



Manure and straw returning benefit for stability of soil microbial ecosystem



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Webinar series SUSTAINABLE MANAGEMENT OF BLACK SOILS



Background

• Manure with soil microbial diversity

Straw returning with soil microbial diversity

Global Black Soil Distribution map (GBSmap)



Table 2.2 Land cover and population in black soils

| | Black soils | World | Percentage |
|-----------------------------|-------------|--------|------------|
| Area (million hectare) | 725 | 12 995 | 5.58 |
| Cropland (million hectare) | 227 | 1 308 | 17.36 |
| Forest (million hectare) | 212 | 4 496 | 4.72 |
| Grassland (million hectare) | 267 | 3 129 | 8.52 |
| Population (million people) | 223 | 7 788 | 2.86 |
| | | | |

- Soils are the source of 95 percent of our food.
- Black soil represents only 5.6 percent of the global land area.
- While 31 percent of global black soils are cultivated, great portions remain with their natural land cover of forests and grasslands.

Table 2.1 Top ten countries with the largest black soil areas

| Country | Black soil area (million hectare) | Country area (million hectare) | Black soil proportion (percentage) |
|--------------------------|--------------------------------------|-----------------------------------|---------------------------------------|
| Russian Federation | 326.8 | 1700.2 | 19.22 |
| Kazakhstan | 107.7 | 283.9 | 37.93 |
| China | 50 | 934.6 | 5.35 |
| Argentina | 39.7 | 278.1 | 14.28 |
| Mongolia | 38.6 | 156.5 | 24.67 |
| Ukraine | 34.2 | 60 | 57.01 |
| United States of America | 31.2 | 950.1 | 3.28 |
| Colombia | 24.5 | 113.8 | 21.54 |
| Canada | 13 | 997.5 | 1.30 |
| Mexico | 11.9 | 196.4 | 6.04 |

World's soil under threat





- 2-3mm/a, 50-60cm to 30cm.
- Soil pH decreased 1.28-1.51 after continuous input of chemical fertilizers for 19 to 24 years.









Zhang *et al*, 2023



Soil organic carbon (SOC) contents in China



Soil Microbial feed humans



Biodiversity determined the stability of ecosystem

Dolfing et al. Sciences, 2004

Science The Microbial Engines That Drive Earth's Biogeochemical Cvcles MAAAS Science 320, 1034 (2008) DOI: 10.1126/science.1153213



The Microbial Engines That Drive Earth's Biogeochemical Cycles





Soil biodiversity and soil community composition determined ecosystem **multifunctionality** Webinar series | SUSTAINABLE MANAGEMENT OF BLACK SOILS

GLOBAL SOIL

Falkowski et al., Sciences, 2008

Wang, et al., PNAS, 2014

Biodiversity decreased by human activities



Organic materials are considered as remediation measures of degraded soil

- Main husbandry production areas.
- Abundant organic materials, straw and manure.
- Reduce environment pollution and GHG emission.



How organic materials affect the diversity of soil microbial community in black soil? How organic materials affect the ecological functions of soil microbial community in black soil?



Diversity and composition

- ➢Organic manure increased the abundance of soil bacterial community and diversity.
- ➢Organic manure application benefit for stability of bacterial community.



PC1(71.02%)





| | Ace | Chao | Coverage | Shannon | Simpson |
|-----|------------------------|----------------------------|-------------------------|------------------|-----------------------|
| NoF | 2538.24±130.92ab | 2516.57±144.28ab | 0.9850 ± 0.001 | 6.19±0.07a | $0.0057 \pm 0.0008b$ |
| М | $2607.90 \pm 16.42 ab$ | $2609.76 \!\pm\! 29.10 ab$ | 0.9847 ± 0.0008 | $6.25 \pm 0.06a$ | $0.0045 \pm 0.0003 b$ |
| CFM | 2720.91±95.70a | $2722.38 \!\pm\! 66.74a$ | $0.9875 \!\pm\! 0.0066$ | $6.18 \pm 0.04a$ | $0.0059 \pm 0.0005b$ |
| CF | 2386.63±126.4b | 2388.23±136.56b | 0.9858 ± 0.00145 | $5.92 \pm 0.12b$ | $0.0085 \pm 0.0012a$ |

Diversity and composition

> The relative abundance of *Acidobacteria*, *Entotheonellaeota* increased by manure application



Diversity and composition





- Long-term inputs of manure contributed to increases in soil nutrient levels and SMBC in comparison to no fertilization, and also had positive effects on copiotrophic taxa.
- With respect to increasing root exudates and root biomass due to crop growth, readily metabolic C in manure .
- Higher diversity index were found in M and CFM Hu *et al.*, Pedosphere, 2018 Shen *et al*, Applied Soil Ecology, 2010

Diversity and composition



- Organic fertilization significantly increased the diversity and abundance of fungi community.
- \succ High total carbon content led to high fungi biomass and gene abundance.
- The key species was *Penicillium*
- Complex fungal network.



Hu et al, Agriculture, Ecosystems and Environment, 2017 Hu et al, Chinese Journal of Applied Ecology, 2018 Ding et al, Applied Soil Ecology, 2017

Organic fertilization with soil P



Soil organic matter and pH have important effects on the change in soil Olsen P by 1 kg ha⁻¹ of P balance

Zhan et al., Plos One, 2015

P activation coefficient (PAC)

Organic fertilization with soil carbon



Long-term manure applications improved the C sequestration not only in the topsoil but also in the deep layers;

Abra et al., Journal of Soils and Sediments, 2020

The molecular structure of HA in Black Soil tends to be aliphatic, simpler, and younger after the application of manure Zhang et al., Plos One, 2017

C+J

350

350

400

Em. (nm)

450

500

400

Em. (nm)

C+K

450

500

Ecosystem functions



- Fertilization significantly increased the capability of C, N and P transformation.

M/CF

- Organic fertilization increased the relative abundance of functional genes involved in Cellulose, C degradation stage and C fixation.
- Manure application increased the course of soil denitrification.





Functional genes diversity



| Treat- SB | | | MB | | NB | |
|---------------|-----------------|----------------------|-----------------|----------------------|-----------------|----------------------|
| ment | OTU richness | Shannon diversity | OTU richness | Shannon diversity | OTU richness | Shannon diversity |
| NoF | $257a^{b)}$ | 3.01b | 162a | 3.68a | 195b | 3.74a |
| \mathbf{CF} | 244a | 3.32ab | 133b | 2.97b | 138c | 3.03b |
| Μ | 260a | 3.61a | 125b | 3.39a | 234a | 3.80a |
| CFM | 257a | 3.65a | 112b | 2.63b | 206b | 3.03b |

M and CFM regimes significantly increased the abundances and the diversity of *nir*S-type denitrifiers compared with NoF

Hu et al., Pedosphere, 2020

Ecosystem functions



- Organic fertilization enriched the relative abundances of *Proteobacteria* and *Planctomycetes* that carry C and N cycling genes, while inhibited the growth of oligotrophic groups such as Verrucomicrobia.
- Manure fertilization, particularly the combination of chemical and organic fertilizers (CFM), significantly enhanced the abundance of icd (rTCA cycle), which are involved in C fixation.
- The substantial enhancement in soil P contents induced by manure addition predominantly affected the C and N cycling profiles, abundance of functional genes and microbial taxa Hu *et al.*, Geoderma, 2022

GC

GS

WC

Diversity and composition



Alpha diversities of soil microbial metabolic function in different amounts of corn stover mulch treatments

| 处理 | Simpson多样性指数 | Shannon多样性指数 | Evenness均匀度指数 |
|-----------|-----------------------|------------------------|------------------------|
| Treatment | Simpson_1-D | Shannon_H | Evenness_e^H/S |
| NT0 | $0.773 \pm 0.005 \ b$ | $1.837 \pm 0.048 \; b$ | $0.485 \pm 0.024 \ b$ |
| NT33 | 0.846 ± 0.025 a | 2.161 ± 0.098 a | 0.677 ± 0.063 a |
| NT67 | 0.855 ± 0.017 a | 2.161 ± 0.083 a | 0.675 ± 0.054 a |
| NT100 | 0.862 ± 0.013 a | $2.200 \pm 0.066 \; a$ | $0.698 \pm 0.045 \; a$ |



Long-term corn straw cover (14 a) could improve soil microbial carbon metabolic functions by reducing microbial urs metabolic demand for readily available carbon sources and increasing microbial carbon metabolic diversities. Liu et al., Chinese Journal of Soil Sciences, 2023

- Arable soils with a low organic carbon content have a potential to get a higher and more active microbial diversity with amendments
- The majority of the differentially expressed transcripts produced at straw addition belonged to Bacteria.

Ecosystem functions







The straw addition triggered the upregulation of a set of enzyme families catalysing the organic matter degradation

Xie et al. Field Crops Research, 2021

Kozjek et al., SBB, 2023

Ecosystem functions

Predicted annual changes in topsoil (0-20 cm) SOC stocks (Mg ha⁻¹ yr⁻¹) under four scenarios at each experimental site over a 30 yar period

| Scenario | Heihe | Hailun | Harbin | Gongzhuling |
|-------------|-------|--------|--------|-------------|
| Straw 40% | -0.07 | -0.07 | 0.05 | 0.03 |
| Straw 100% | 0.36 | 0.39 | 0.53 | 0.53 |
| Manure 100% | -0.10 | -0.04 | 0.08 | 0.10 |
| Manure 200% | 0.01 | 0.05 | 0.17 | 0.19 |

• Increasing inputs of carbon sources such as straw and manure, together with appropriate tillage, could substantially improve SOC sequestration in the black soil region of northeastern China.

Wang et al., Journal of Soils and Sediments, 2019



Diversity and composition



1979, Heihe, soil type is black soil



Compared with CF, straw addition didn't significantly change the composition and diversity of bacterial community.
Yan et al., 2020 confirmed no significant affect on diversity of soil microbial community.

Diversity and composition



- *Proteobacteria, Acidobacteria, Actinobacteria, Gemmatimonadetes* and *Chloroflexi* are dominant group, accounting for more than 80%.
- The abundance of *Proteobacteria*, *Acidobacteria* increased in S, CF and CFS.
- The most significant difference species of S is Thermoleophilia in Phylum of *Actinobacteria*
- The most significant difference species of CFS is Acidobacteriia in Phylum of *Acidobacteria*.





Unpublished data

S CF CFS

Ecosystem functions



By identifying the gene node with the highest degree of network as a key node

The key node of straw application alone (S) treatment is *celF* gene related to cellulose decomposition, and the two genera with the highest degree of positive correlation are *Actinobacteria* g_{L} Kutzneria and the g_{L} Candidatus_Koribacters of *Acidobacteria*

The key node of CF is the malQ gene associated with starch metabolism and glycogen degradation, belonging to the phylum of *Acidobacterium*__Unclassified_ C__Acidobacteria

The key node of CFS is the cbbL gene related to carbon sequestration, belonging to the phylum of Proteobacteria____ Nitrobacter.

Ecosystem functions





代谢通路组间差异检验图

Unpublished data

Compared to the application of chemical fertilizers, the addition of straw increased the relative gene abundance of nitrite reductase (EC: 1.7.2.2) and glutamine synthase (EC: 6.3.1.2)

G1: NOF G2: S G3: CF G4: CFS

CK SOILS

Summary

- Organic fertilization increased the diversity of bacterial community, then straw returning almost no effect on diversity.
- Organic fertilization inhibited the growth of oligotrophic groups, then straw promote the copiotrophic groups.
- Organic fertilization enhanced the metabolic of carbon cycle, then straw retuning enhanced N fixation.
- Manure application decreased the course of soil denitrification







Thanks for your attention!

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