Enzyme patterns in topsoil and subsoil horizons along a latitudinal transect in Western Siberia

Jörg Schnecker

Department of Microbiology and Ecosystem Science University of Vienna



Co-Authors











Field: Ricardo J. Eloy Alves Norman Gentsch Antje Gittel Nikolay Lashinskiy Robert Mikutta











Field:

Ricardo J. Eloy Alves Norman Gentsch Antje Gittel Nikolay Lashinskiy Robert Mikutta

Lab:

Angelika Hofer Karoline Klaus











Field: Ricardo J. Eloy Alves Norman Gentsch Antje Gittel Nikolay Lashinskiy Robert Mikutta

Lab:

Angelika Hofer Karoline Klaus Field/Lab/Data evaluation/Discussion:











Field:

Ricardo J. Eloy Alves Norman Gentsch Antje Gittel Nikolay Lashinskiy Robert Mikutta

Lab:

Angelika Hofer Karoline Klaus

Field/Lab/Data evaluation/Discussion:

Birgit Wild Mounir Takriti Anna Knoltsch Andreas Richter







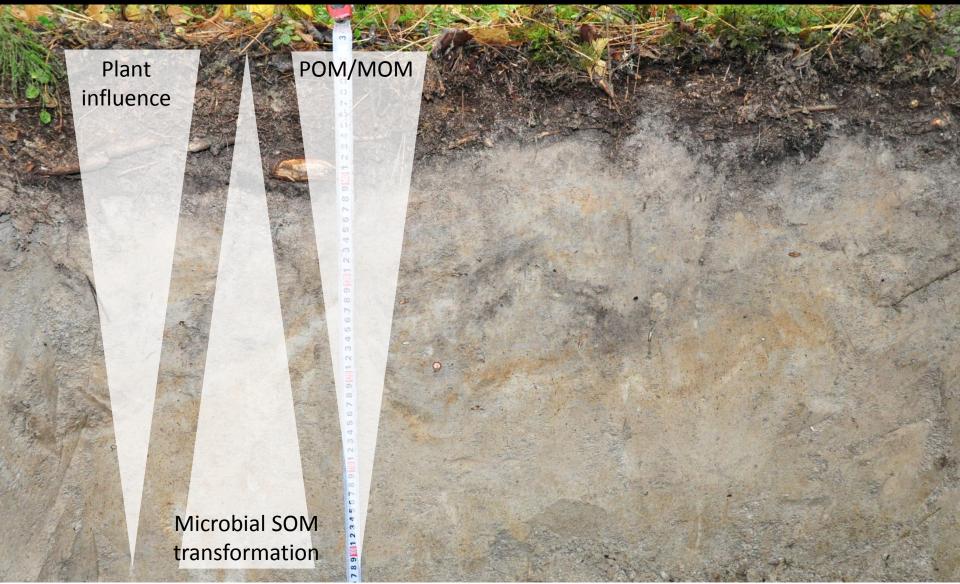


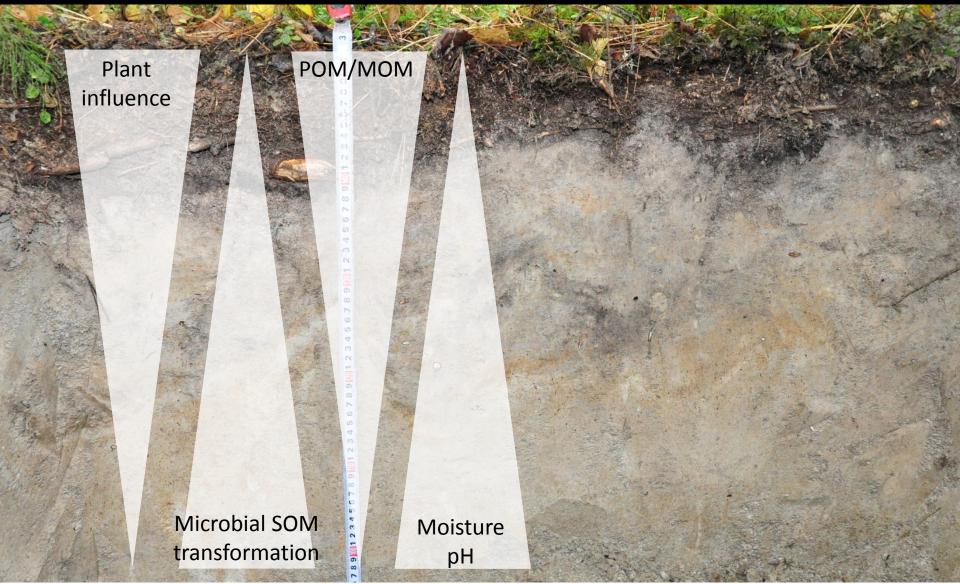


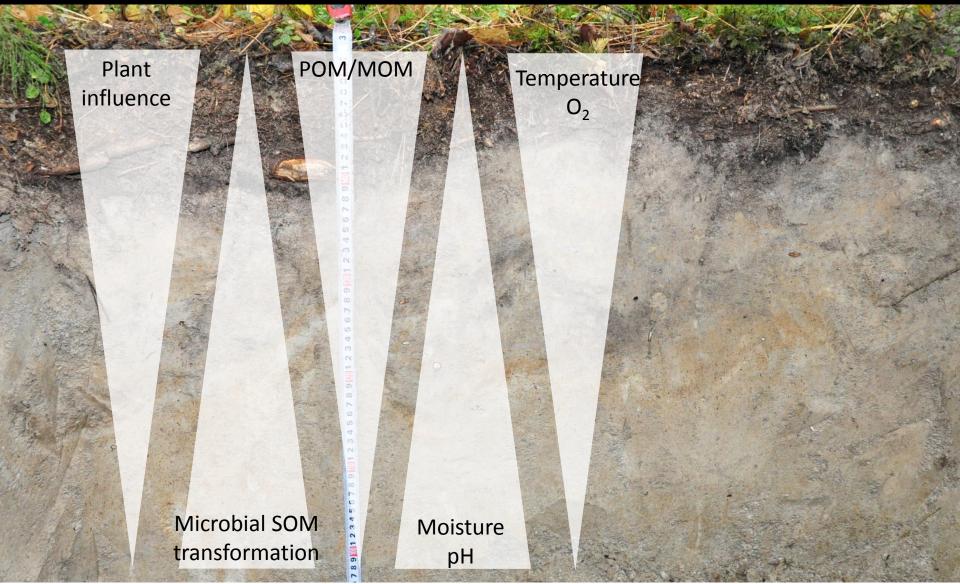


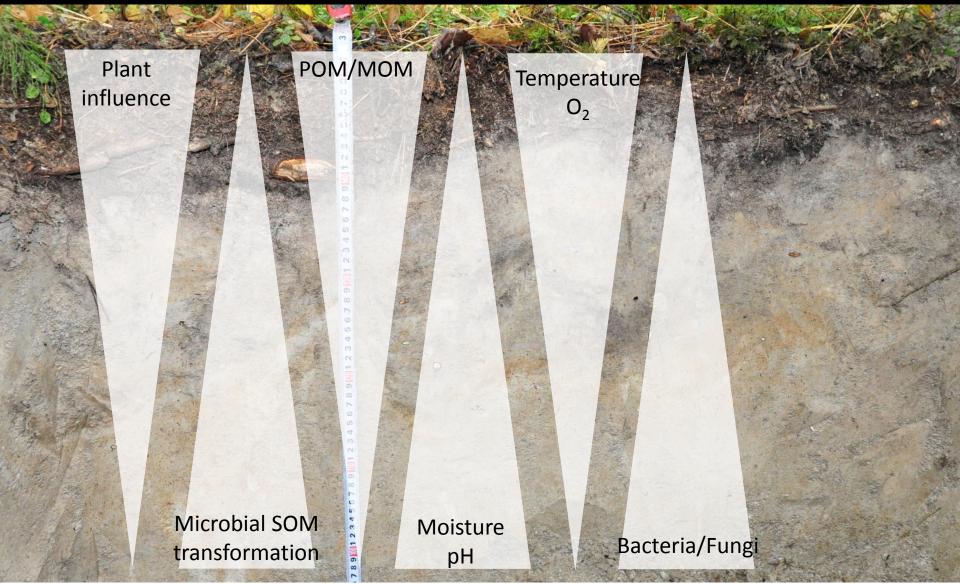


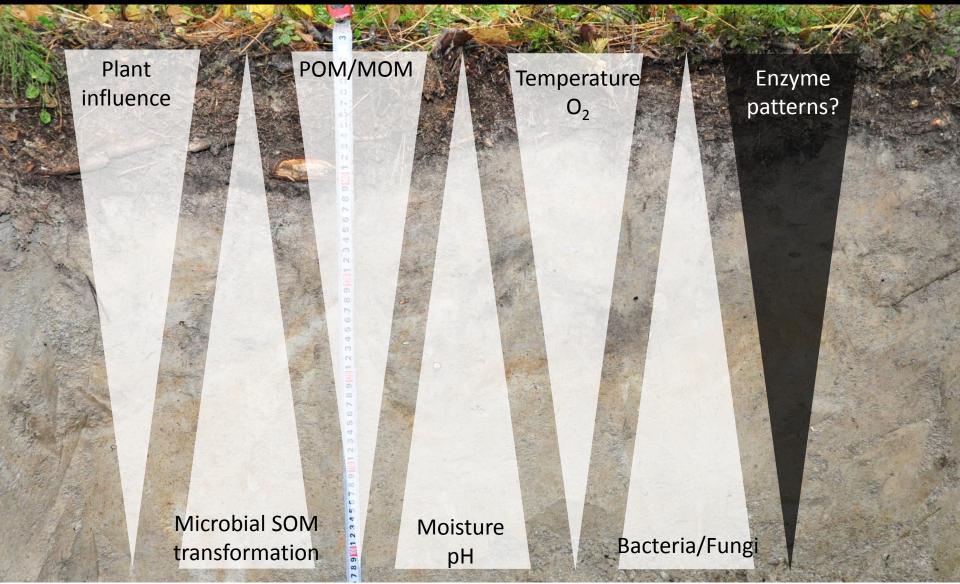




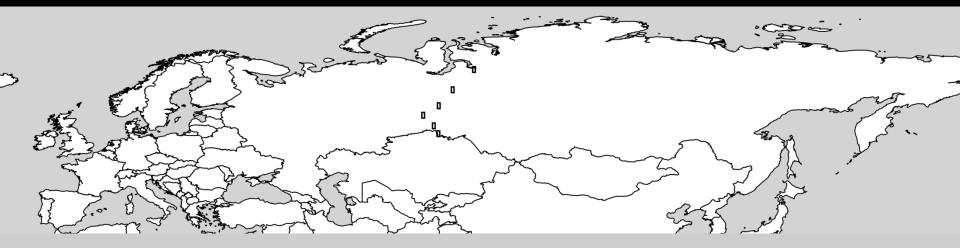








Sites



Tundra



Northern



Tundra



Middle taiga



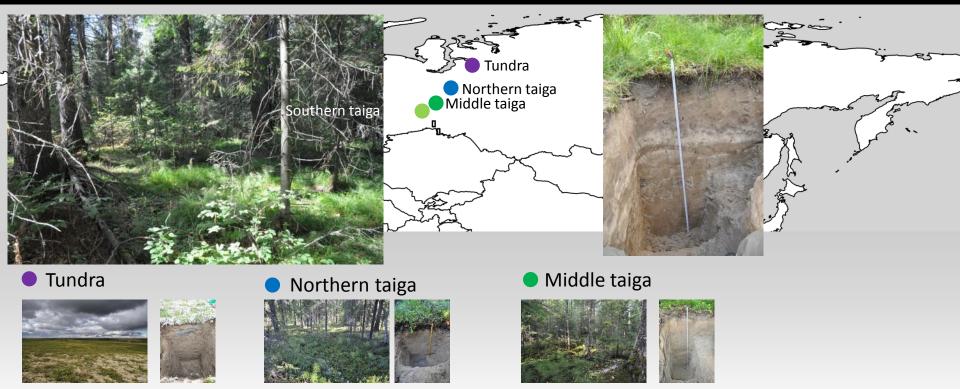
Tundra



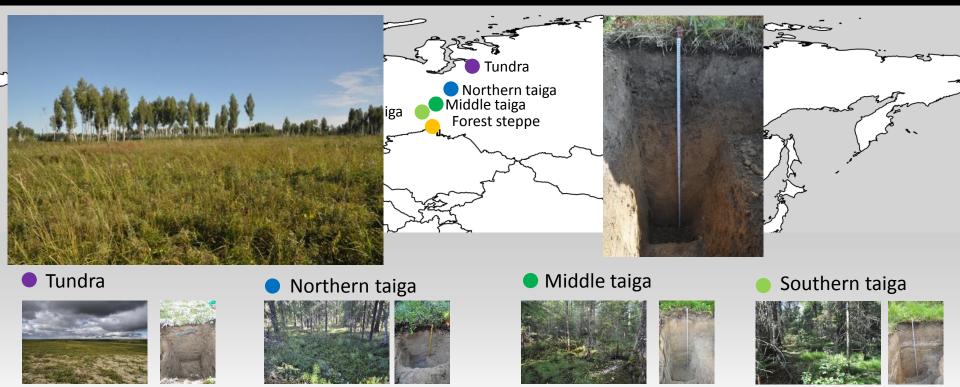
Northern taiga



Southern taiga



Forest steppe



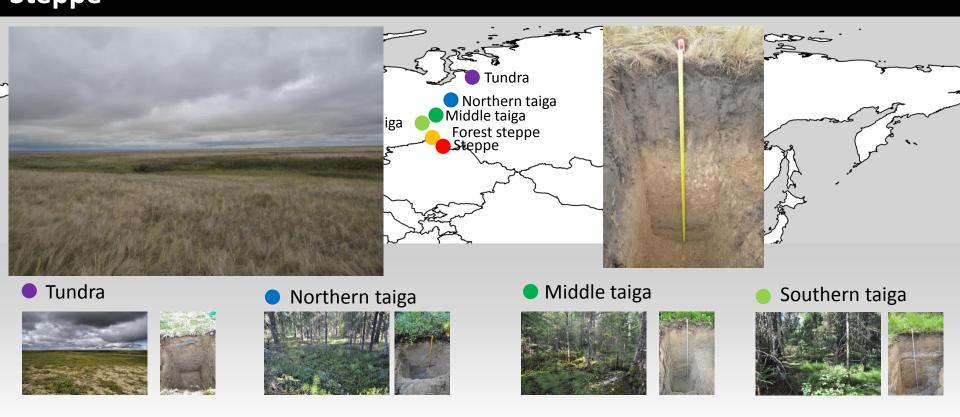
Forest steppe







Steppe



Forest steppe -meadow

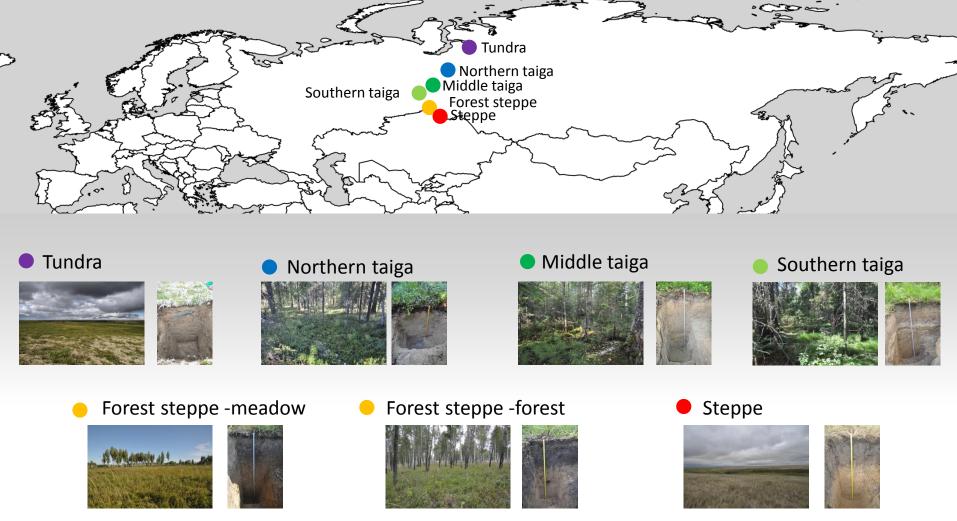




Forest steppe -forest



Sites

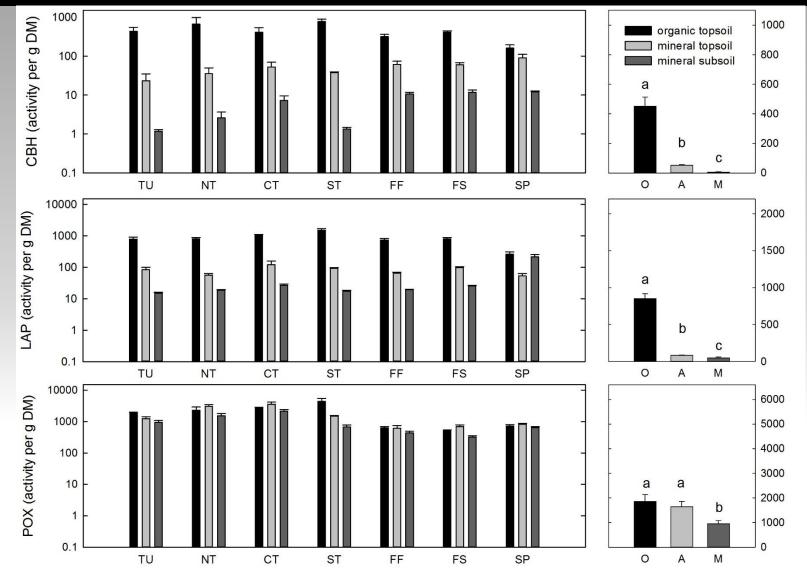


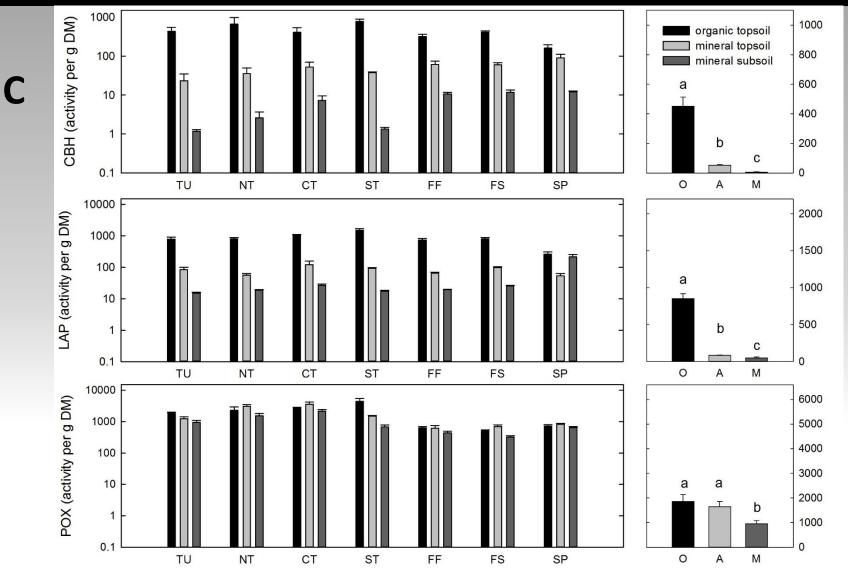


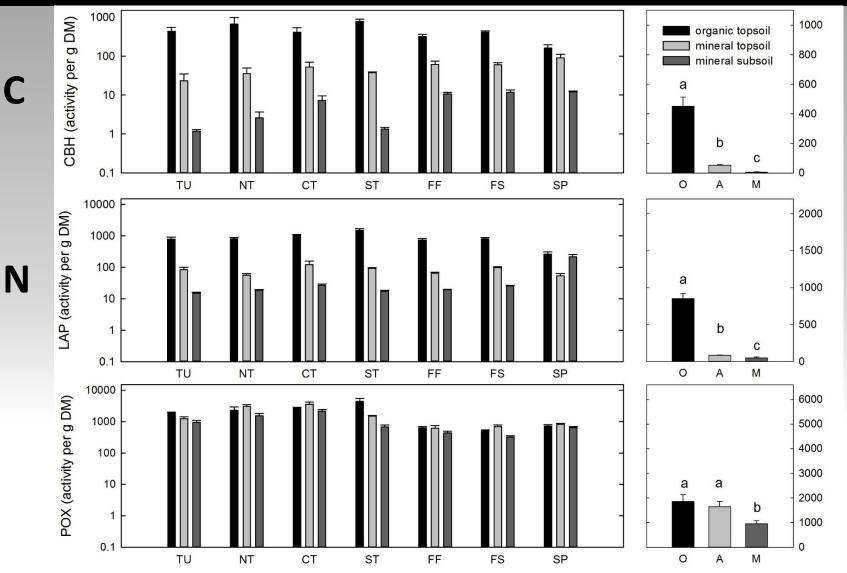


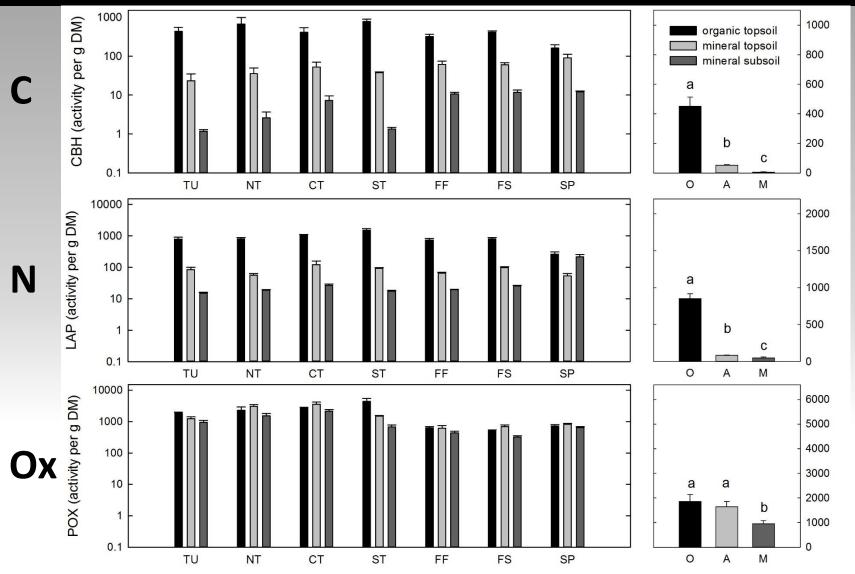




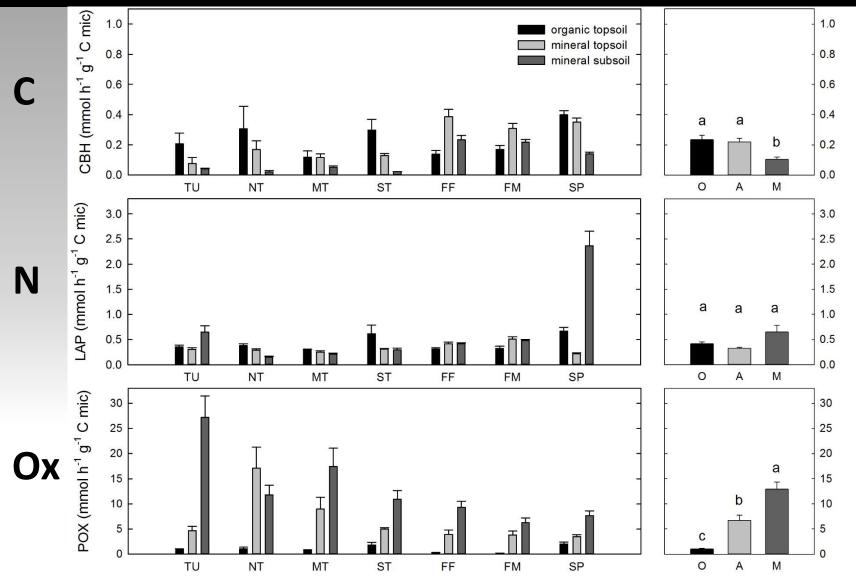


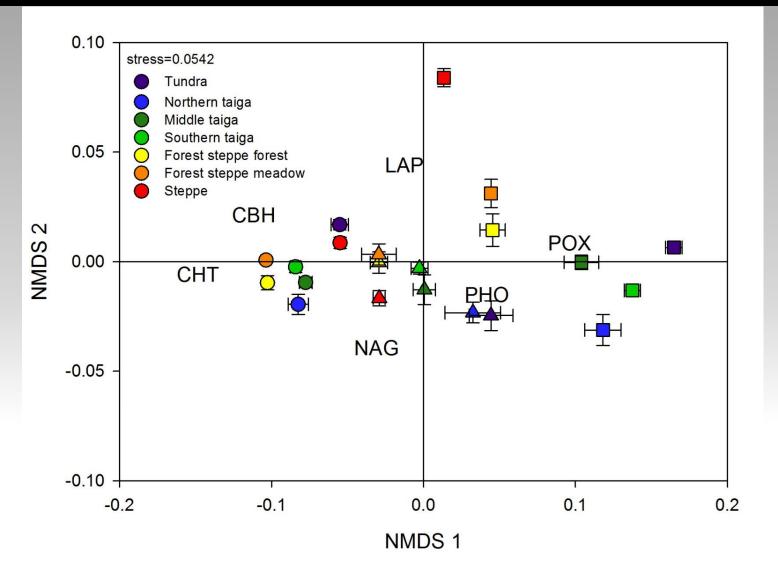


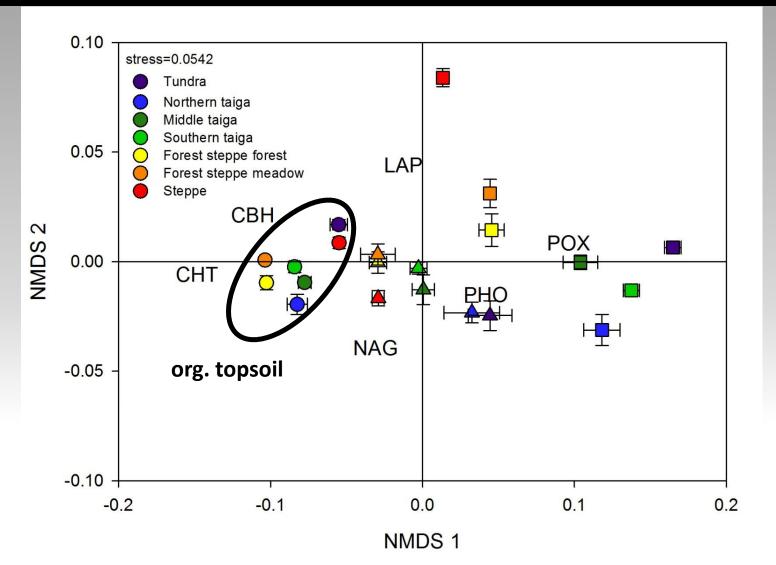


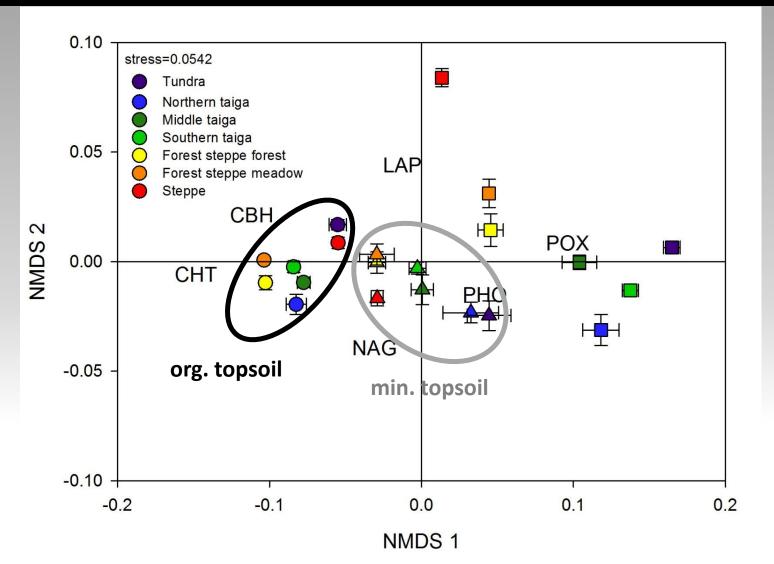


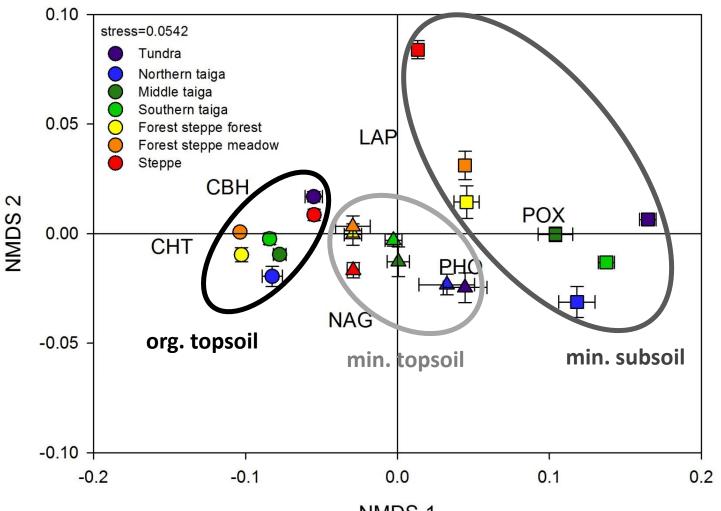
Enzyme activities per g mic C





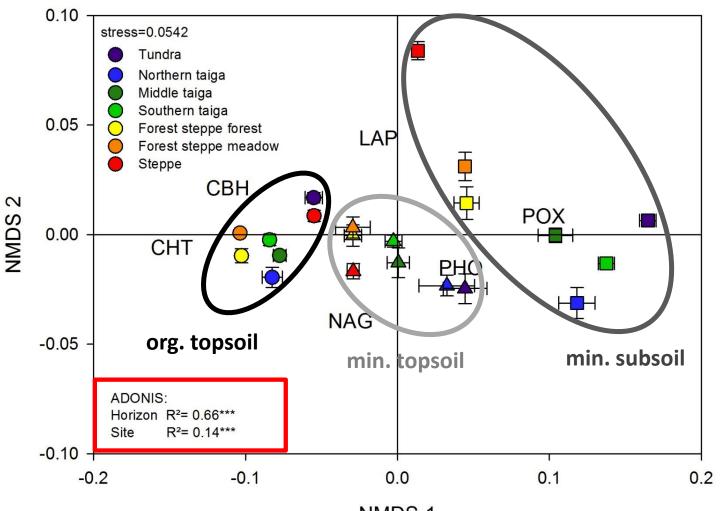






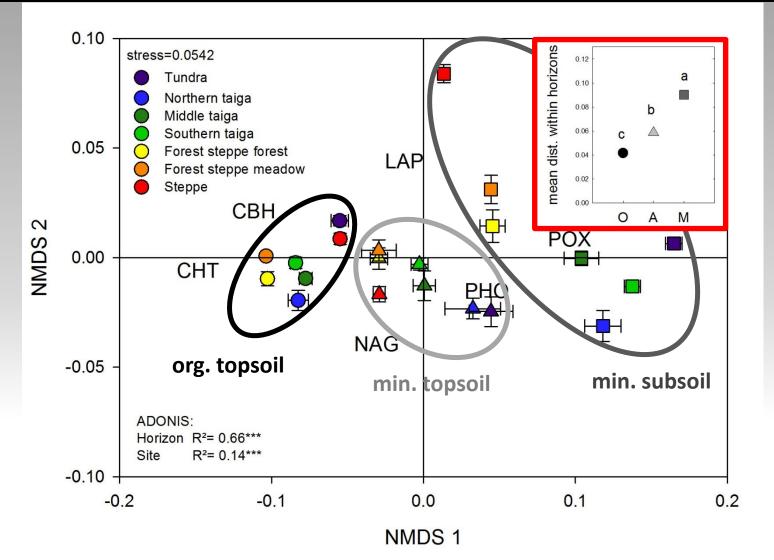
NMDS 1

Enzyme patterns

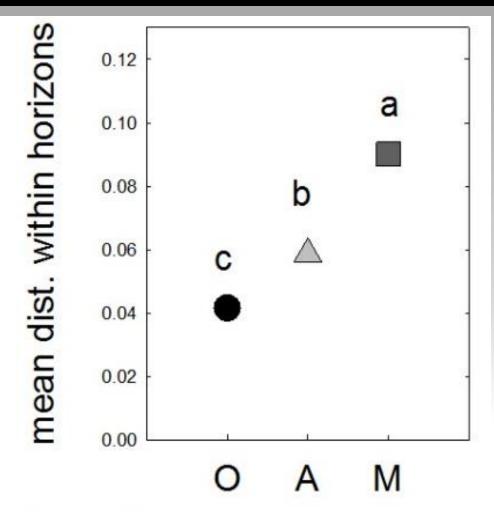


NMDS 1

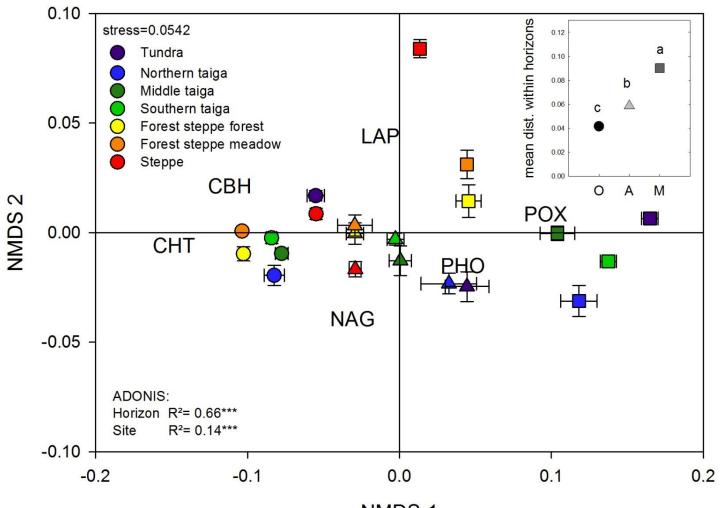
Enzyme patterns



Enzyme patterns



Enzyme patterns



NMDS 1

Mantel tests

soil C soil N soil CN microbial C microbial N microbial CN pН mic. comm. comp. fungi:bacteria

	organic
	topsoil
soil C	0.06
soil N	0.22
soil CN	0.05
microbial C	
microbial N	0.03
microbial CN	
рН	0.02
mic. comm. comp.	0.16
fungi:bacteria	0.10

	organic	mineral
_	topsoil	topsoil
soil C	0.06	
soil N	0.22	
soil CN	0.05	0.04
microbial C		
microbial N	0.03	
microbial CN		0.05
рН	0.02	0.03
mic. comm. comp.	0.16	0.04
fungi:bacteria	0.10	

	organic	mineral	mineral
	topsoil	topsoil	subsoil
soil C	0.06		
soil N	0.22		
soil CN	0.05	0.04	
microbial C			
microbial N	0.03		0.02
microbial CN		0.05	
рН	0.02	0.03	0.29
mic. comm. comp.	0.16	0.04	0.17
fungi:bacteria	0.10		0.24

	organic	mineral	mineral
	topsoil	topsoil	subsoil
soil C	0.06		
soil N	0.22		
soil CN	0.05	0.04	
microbial C			
microbial N	0.03		0.02
microbial CN		0.05	
рН	0.02	0.03	0.29
mic. comm. comp.	0.16	0.04	0.17
fungi:bacteria	0.10		0.24

Summary



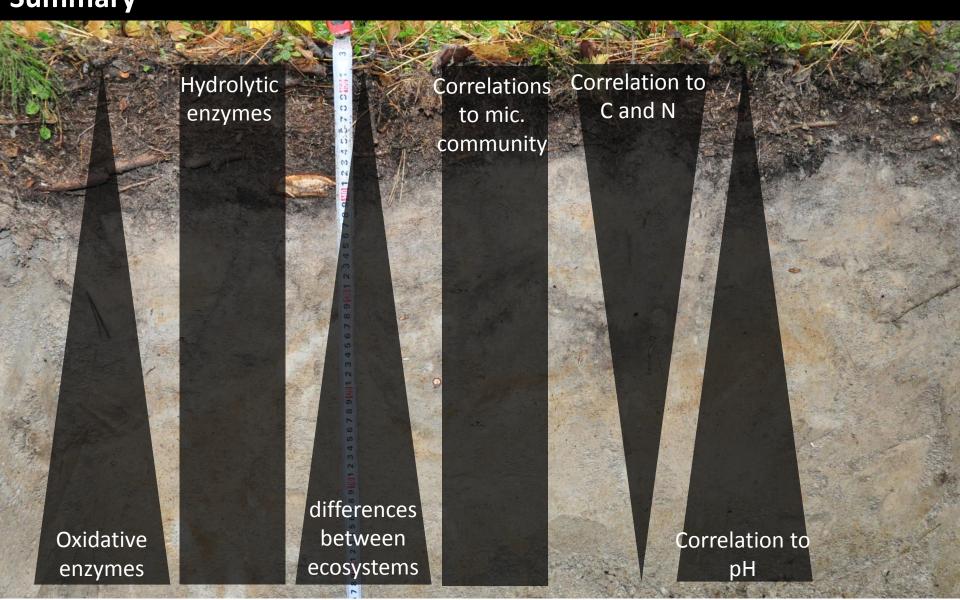
Summary

Hydrolytic enzymes differences Oxidative

enzymes

between ecosystems

Summary



Conclusions



Conclusions

Microbial community composition determines how soil organic matter is decomposed.



Conclusions

Microbial community composition determines how soil organic matter is decomposed.

In mineral subsoils C and N are weak predictors for how soil organic matter is decomposed.

Conclusions

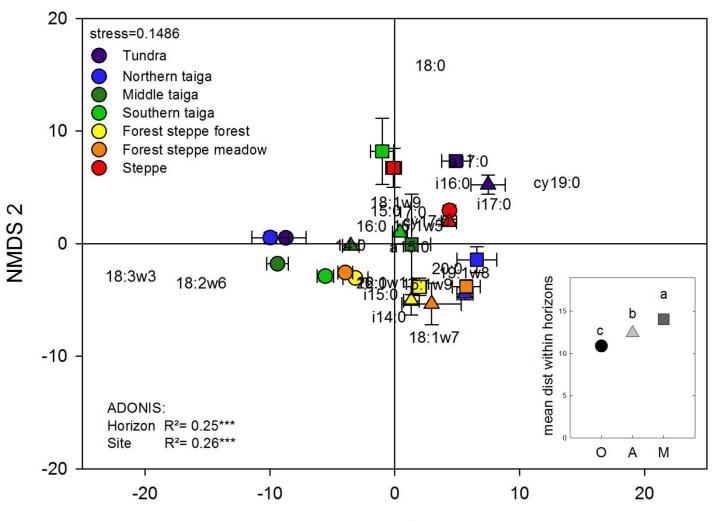
joerg.schnecker@univie.ac.at

Microbial community composition determines how soil organic matter is decomposed.

Thank you for your attention.

In mineral subsoils C and N are weak predictors for how soil organic matter is decomposed.

Microbial community composition



NMDS 1

Microbial community composition

