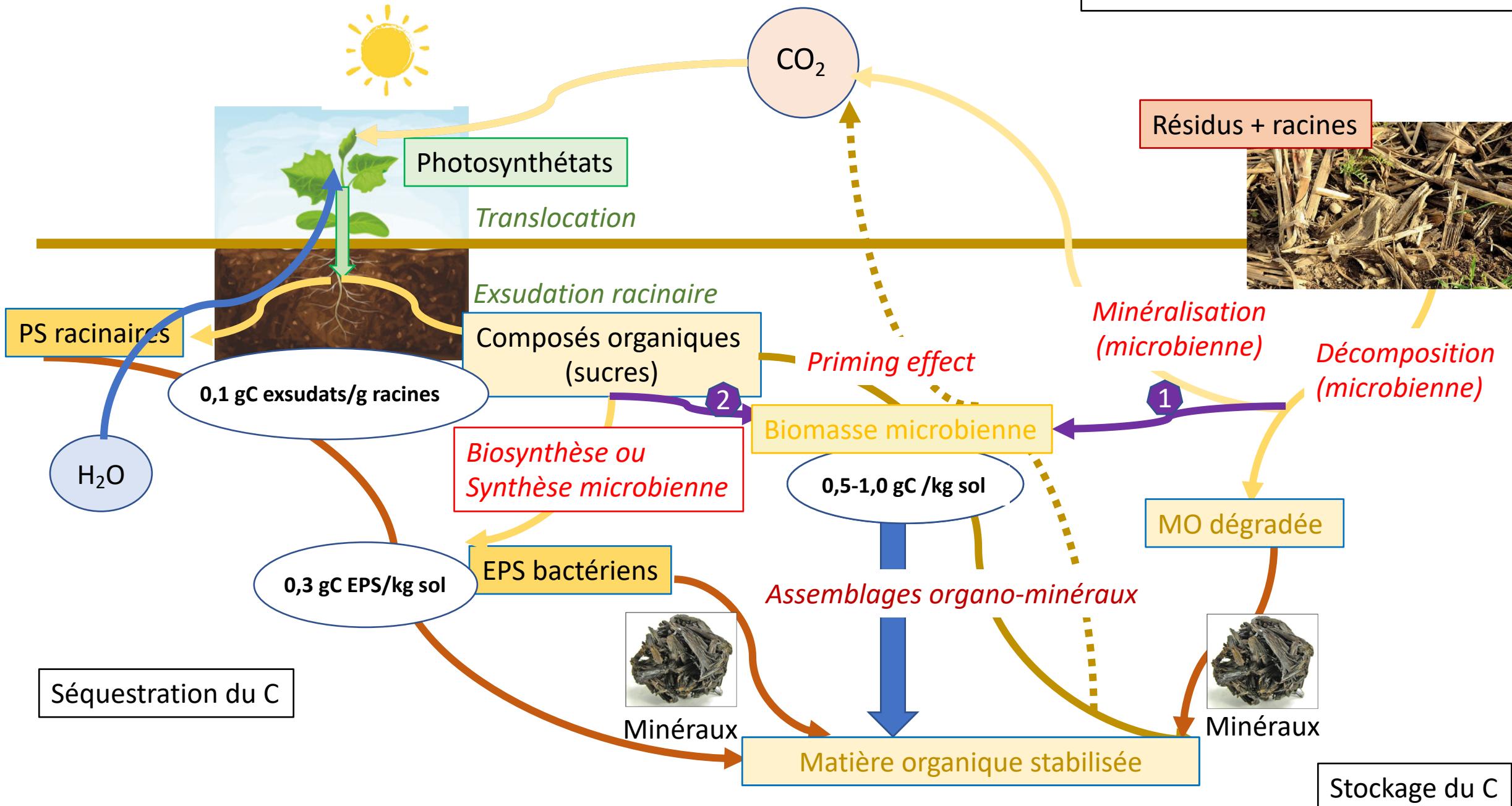
Cycle du C : 2. Séquestration du C



Cycle du C : 2. Séquestration du C



Cycle du C dans les sols



Article

Microbial carbon use efficiency promotes global soil carbon storage

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Feng Tao^{1,2,3}, Yuanyuan Huang⁴, Bruce A. Hungate^{5,6}, Stefano Manzoni⁷, Serita D. Frey⁸, Michael W. I. Schmidt⁹, Markus Reichstein², Nuno Carvalhais^{2,10}, Philippe Ciais¹¹, Lifen Jiang¹², Johannes Lehmann¹³, Ying-Ping Wang¹⁴, Benjamin Z. Houlton¹⁵, Bernhard Ahrens², Umakant Mishra^{16,17}, Gustaf Hugelius⁷, Toby D. Hocking⁶, Xingjie Lu¹⁸, Zheng Shi¹⁹, Kostiantyn Viatkin^{3,13}, 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 Huang¹✉ & Yiqi Luo¹²✉

Microbial carbon use efficiency promotes global soil carbon storage

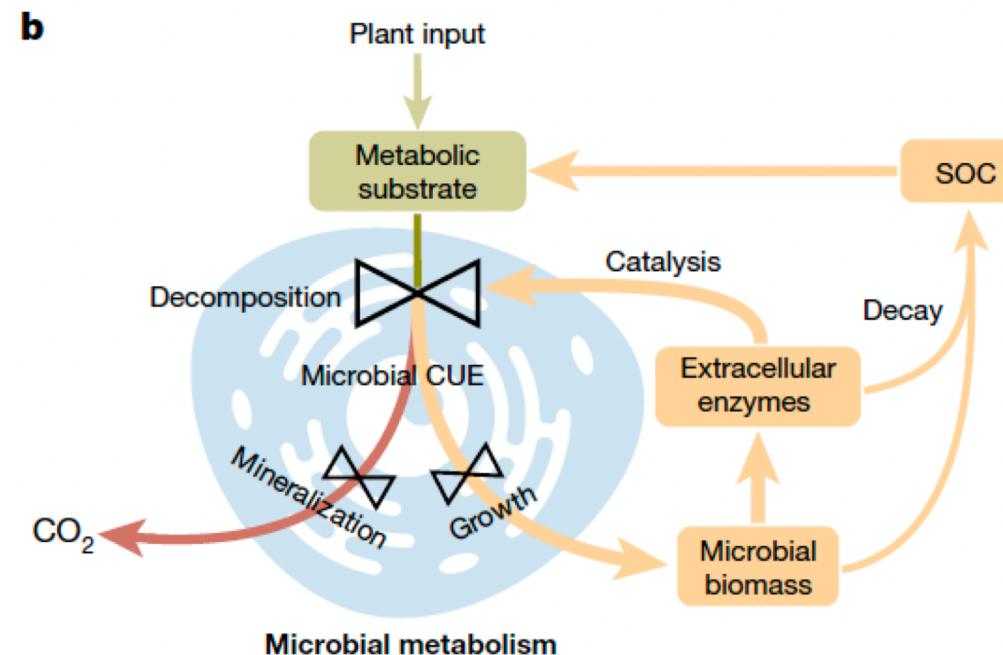
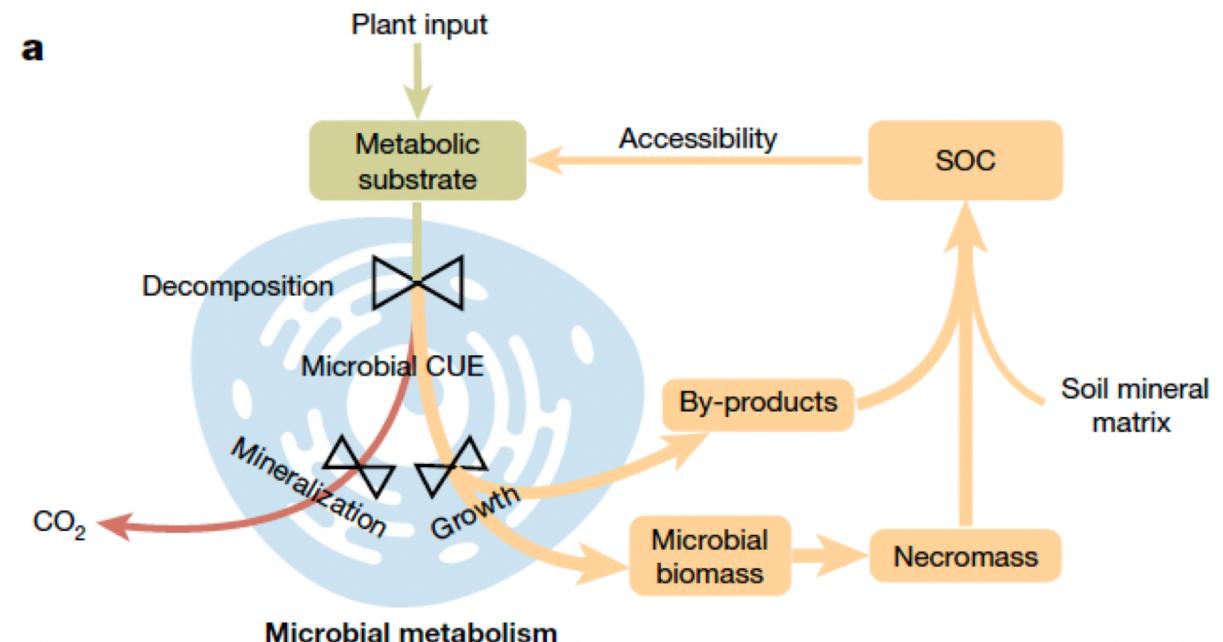
Nature Com 2023

<https://doi.org/10.1038/s41586-023-06042-3>

Two contrasting pathways in determining the relationship between microbial CUE and SOC storage.

a, **The first pathway** indicates that a **high CUE** favours the accumulation of **SOC storage** through increased **microbial biomass** and **by-products**.

b, **The second pathway** emphasizes that a **high CUE stimulates SOC losses** via increased microbial biomass and subsequent **extracellular enzyme production** that enhances SOC decomposition



Microbial carbon use efficiency promotes global soil carbon storage

They find that **CUE** is at least **four times as important as other evaluated factors**, such as carbon input, decomposition or vertical transport, in determining SOC storage and its spatial variation across the globe. In addition, **CUE shows a positive correlation with SOC content**. Our findings point to microbial CUE as a **major determinant of global SOC storage**. Understanding the microbial processes underlying CUE and their environmental dependence may help the prediction of SOC feedback to a changing climate.

Merci pour votre attention