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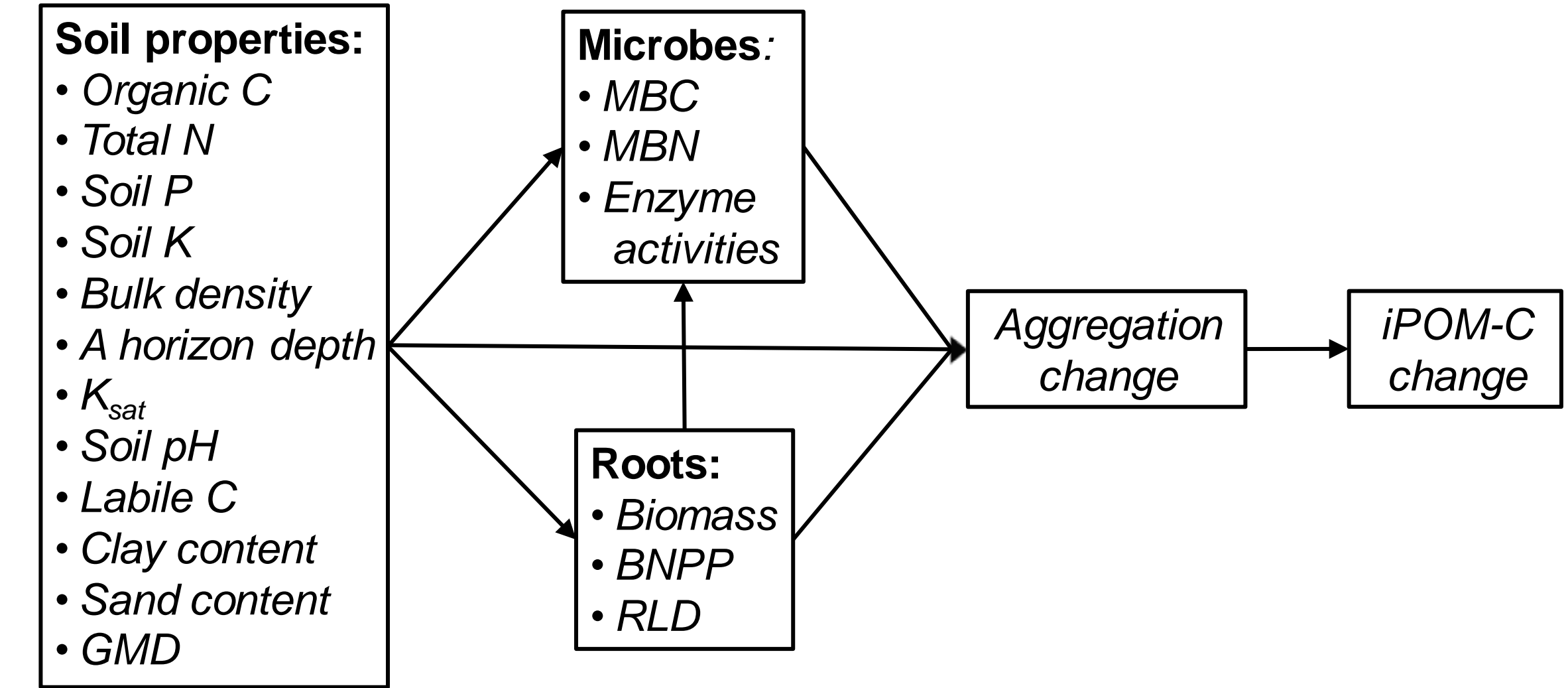
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Bioenergy: potential impacts on soil C storage

- Land use changes prompted by bioenergy may alter belowground carbon (C) dynamics leading to shifts in soil C storage.
- 'Marginal' sites not well-suited for annual row crops are targeted for bioenergy production with perennials because of the expected benefits from increased soil organic matter (SOM).
- Model predictions of soil C responses to bioenergy crops are dependent on understanding ecosystem drivers of C cycling processes operating at multiple scales.
- Short-term accumulation of soil C occurs primarily through the physical protection of organic matter resulting from increased soil aggregation.
- We investigated the importance of multiple environmental drivers of short-term (3 years) changes in physically protected C across a heterogeneous agroecosystem—containing both highly productive and 'marginal' soils—using structural equation modeling.

Our *a priori* model included...

- 11 soil chemical, physical & biological properties (Ontl et al. 2013).
- Three microbial community variables (Hargreaves & Hofmockel 2014).
- Three variables describing root characteristics (Ontl et al. 2013).
- Changes in soil aggregation (Ontl et al. *In review*).
- Changes in iPOM-C (Ontl et al. *In review*).

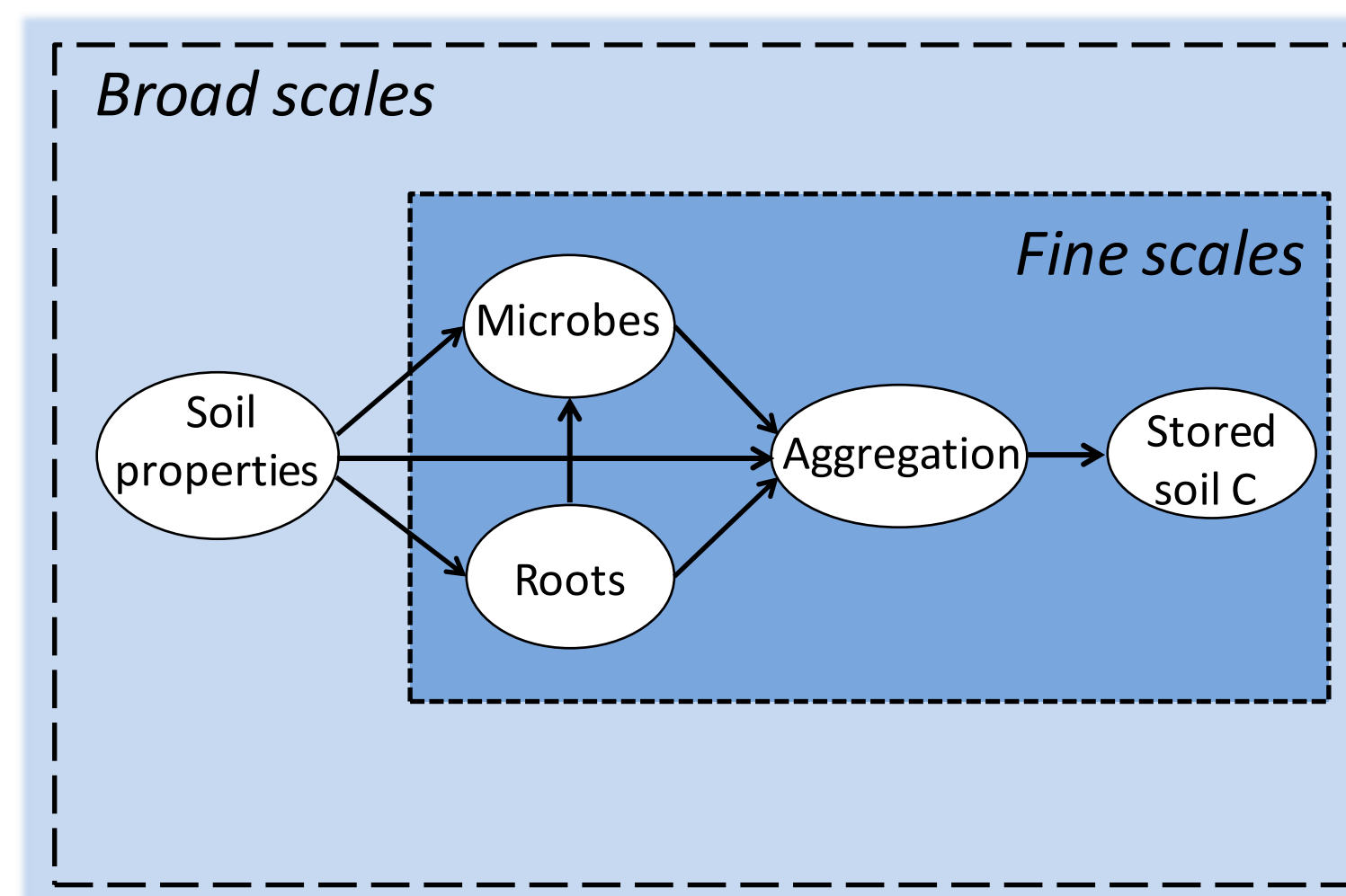


Fine- & broad-scale processes drive formation of iPOM-C

Soil aggregation dynamics drive C storage: aggregates trap **particulate organic matter (iPOM)**, physically protecting C within microsites & reducing decomposition.

At **broad scales**, soil physical, chemical & biological properties vary.

Both fine- and broad-scale drivers influence iPOM, **but the relative influence of each are not well understood.**



Fine-scale rhizosphere processes (root-microbe interactions, microbial biomass) influence aggregation & iPOM-C.

Enzymes break down SOM, **but the role of microbial enzyme activities on iPOM-C formation processes are unknown.**

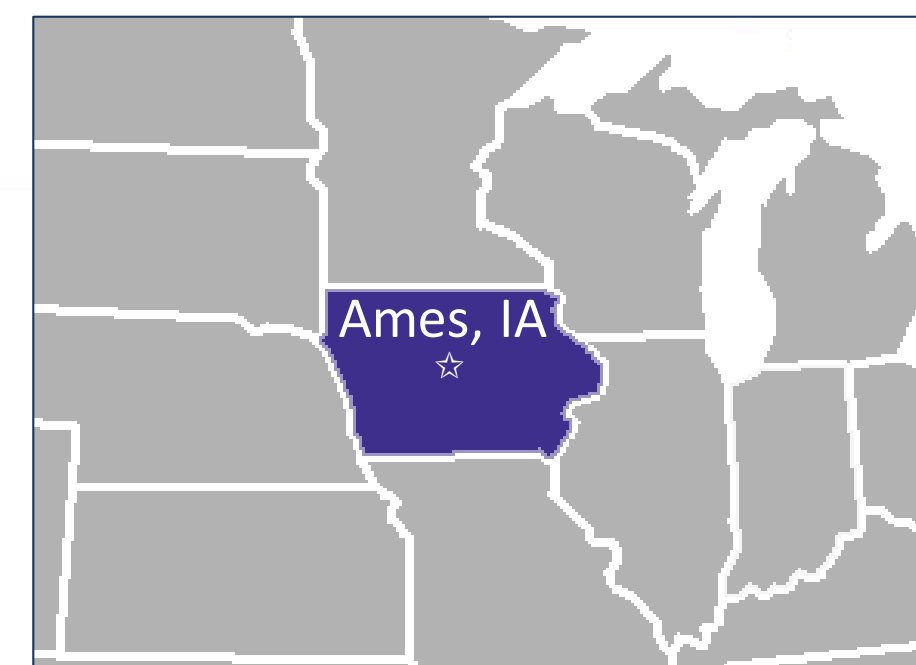
We tested two hypotheses:

H₁: Soil variation at broad scales (e.g. nutrients, texture) have **direct** impacts on aggregation, and **indirect** impacts through fine-scale processes.

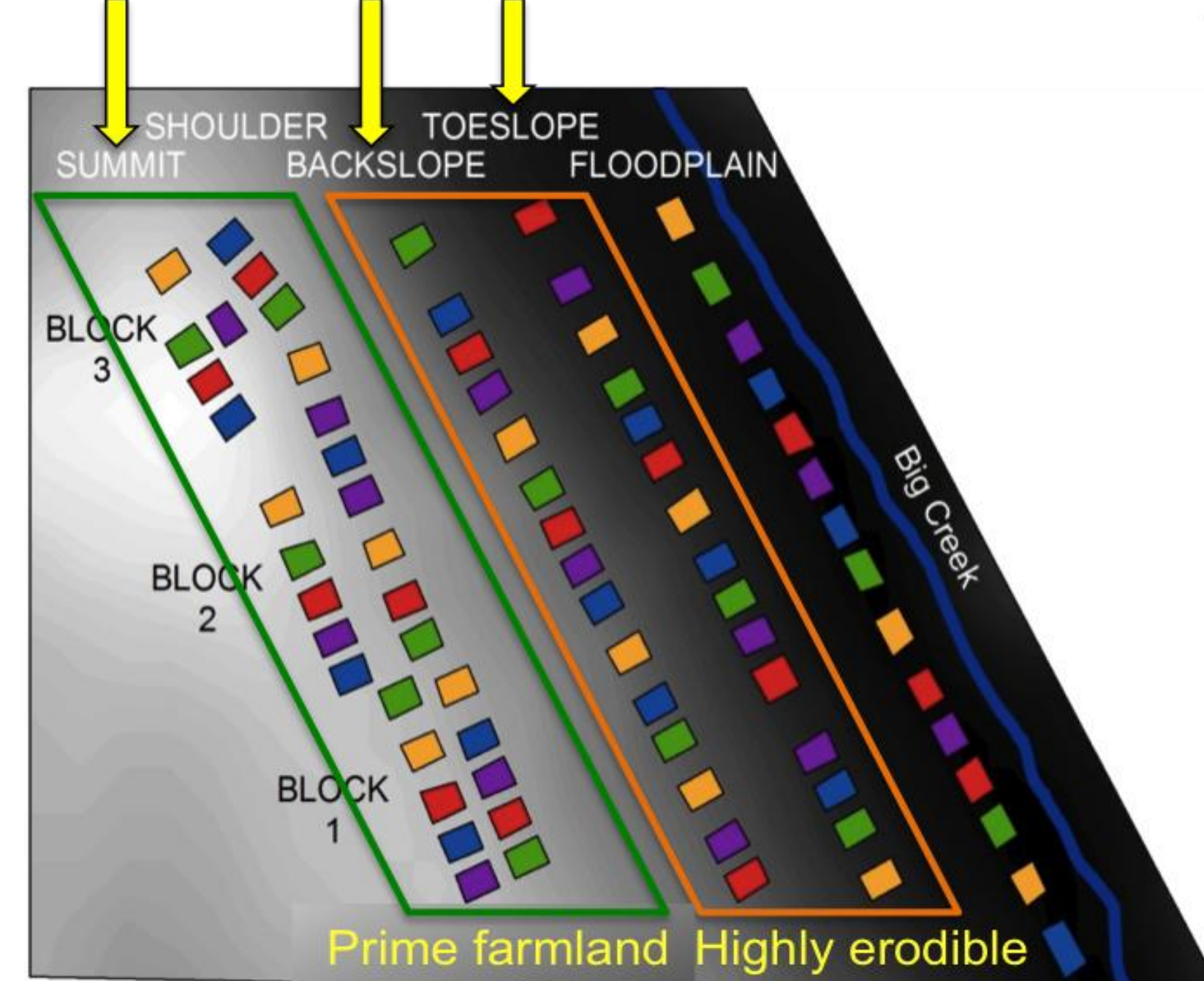
H₂: Microbial biomass and extracellular enzyme activities positively impact iPOM-C levels through effects on aggregate formation.

The Landscape Biomass Project

- Data were collected from 27 plots (0.06 ha in size) from 2009-2012
- Located on three positions sited across a **topographic gradient**
- Three bioenergy cropping systems were included



Three landscape positions



Elevation
325 meters
305 meters

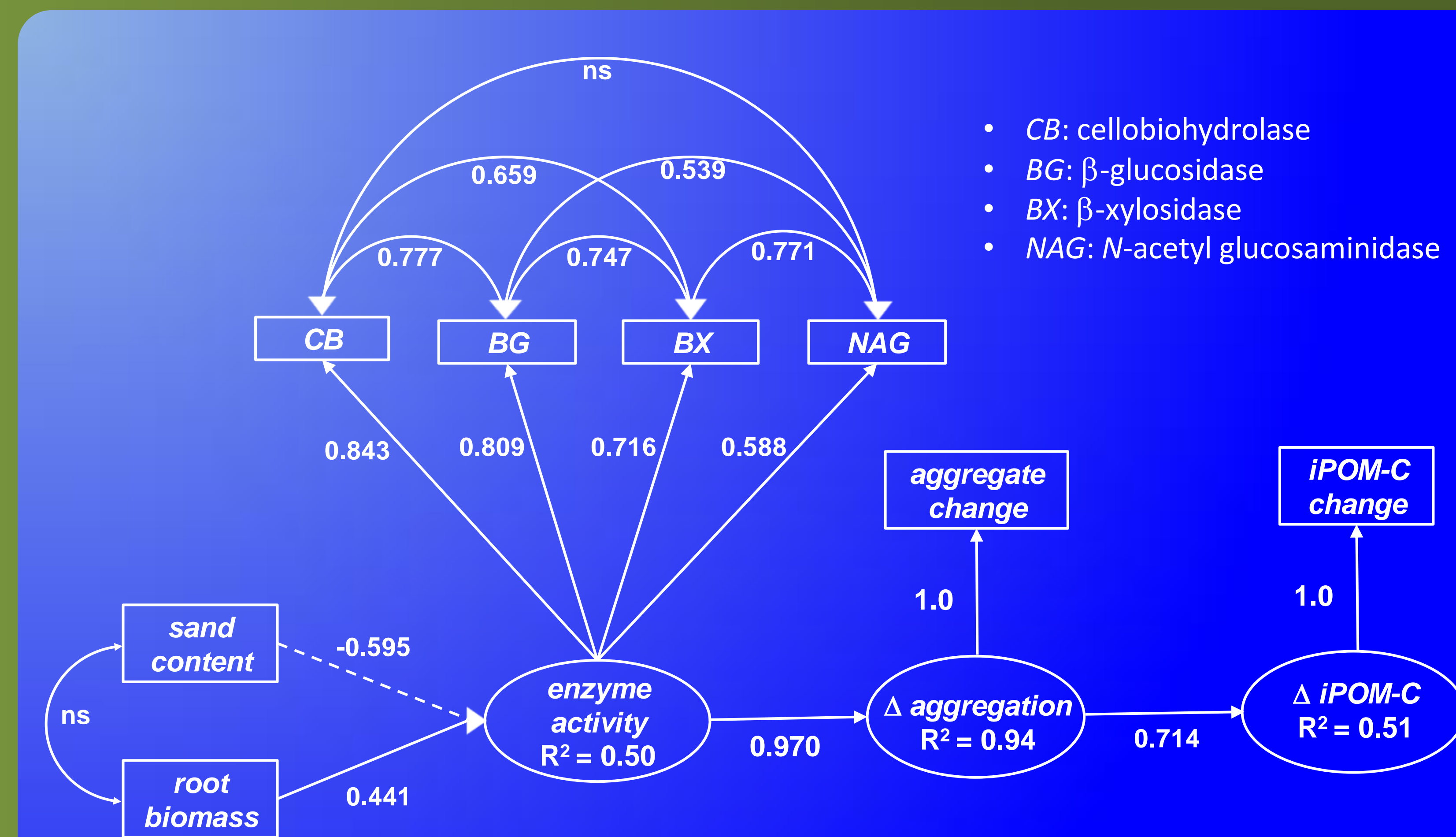
Cropping system

- Continuous corn
- Modified rotation
- Switchgrass
- Triticale/sorghum
- Aspen/triticale

0 50 100 200 300 Meters

Structural Equation Modeling Results

- H₁:** Broad-scale drivers affected C storage, but only **indirectly** through influences on fine-scale root-rhizosphere processes: • Sand content negatively influenced potential enzyme activities. • Land use had significant effects on root biomass (data not shown, Ontl et al. 2013a).
- H₂:** Fine-scale drivers strongly affect iPOM-C changes: • Root biomass positively influenced enzyme activities. • Enzyme activities were strongly linked to aggregation change. • Aggregation changes positively influenced shifts in iPOM-C.



- CB: cellobiohydrolase
- BG: β-glucosidase
- BX: β-xylosidase
- NAG: N-acetyl glucosaminidase

Model goodness-of-fit measures

χ^2	15.3	CFI ^c	0.99
Df ^a	13	RMSEA ^d	0.08
P-value ^b	0.291	SRMR ^e	0.047

Criteria

^a Model degrees of freedom	>0
^b χ^2 P-value	>0.05
^c Comparative Fit Index	>0.90
^d Root Mean Square Error of Approximation	<0.08
^e Standardized Root Mean Square Residual	<0.08

Straight arrows indicate causal paths between variables; curved arrows indicate covariance. Dashed arrows signify negative relationships. Standardized parameter estimates are shown; parameter estimates are significant unless otherwise indicated (ns). Numbers inside circles represent variance explained by the model (R²).

Pathway	Unstandardized Estimate	SE	Critical ratio	P-value	Standardized Estimate
LATENT VARIABLES					
BG → enz	1.000	0	1	0	0.809
CB → enz	1.746	0.160	10.884	<0.001	0.843
BX → enz	1.801	0.231	7.778	<0.001	0.716
NAG → enz	0.591	0.136	4.362	<0.001	0.588
REGRESSIONS					
stcrop → enz	0.275	0.101	2.716	0.007	-0.595
sand → enz	-0.625	0.185	-3.368	0.001	0.441
enz → agg	0.137	0.027	5.085	<0.001	0.970
agg → ipom	5.192	0.979	5.302	<0.001	0.714

References:

- Hargreaves SK & Hofmockel KS. 2014. Physiological shifts in the microbial community drive changes in enzyme activity in a perennial agroecosystem. *Biogeochemistry* 117: 67–79.
- Ontl TA, Hofmockel KS, Cambardella CA, Schulte LA, Kolka RK. 2013a. Topographic and soil influences on root productivity of three bioenergy cropping systems. *New Phytologist* 199: 727–737.
- Ontl TA, Cambardella CA, Schulte LA, Kolka RK. *In review*. Soil aggregation and carbon pools respond quickly to changes in land use and tillage across a heterogeneous agroecosystem. *Soil Science Society of America Journal*.