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Agroscope

Diversity of arbuscular mycorrhizal fungi in agricultural systems

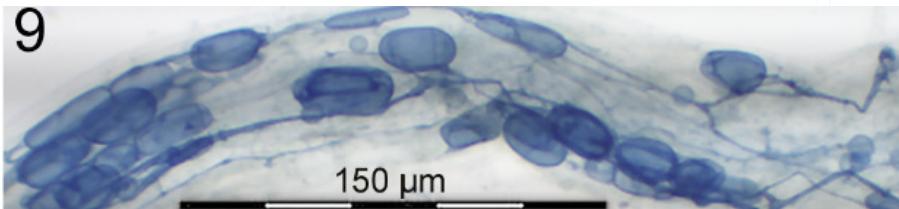
Fritz Oehl

Monte Verità, 8.10.2019

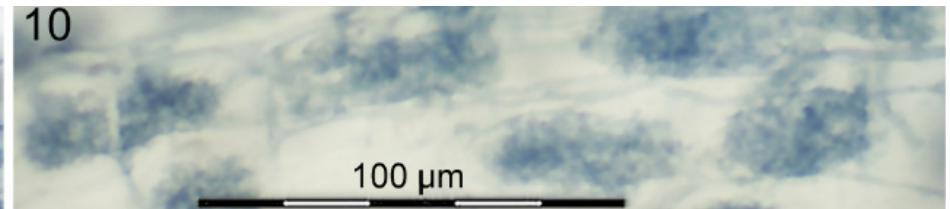
www.agroscope.ch | gutes Essen, gesunde Umwelt



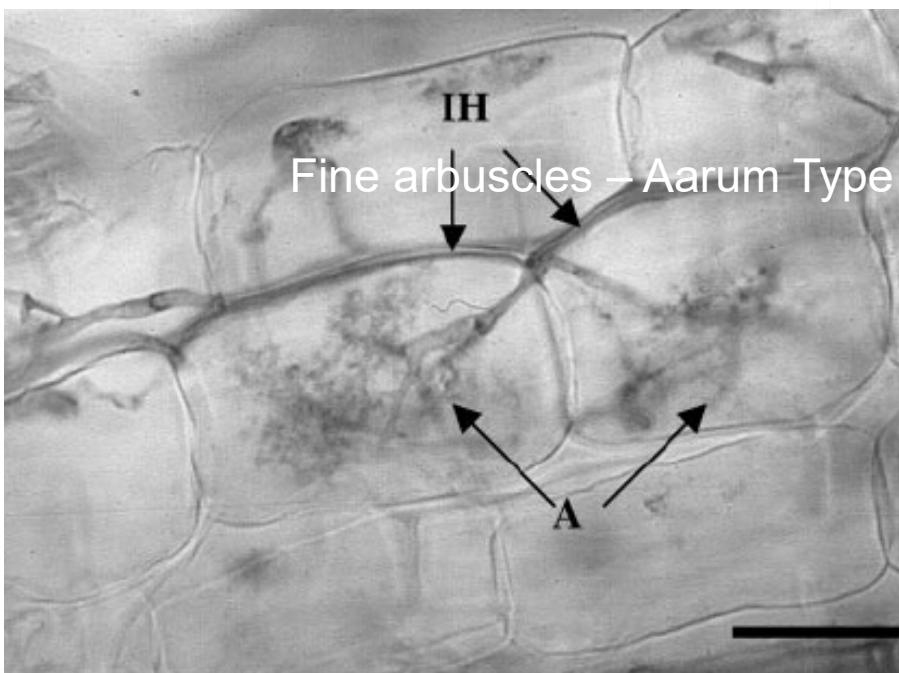
(Vesicular-)Arbuscular Mycorrhizal structures



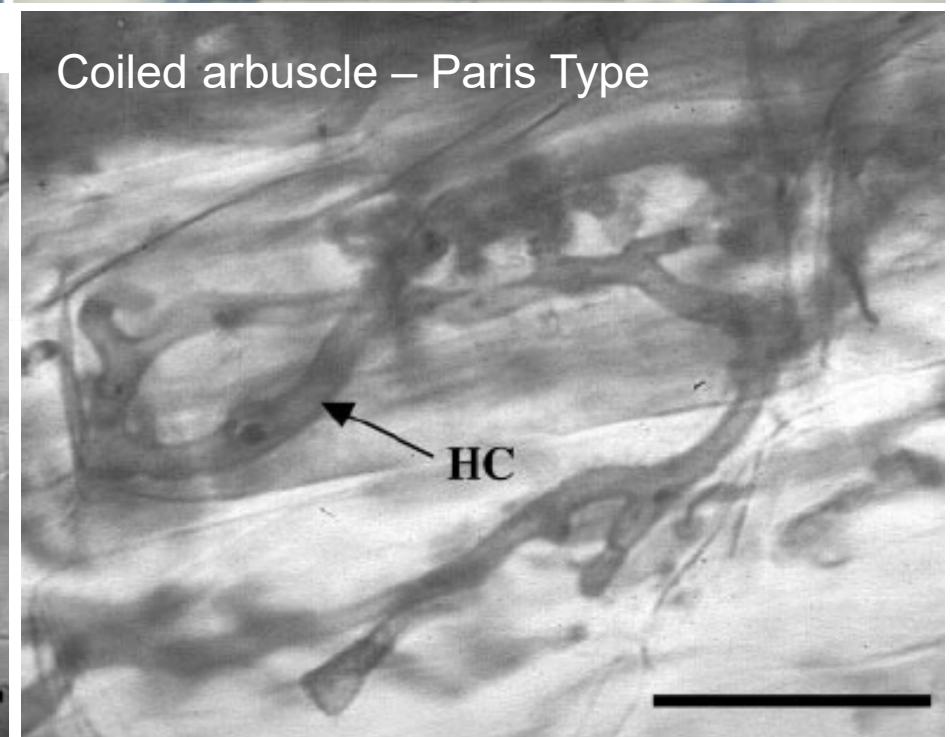
Corazon-Guivin et al. 2019



Coiled arbuscle – Paris Type



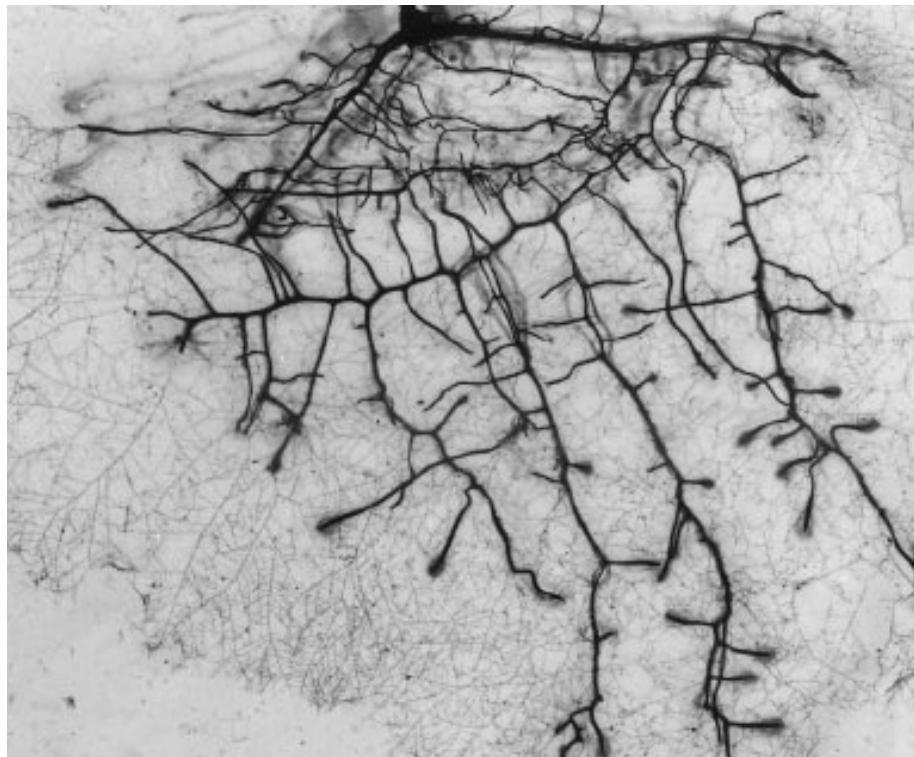
Cavagnaro et al. 2001





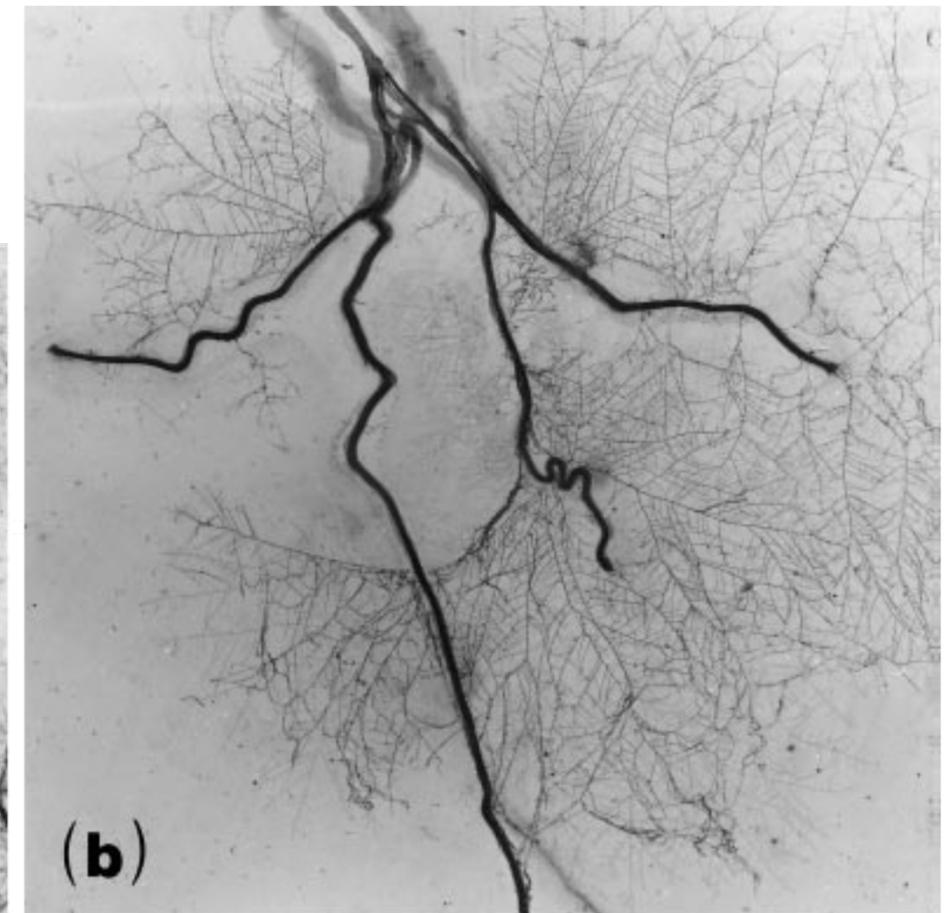
Extraradical mycelia

Cherry - *Prunus avium* -roots



AM fungi in agricultural systems
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Leek - *Allium porrum* - roots

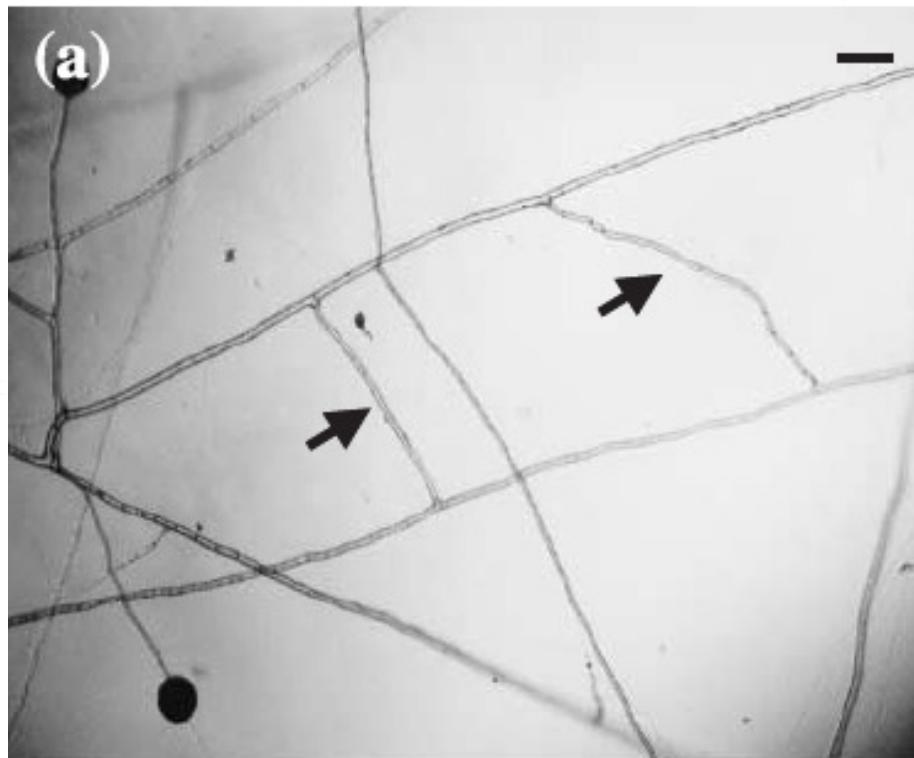


Giovannetti et al. 2001



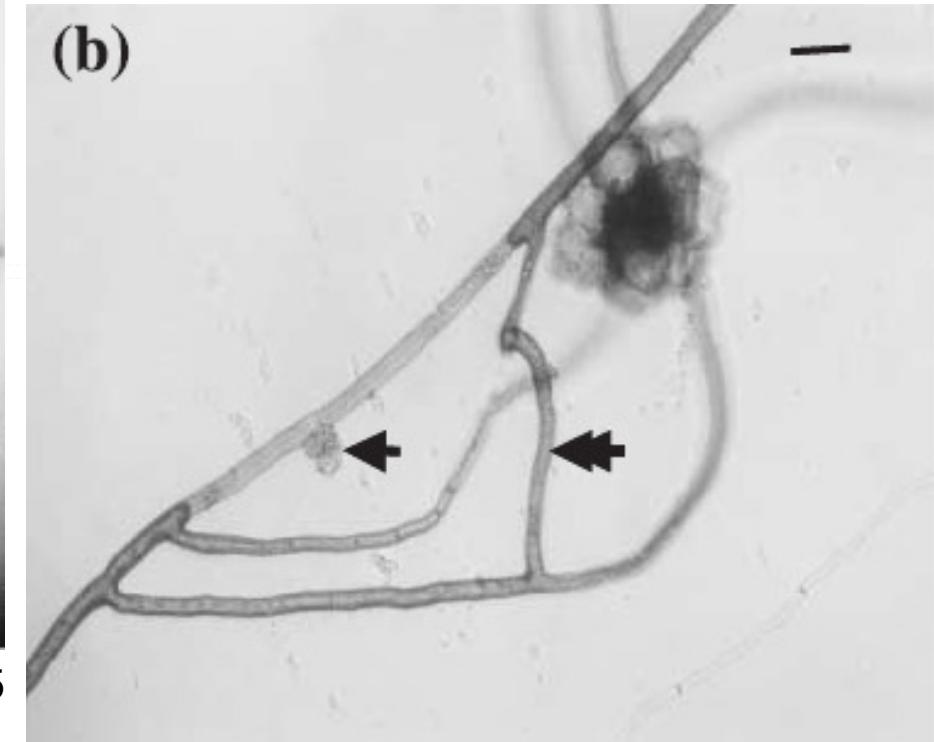
Extraradical mycelia

Glomerales



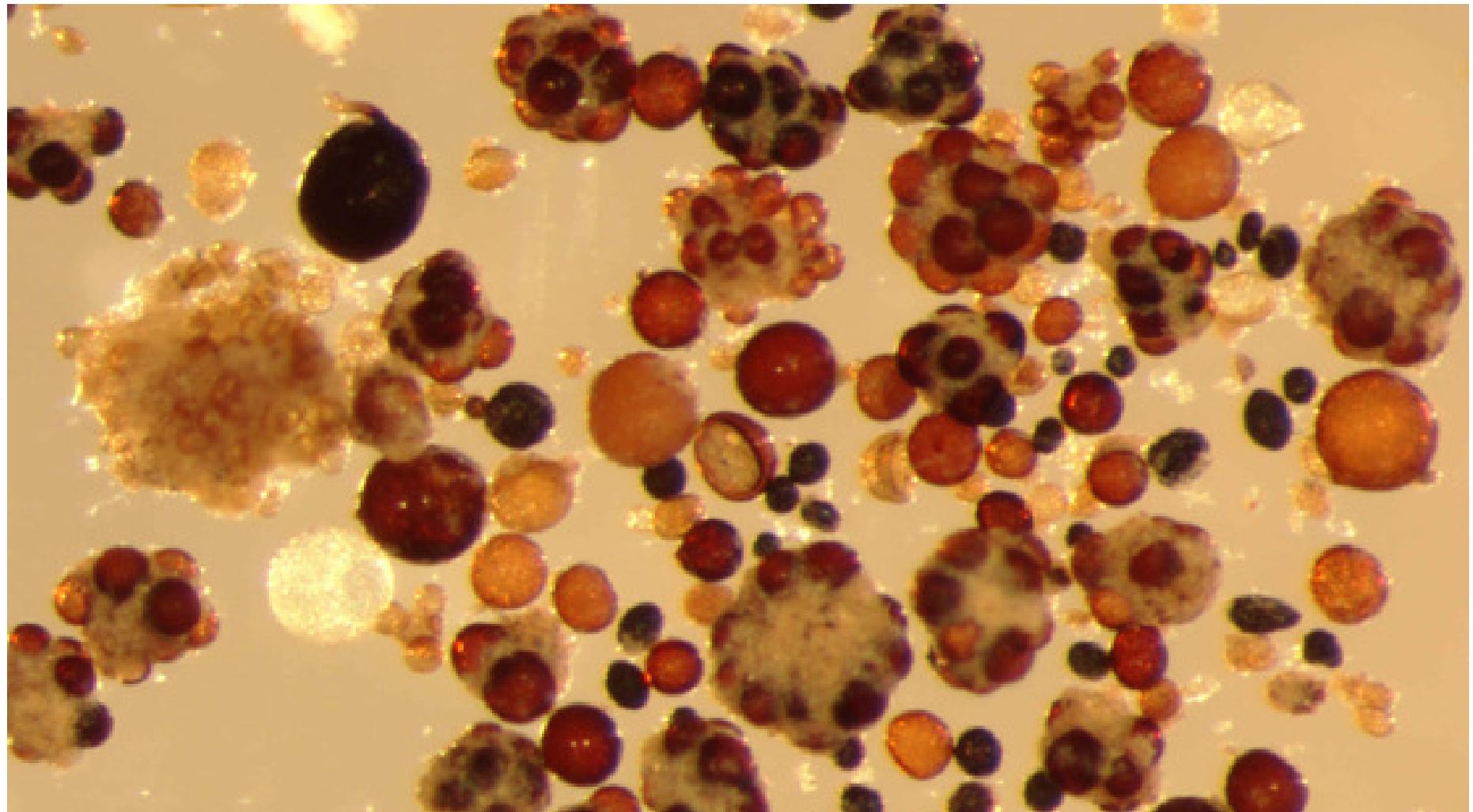
De la Providencia et al. 2005

Gigasporales



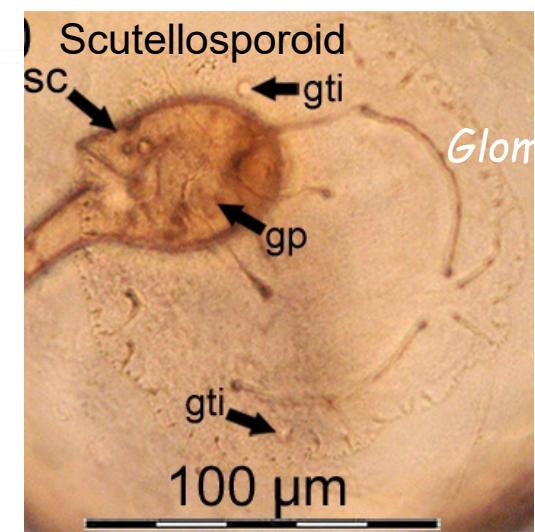
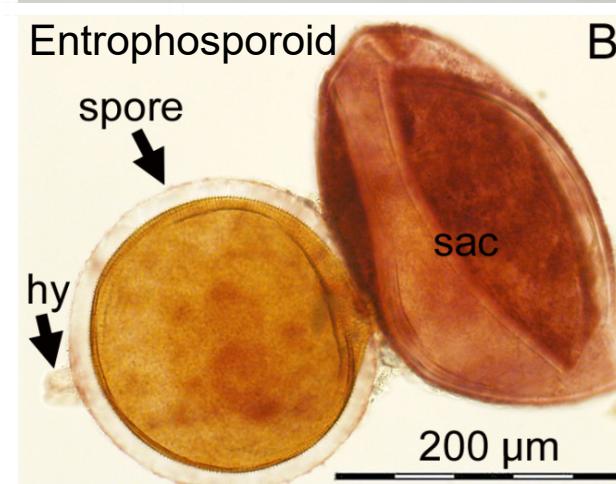
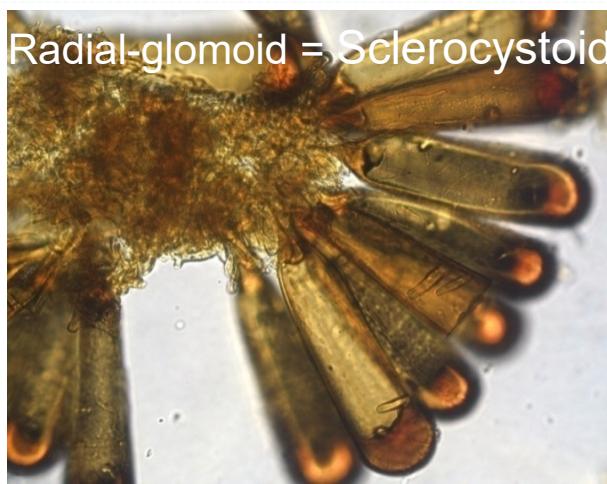
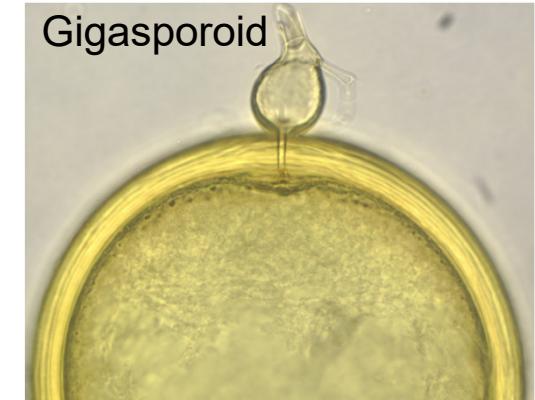
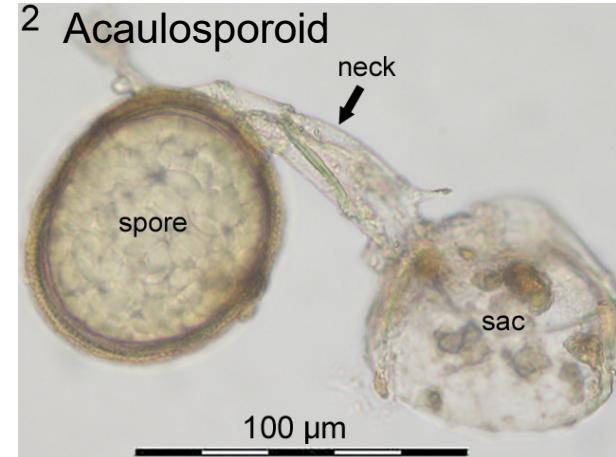


AMF spore population in a natural grassland in Europe



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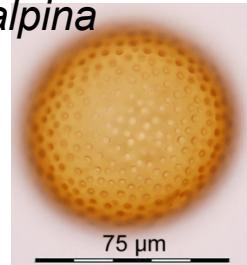
1999: 2x3 Spore formation types = 3 AMF families, 6 genera; ca. 150 species



2019: 16 AMF families, 49 genera, ca. 320 species



*Acaulospora
alpina*



Dominikia aurea



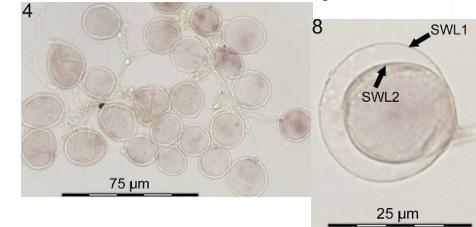
*Funneliglomus
sanmartinensis*



*Archaeospora
europaea*



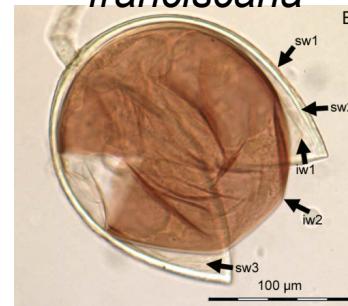
Microkamienskia peruviana



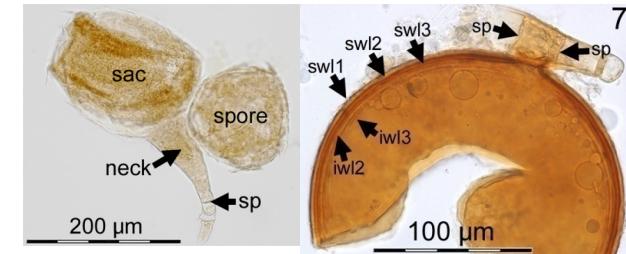
*Pacispora
robicina*



*Pacispora
franciscana*



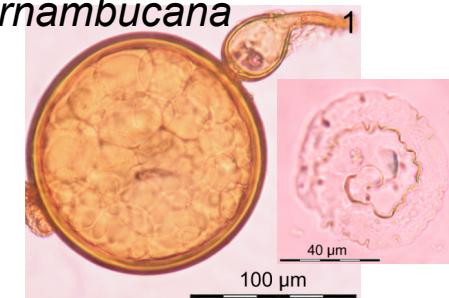
*Otospora
bareae*



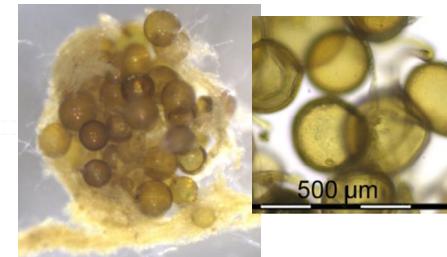
Glomus badium



*Orbispora
pernambucana*



Rhizoglomus venetianum



*Cetraspora
helvetica*



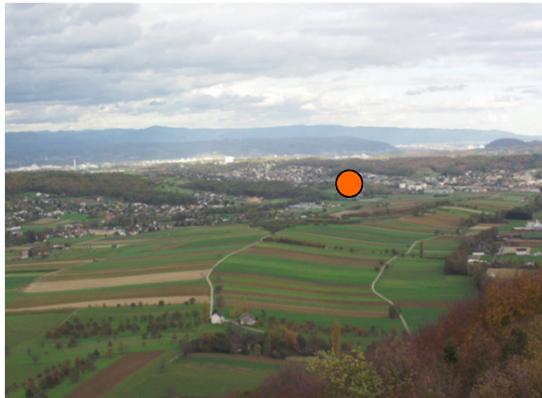
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Baltruschat et al. 2019 7



1. DOK Long-term field experiment in Therwil BL

Organic versus conventional farming since 1976/78



Geology: periglacial Loess

Soil type: Haplic Luvisol

Site: DOC field trial, Therwil (BL)

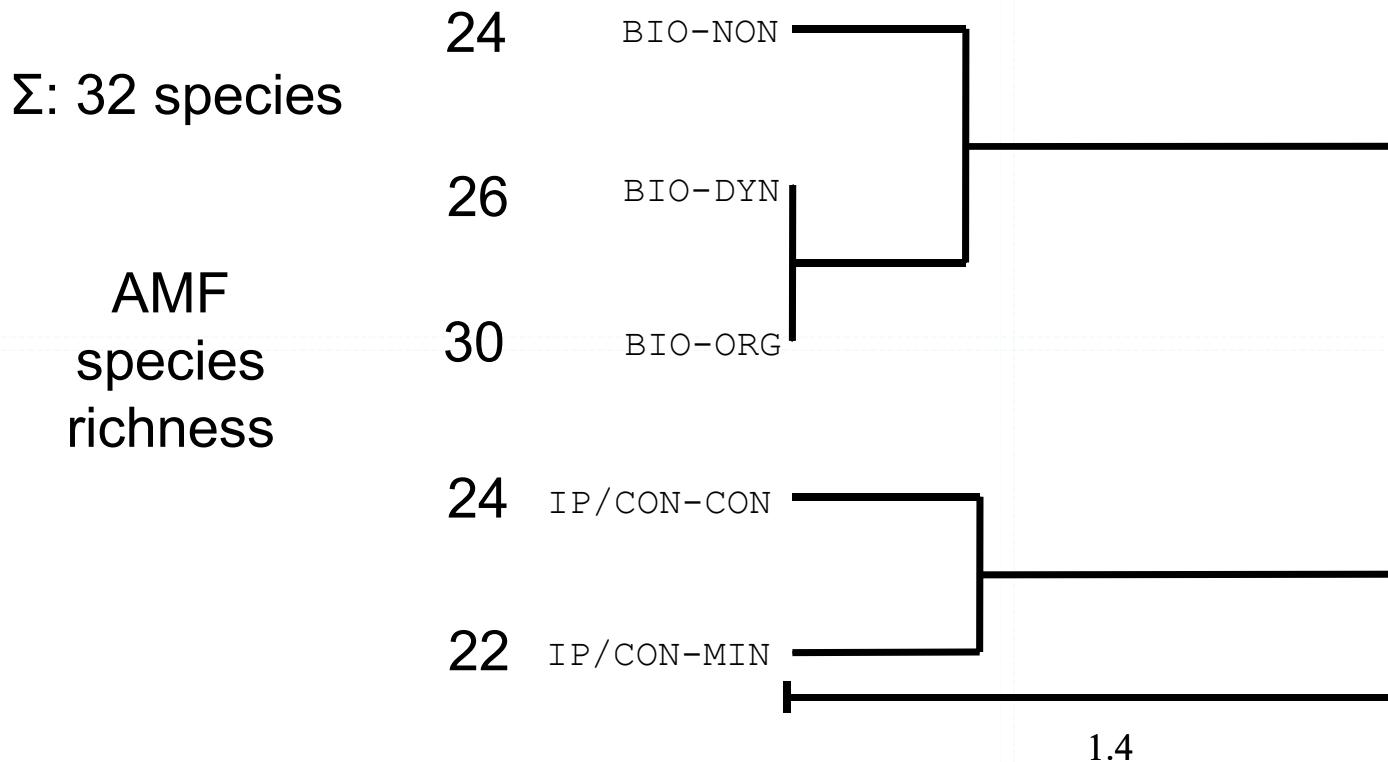


Land use intensity





Hierachical cluster analysis on AMF spore populations in conventional and organic farming systems



Oehl et al. 2004



Impact of chemical soil parameters on spore density of different AMF species

AMF orders / AMF species

		r (linear regression)			Weed species
		pH (H ₂ O)	Organic carbon	Available P (E ₁)	Available K
Glomerales/Paraglomerales					
<i>Oehlia diaphana</i>		-0.26	-0.48*	0.51*	0.42
<i>Funneliformis caledonius</i>		-0.36	-0.21	0.56*	0.63*
<i>Claroideoglomus etunicatum</i>		0.19	0.09	-0.33	-0.36
<i>Rhizoglomus fasciculatum</i>		0.06	0.09	-0.16	0.19
<i>Fu. mosseae</i>		0.28	0.08	-0.05	-0.10
<i>Dominikia compressa</i>		0.10	0.26	-0.14	-0.09
<i>Fu. geosporus</i>		0.00	0.08	-0.09	0.16
<i>Paraglomus laccatum, albidum & occultum</i>		0.29	-0.19	-0.27	0.46
<i>Septoglomus constrictum</i>		0.37	0.31	0.08	0.03
<i>Rh. invermaium</i>		0.19	-0.03	-0.20	-0.3
Diversisporales/Gigasporales					
<i>Pacispora dominikii</i>		0.62*	0.21	-0.51*	-0.20
<i>Scutellospora calospora</i>		0.10	0.24	-0.48*	-0.55*
<i>Cetraspora pellucida</i>		-0.27	-0.28	-0.48*	-0.58*
<i>Acaulospora paulinae & sieverdingii</i>		0.09	-0.14	-0.62*	-0.67*
<i>Ac. thomii</i>		0.13	-0.24	-0.49*	-0.55*
<i>Ac. laevis</i>		0.04	-0.15	-0.53*	-0.57*
<i>Ac. longula</i>		0.23	0.26	-0.70*	-0.58*
<i>Ac. scrobiculata</i>		0.21	-0.42	-0.66*	-0.57 *

AM fungi in agricultural systems
Fritz Oehl

Oehl et al. 2011

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2. 'Oberacker' Long-term field experiment in Rütti BE Tillage versus No-tillage, ÖLN/PEP conventional

- Since 1994
- Soil type: Sandy-loamy Luvisol/Cambisol, sandig-lehmige Parabraun-/Braunerde
- pH 5.7-6.2 in topsoil, pH 6.3-6.5 in subsoil
- 6y crop rotation:
- Sugar beet, winter wheat, winter protein peas, corn, broad bean, winter barley
- In collaboration with Wolfgang Sturny, Claudia Maurer, Andreas Chervet, Murielle Rüdy, Urs Zihlmann



Maurer et al. 2014



AM fungal species richness in 'Oberacker' long-term field experiment

	Species richness no-till	Stdev	Species richness tillage	Stdev
Winter protein peas	21		17	
Winter wheat	17		15	
Interim crops after wheat	17		14	
Winter barley	15		11	
Interim crops after barley	21		12	
Broad bean	20		10	
Mean species richness	18.5	2.5	13.2	2.6
Total species richness	33		21	

Maurer et al. 2014

After 20 months culturing in
the greenhouse on
grass/clover from winter
barley field

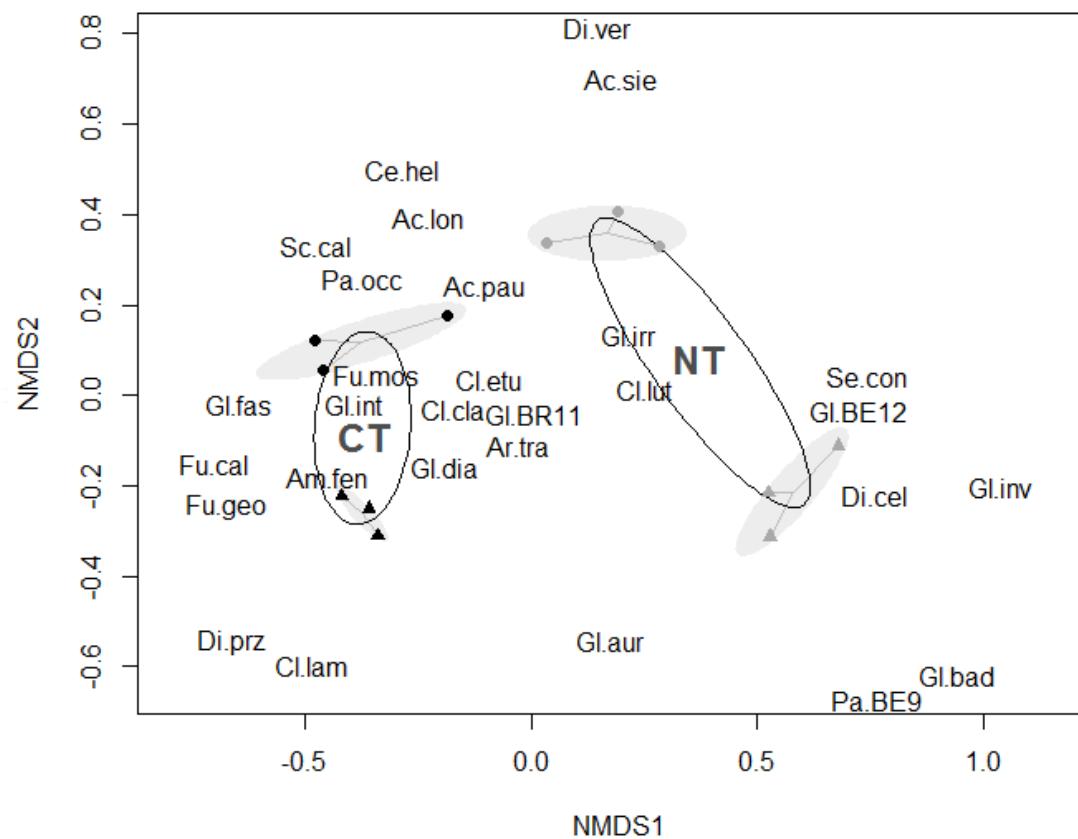
23

24

Köhl et al. 2014



Multidimensional scaling of AM fungal communities from Oberacker, in microcosms after 20 months



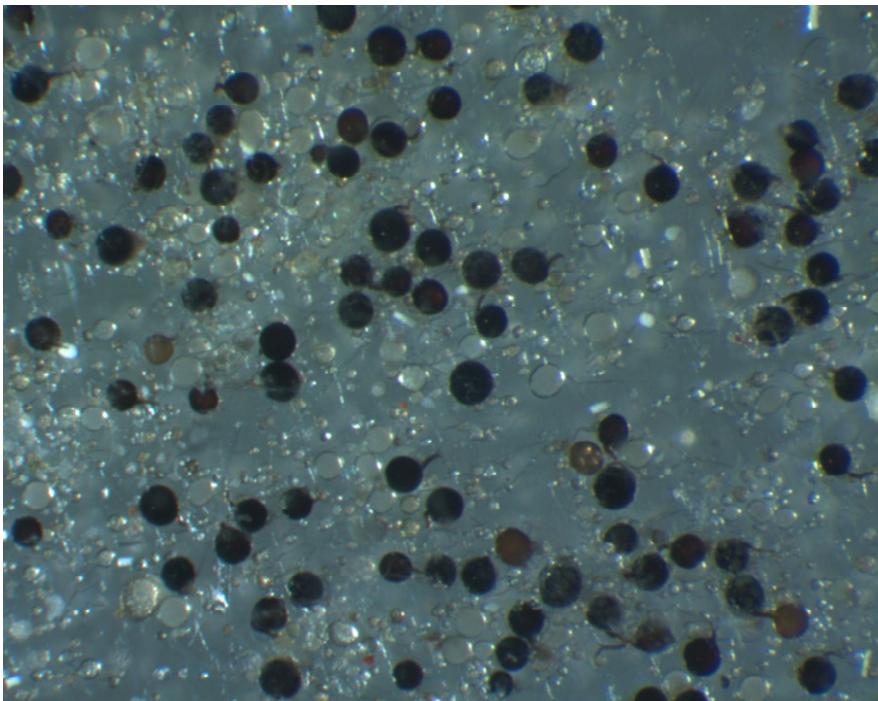
Köhl et al. 2014



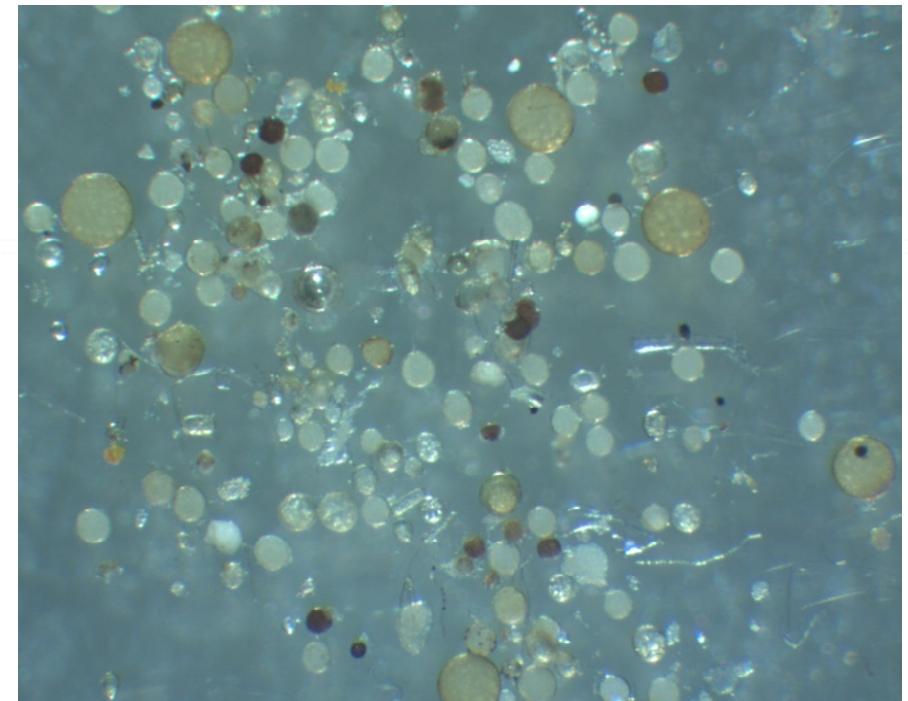


AM fungal spore communities from 'Oberacker' experiment

Conservation tillage



Tillage





3. Pinot Gris - On farm 'experiment' in Hainfeld DE Tillage versus No-tillage after 38 years, conventional

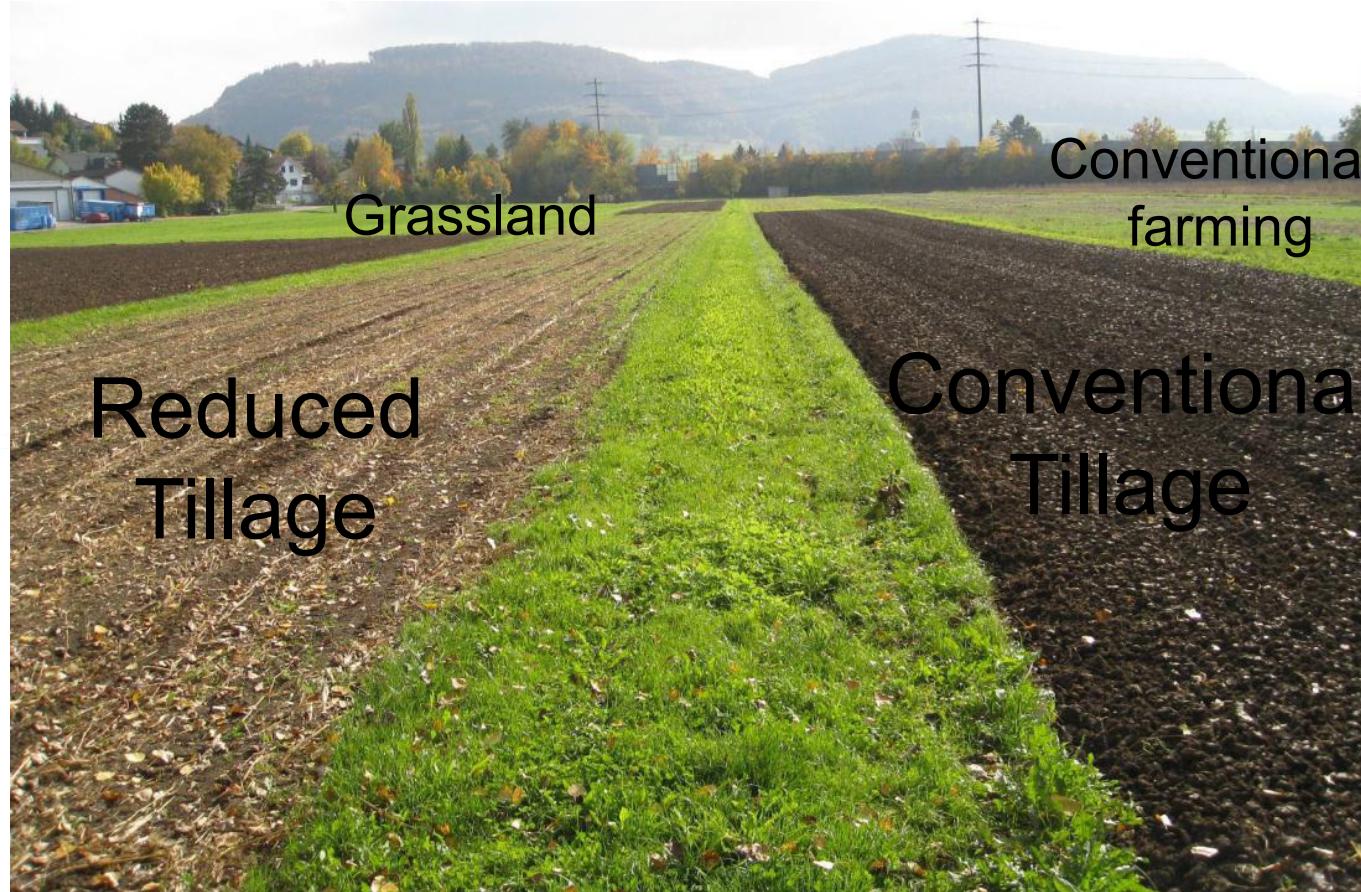


No-till		Tillage
27.2 (1.5)	Mean species richness	17.8 (1.5)
34	Total species richness	24

Oehl & Koch 2018



4. FiBL long-term field experiment in Frick AG Tillage versus Reduced-tillage, Bio-systems





4. FiBL soil tillage experiment



Reduced Tillage RT



Convent Tillage CT

- Since 2002
- In our study, conventional IP systems in the neighbourhood of the experiment were included, and a adjacent grassland from FiBL
- Soil type: clayey Cambisol, tonige Braunerde;
pH 7.5-7.7 in topsoil, pH 7.8-8.2 in subsoil
- 6y crop rotation in FiBL experiment:
> maize, winterwheat, sunflower, spelt, 2y grass-clover
- Part of PhD thesis of Verena Säle,
collaboration with Alfred Berner & Paul Mäder

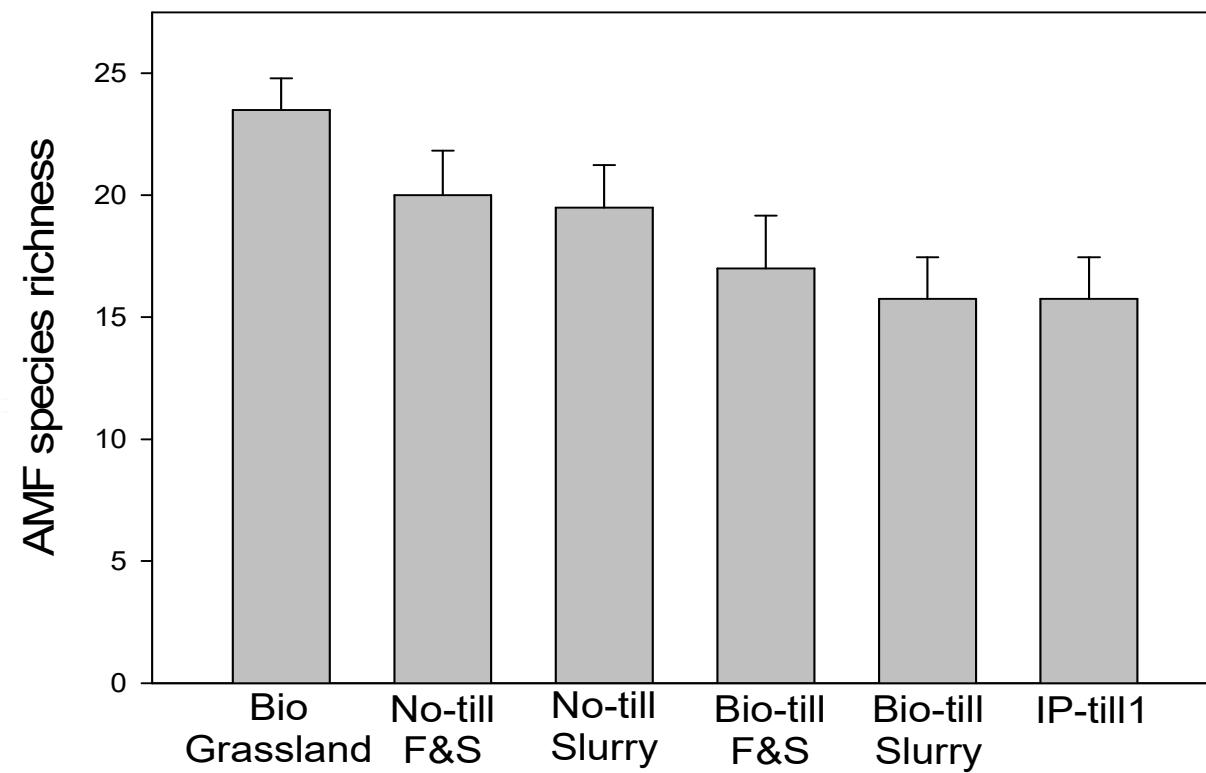


© FiBL



AMF species richness in the FiBL tillage experiment

AMF species richness: 38 33 33 33 28 28 Σ: 53 species



Säle et al. 2015



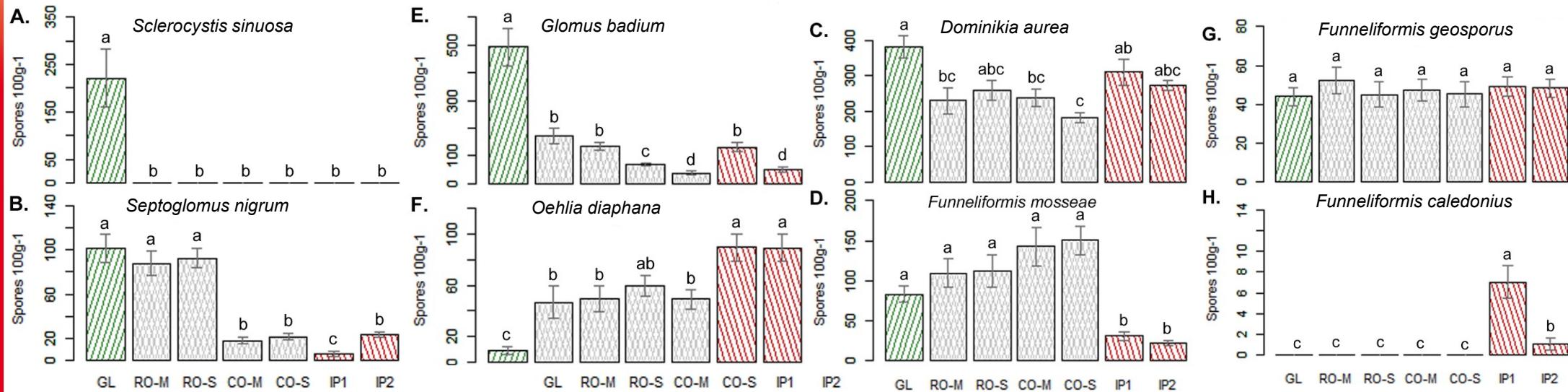
AMF species at study sites in Frick

60-80% of AMF species and > 90% of spores from **Glomerales & Paraglomerales**:
> typical for calcareous sites in Europe

3 Classes	5 Orders	8 Families	17 Genera	53 Species
Glomeromycetes	Glomerales	Glomeraceae	<i>Glomus</i> & <i>Rhizoglomus</i> & <i>Oehlia</i> <i>Funneliformis</i> & <i>Septoglomus</i> <i>Dominikia</i> <i>Sclerocystis</i>	12 11 2 3
		Entrophosporaceae	<i>Claroideoglomus</i> <i>Entrophospora</i>	3 1
	Diversisporales	Diversisporaceae	<i>Diversispora</i> <i>Pacispora</i>	2 1
	Gigasporales	Acaulosporaceae	<i>Acaulospora</i>	4
Archaeosporomycetes	Archaeosporales	Scutellosporaceae	<i>Scutellospora</i>	1
		Ambisporaceae	<i>Ambispora</i>	2
		Archaeosporaeae	<i>Archaeospora</i> <i>Palaeospora</i>	4 1
Paraglomeromycetes	Paraglomerales	Paraglomeraceae	<i>Paraglomus</i>	5



Selected AMF species in the FiBL tillage experiment - with or without indicator potential





AMF species richness in different soil types, climates & land use intensities

Soil type	Natural ecosystem type	Natural systems	Organic farming/ Low input	Reduced tillage systems	High-input systems
Calcaric Leptosol Oehl et al. 2010	Grasslands	27-33	20-25		21-23
Calcaric Regosol Oehl et al. 2003	Grasslands	24-31			22-24/13
Calcaric Chernosem Baltruschat et al.			26-33	23-27	16-19
Haplic Luvisol Wetzel et al. 2014 Oehl et al. 2003, 2004, 2005, 2009	Grasslands	26-32	25-31	25-33*	22-24 (IP Suisse) 16-19 (Conv.)
Humic Cambisol Oehl et al. 2010	Grasslands	32-39			21-25
Vertic Cambisol Säle et al. 2015	Grasslands	38	33-33	28-32	
Cambisol/Luvisols Maurer et al. 2018, *	Grasslands	33-35	26-30	25-27	20
Ferralsol (semi-humid to semi-arid) Tschabi et al. 2008	Sudan and Guyana savanna (forests)	28-38	15-19 (Yam fields)		5-10 (Cotton fields)
Ferralsol (semi-humid) Pontes et al. 2017a	Cerrado savanna forest	26-33		24-26	15-21 (-28)
Ferralsol (semi-arid) Pontes et al. 2017b	Caatinga dry savanna (forest)	44	29-36		
Ferralsol (semi-arid) Marinho et al. 2019	Caatinga dry savanna (forest)	51-56	25-42		



Summary and Conclusions

- Land use intensity, soil type and climate strongly affect AMF communities
- Low input systems generally have high AMF species richness and diversity, comparable with those of natural systems
- AMF indicator species can be named for different land use intensities, soils and climates
- In Central Europe, *Funneliformis caledonius* and *Oehlia diaphana* are representative AMF species for intensively managed agricultural systems
- Several AMF species are indicators for low-input agricultural systems, such as *Cetraspora helvetica* and *Gigaspora margarita*
- Others, such as *Glomus badium* and *Septoglomus nigrum*, are indicators for no- or reduced tillage systems
- It is still difficult to predict the beneficial potential of single AMF species in respect to their different ecosystem services and their environments
- A higher diversity of AMF fungi in soils usually is accompanied by a higher general soil biodiversity
- Both should lead to more active and biologically more buffered soils, and thus to a higher biological soil fertility and stability, and to enhanced plant growth and health





Danke für Ihre Aufmerksamkeit

Thank you for your attention!



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