

Long-term effects of organic plant protection strategies in viticulture on soil quality and soil microbial communities

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Agricultural soils

Quality of agricultural soils is mainly measured by their productivity

• Productivity depends on their physical, chemical, and biological soil parameters

Grapevine microbiome consists of microbial communities in multiple habitats

- Its structure directly affects crop productivity and quality
- Different ways of managing crops can alter microbial community composition

Question: Does organic (ORG) and biodynamic (BD) management have a long-term impact on

- 1) Soil quality
- 2) enzymatic activity and microbial biomass in the soil
- 3) the soil + aboveground microbial community



Management systems

- Integrated (INT) according to good agricultural practice
- ORG according to Regulation (EU) No. 2018/848 and ECOVIN guidelines
- BD according to Regulation (EU) No. 2018/848 and demeter standards



	INT	ORG	BD
cover crop	winter cover crops + gras mixture (alternating)	winter cover crops + perennial mixture (Wolff-mixture or others)	
under-vine management	herbicides	mechanically	
fertilization	composted biological waste, mineral fertilization (N _{min})	composted farmyard manure	composted farmyard manure + biodynamic compost preparations (or cow pat pit preparation)
		ploughing of cover crops	
plant protection	synthetic fungicides, mating	copper (max. 3 kg/ha and year), sulfur / bicarbonate +	
	disruption	adhesives, mating disruption	
biodynamic	_	_	horn manure + horn silica,
preparations			compost preparations

Trial setup

- Trial site: Geisenheimer Mäuerchen (M4)
- Geisenheim/ Rheingau (49°59'; 7°56')
- Planted in 1991; ~ 1 ha
- Spacing of 2 m between rows
- Spacing of 1.2 m within rows
- VSP system
- Riesling Gm 198-30
- rootstocks: Börner, SO4 (randomly distributed)
- Soil: sandy/clayey loam
- hortic anthrosol (due to long viticultural use)
- 4 field replicates







Pictures: J. Döring



Assessment of soil quality and soil microbial biodiversity





- Bulk density in INT viticulture significantly enhanced
- BD treatment shows lowest bulk density
- Systems did not differ in soil pH, C/N, %N, available water capacity (AWC)



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- Bulk density in INT viticulture significantly enhanced
- BD treatment shows lowest bulk density
- Soil organic carbon (SOC) content significantly higher in BD viticulture → higher SOC in all three positions





- Copper content (total + bioavailable) significantly enhanced in ORG and BD treatments
- Total copper:
 - int 74 mg/kg soil org and bd 87 mg/kg soil
- Bioavailable copper:
 - int 19 mg/kg soil org 24 mg/kg soil bd 22 mg/kg soil



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- Copper content (total + bioavailable) significantly enhanced in ORG and BD treatments
- Total copper:
 - int 74 mg/kg soil org and bd 87 mg/kg soil
- Bioavailable copper:
 - int 19 mg/kg soil org 24 mg/kg soil bd 22 mg/kg soil
- No significant effects of total + bioavailable copper content in inter-row



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Enzymatic activities May-August 2016

• 5 enzymatic activities:

urease UR

catalase CAT

phosphatase PHO



- substrate is added to the soil and is incubated; amount of fission products are measured photometrically (400nm)
- Sample preparation according to Alef et al. (1991)

Microbial community analysis (PLFAs and NLFAs) May-August 2016

- Phospholipid-derived fatty acids and neutral ٠ lipid fatty acids: chemotaxonomic markers in microbial ecology
- PLFAs are structural parts of cellular membranes \rightarrow estimation of total biomass: `population indicators`
- different markes for bacteria, fungi, protozoa (PLFAs) and arbuscular mycorrhizal fungi AMF (NLFAs)
- sample prepration according to Frostegård et al. (1993), measurements by GC-FID

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Enzymatic activities and PLFAs/NLFAs from May-August 2016

Parameter	INT (mean ± sd)	ORG (mean ± sd)	BD (mean ± sd)	treat
Soil Analysis				
N min $[NO_3-N \text{ kg ha}^{-1}]$	4.42 ± 2.43 b	12.27 ± 11.78 a	11.79 ± 6.81 a	**
RWC [%]	47.74 ± 11.15 -	47.01 ± 13.20 -	48.52 ± 12.73 -	ns
Enzymatic activity				
GLU [μg ρNP g ⁻¹ h ⁻¹]	564.52 ± 163.10 b	730.96 ± 176.79 a	753.96 ± 192.63 a	***
CAT [% of O ₂ released]	8.76 ± 3.35 b	13.38±6.78 a	12.61 ± 6.98 ab	*
UR [µg NH ₄ -N g ⁻¹ h ⁻¹]	24.07 ± 5.44 b	27.31 ± 5.35 ab	29.44 ± 4.92 a	*
DHA [μ g TPF g ⁻¹ h ⁻¹]	0.656 ± 0.137 b	0.714±0.152 ab	0.756±0.132 a	*
ΡΗΟ [μg ρΝΡ g ⁻¹ h ⁻¹]	217.81 ± 82.03 -	222.09 ± 73.95 -	227.8 ± 82.7 -	ns

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Enzymatic activities and PLFAs/NLFAs from May-August 2016

Relative quantities of bacteria (PLFA 16:1ω7), fungi (PLFA 18:2ω6), arbuscular mycorrhizae (NLFA 16:1ω5) and protozoa (PLFA 20:4 ω 6) markers in the cover-cropped rows of the different management systems across four sampling dates in 2016

Management

Biodynamic Integrated

- Fungal and bacterial biomass significantly enhanced in ORG and **BD** viticulture
- AMF biomass significantly increased in INT viticulture

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Enzymatic activities and PLFAs/NLFAs from May-August 2016





Soil microbial community

Species richness and community composition of fungi and bacteria in soil in August 2015

- · Soil sampling: management system, position, depth
- DNA extraction

PowerSoil DNA Isolation Kit, MoBio Laboratories Inc., Carlsbad, CA, USA)

- DNA amplification: ITS2, 16s-rRNA fungi: primer mix according to Ihrmark et al. 2012 bacteria: primer mix as modified by to Caporaso et al. 2012
- sequencing: Illumina Miseq
- bioinformatic data management
- comparison with database



https://www.scinexx.de/wp-content/uploads/0/1/01-35285-barcode_1.jpg

Hendgen et al. 2018

https://www.hellblaurosa.de/media/image/f0/7d/ba/selecta-memo-pepito-und-seine-freunde-memoryspiel-38380-40239.jpg https://jeltsch.org/sites/jeltsch.org/files/styles/large/public/field/image/PCR_tube_0.png?itok=V0DRJe9M

Soil microbial community: bacteria



Species richness of bacteria in soil in August 2015



- management system: org more species compared to int
- position: no difference
- depth: topsoil more species

compared tosubsoil

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Soil microbial community: bacteria

community composition of bacteria in soil in August 2015





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Soil microbial community: fungi

Species richness of fungi in soil in August 2015





• management system: no

difference

- position: inrow richer than undervine
- **depth**: topsoil richer than subsoil

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Soil microbial community: fungi



community composition of fungi in soil in August 2015



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Soil microbial community: fungi

Taxonomc classification





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Conclusion

- Management system highly affects soil quality as well as soil microbial biomass, enzymatic activities in the soil and fungal community composition below- and aboveground
- ORG and BD management are caracterized by
 - > higher SOC, lower bulk density, higher Cu contents and higher Nmin in soil
 - Higher enzymatic activities, higher microbial biomass of bacteria and fungi and lower biomass of AMF
 - Higher species richness of bacteria in soils (ORG)
 - > Completely different community composition of fungi in soil, on leaves and on grapes (BD)
- Main reasons for changes: spraying regime and cover crop mixtures → effects on crop productivity?



Thanks

- Thank you for your attention
- More questions? Please don't hesitate to contact me: johanna.doering@hs-gm.de
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