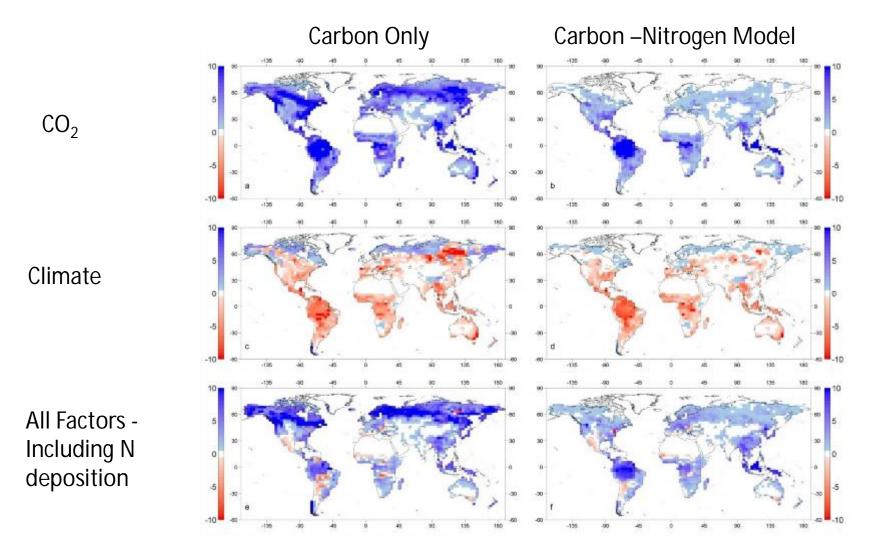
Representation of Soil Carbon Dynamics and Nitrogen Limitation of Decomposition in Terrestrial Ecosystem Models: Present and Future Directions

> Wilfred Post, Peter Thornton, Gangsheng Wang Oak Ridge National Laboratory Oak Ridge, TN USA

#### Do Climate Models Need Soil C-N Dynamics?

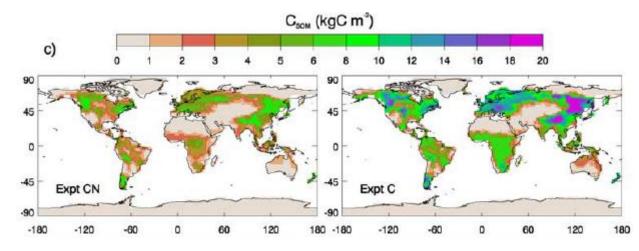
- Latent heat simulation requires stomatal conductance calculation – depends, in part on leaf N concentration.
- Carbon stocks in soil show sensitivity to inclusion of N dynamics.
- Models that include N dynamic in soil C decomposition show a tendency for reduced heterotrophic respiration.

#### OCN Changes in C stocks (kg C m<sup>-2</sup>) 1860-2100



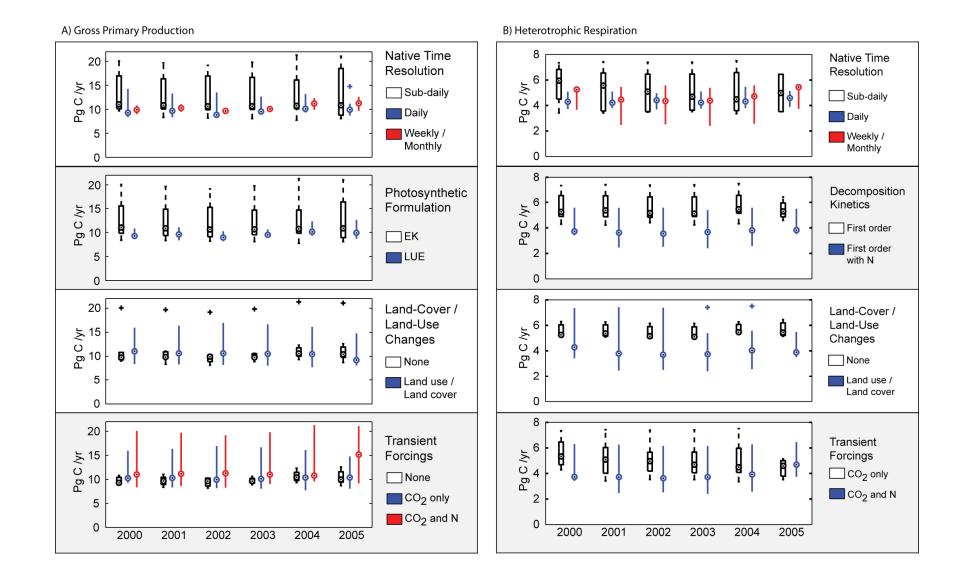
From Zaehle et al. 2010.

#### CLM-CN



Experiment	Wood C <sup>b</sup>	Veg. C	CWD C <sup>c</sup>	Litter C	SOM C <sup>d</sup>	Total C
			Control			
CN	613 (53)	653 (57)	147 (13)	16(1)	334 (29)	1150
C	943 (47)	1014 (50)	247 (12)	28 (1)	736 (36)	2026
Cr	712 (49)	771 (53)	167 (11)	19 (1)	496 (34)	1452
		Y	ears 1976–2000			
CN+co2	649 (54)	690 (58)	153 (13)	16 (1)	339 (28)	1199
CN+ndep	619 (53)	660 (57)	149 (13)	16 (1)	339 (29)	1163
CN+co2ndep	656 (54)	698 (57)	155 (13)	17 (1)	344 (28)	1213
C+co2	1047 (48)	1125 (51)	269 (12)	31 (1)	776 (35)	2201
Cr+co2	803 (50)	870 (54)	184 (11)	22 (1)	537 (33)	1612
		Y	lears 2076–2100			
CN+co2	801 (57)	845 (60)	176 (13)	18 (1)	357 (26)	1397
CN+ndep	642 (53)	684 (57)	154 (13)	17 (1)	352 (29)	1206
CN+co2ndep	847 (57)	895 (60)	186 (13)	19 (1)	379 (26)	1480
C+co2	1527 (52)	1625 (55)	363 (12)	39 (1)	936 (32)	2963
Cr+co2	1225 (54)	1309 (57)	263 (12)	28 (1)	683 (30)	2284

#### **NACP Regional Synthesis**



#### Climate Models Need Accurate Representations of Soil C-N Dynamics

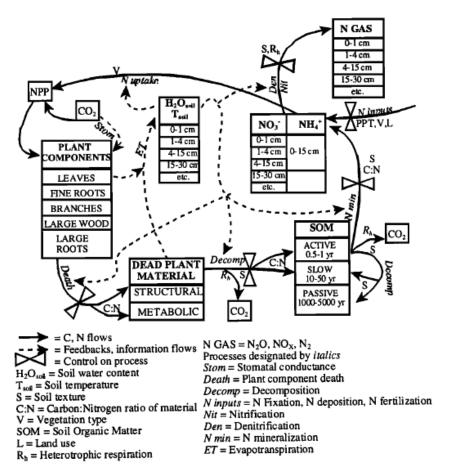
- Soil C decomposition representations have not principally changed in 30 years.
- While soil models differ in many details,
  - N dynamics generally follow C dynamics,
  - N becomes available as a result of stoichiometry constraints.
- All climate models, to my knowledge, used at global scale use first order "donor control" kinetics.

# Jenny/Olson (1963) Paradigm dC / dt = I - kC

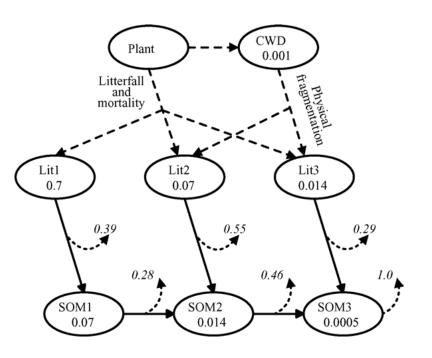
- CENTURY/DAYCENT in OCN, CASA-CNP, CASA-GFED, LM3V, many others at regional scales
- RothC/SUNDIAL in ISAM,
- Biome-BGC in CLM-CN
- In each model the associated N turnover is determined by C dynamics among various C pools and stoichiometry.

#### **Decomposition Model Structures**

#### **CENTURY/DAYCENT**

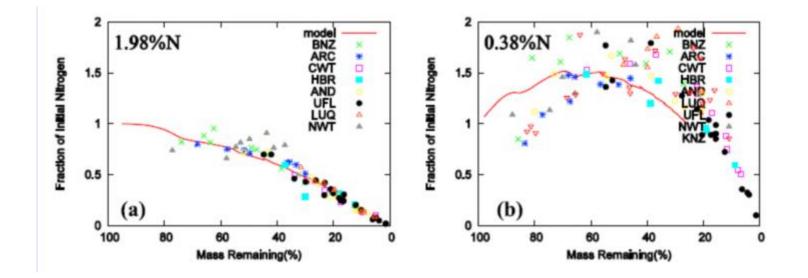


CLM-CN (after Biome-BGC)



Converging cascade model of litter and soil organic matter decomposition. The model includes three litter pools (Lit1, Lit2, and Lit3, see text) and three soil organic matter pools (SOM1, SOM2, and SOM3).

#### ISAM



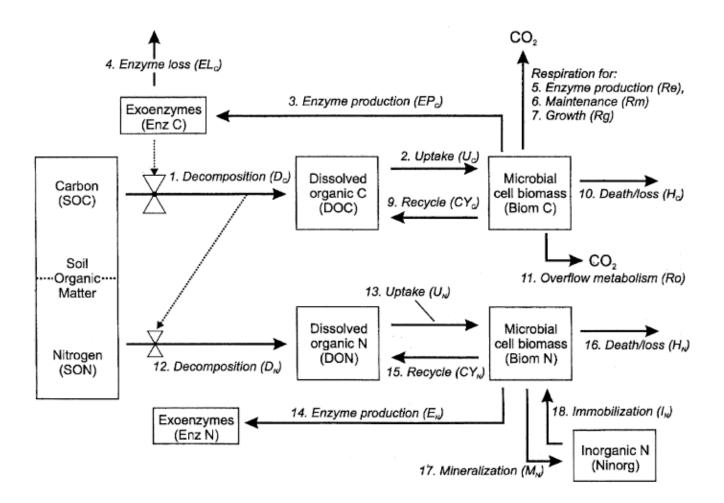
Fraction of initial litter N remaining as a function of the leaf litter mass remaining, with (a) high N litter with 1.98% N and (b) low N litter with 0.38% N. The lines in the graphs are model simulations averaged across all LIDET sites involved.

(Yang et al. 2009)

## Why Do We Need to Consider a New Approach?

- Substrate supply models are immensely useful:
  - Capture major components of observed soil C spatial patterns
  - Kinetics generally consistent with long-term experimental finding
- However, there are important phenomena that are not directly addressed:
  - Priming (Wutzler and Reichstein 2007)
  - Divergent results from N fertilization experiment
  - Mineral soil interactions (Heal et al. in press)
  - Complicated temperature responses (Davidson and Janssens 2006)

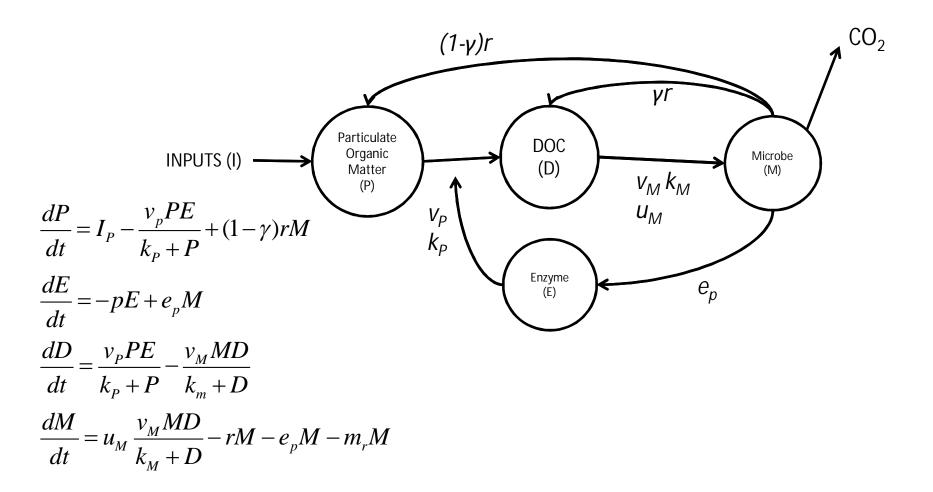
#### Decomposition is an Enzyme Kinetic Process

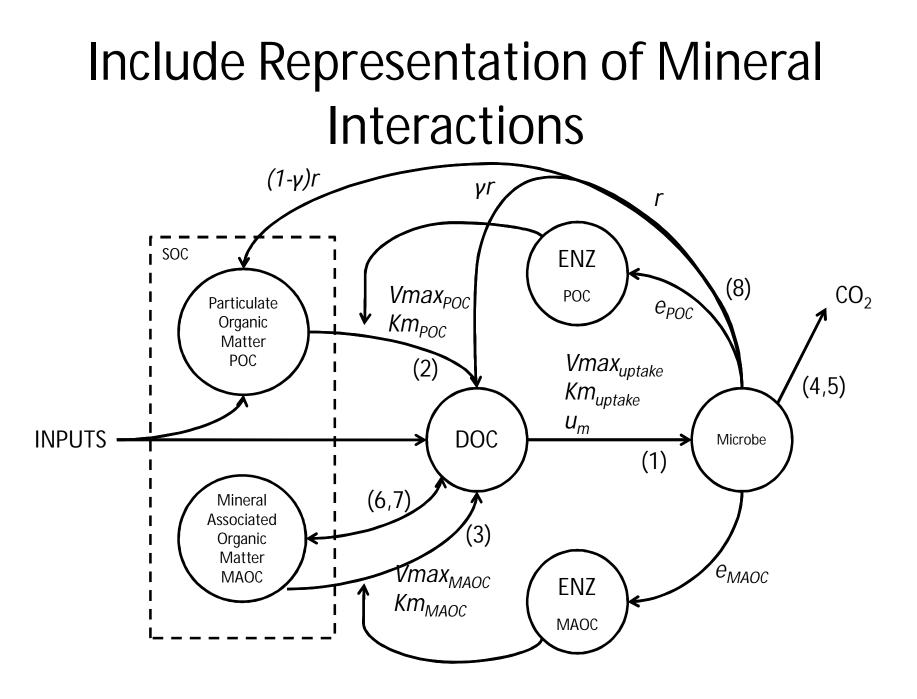


from Schimel and Weintraub (1993)

#### **Carbon Equations**

After Allison et al. (2010)



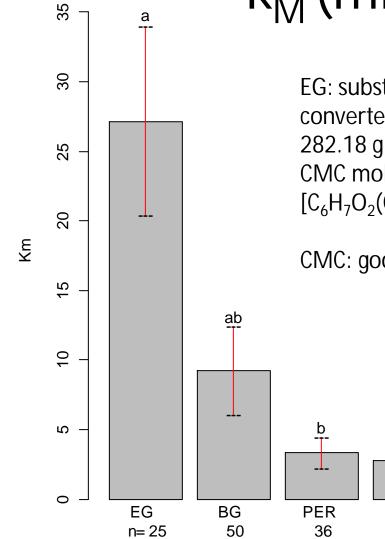


#### Soil Enzymes

Process	Enzyme	Abbr.	EC	Substrate	
Degradation	Laccase (phenol oxidase)	POX	1.10.3.2	L-3,4-	
of				dihydroxyphenylalanine	
Lignin				(DOPA)	
	Peroxidase	PER	1.11.1.7	$DOPA + H_2O_2$	
Degradation	β-1,4-glucosidase;	BG	3.2.1.21	4-methylumbelliferone	
of	Cellobiase			(MUB)-β-D-glucoside;	
Cellulose				p-nitrophenyl (pNP)-	
				glucopyranoside	
	Cellobiohydrolase ;	CBH	3.2.1.91	4-MUB-β-D-cellobioside;	
	Exoglucanase;			pNP-β-D-cellobioside	
	Cellulose 1,4-β-				
	cellobiosidase				
	Endo-glucanase	EG	3.2.1.4	Carboxymethyl cellulose	
	Cellulase			(CMC)	
	Carboxymethylcellulase				
N acquisition	N-acetyl-β-D-	NAG	3.2.1.14	4-MUB-N-acetyl-β-D-	
	glucosaminidase;		3.2.1.52	glucosaminide;	
	β-1,4-N-			pNP-β-N-	
	acetylglucosaminidase			acetylglucosaminide	

72 papers 181 observations

### $k_{M}$ (mM)



EG: substrates are CMC (mg mL<sup>-1</sup>), which is converted to mM by a molecular weight of 282.18 g mol<sup>-1</sup>. [*Ichikawa et al.*, 2005] CMC molecular formula  $[C_6H_7O_2(OH)_x(OCH_2COONa)_y]_n$ 

CMC: good/suitable substrate for EG

b

1

NAG

10

b

POX 12

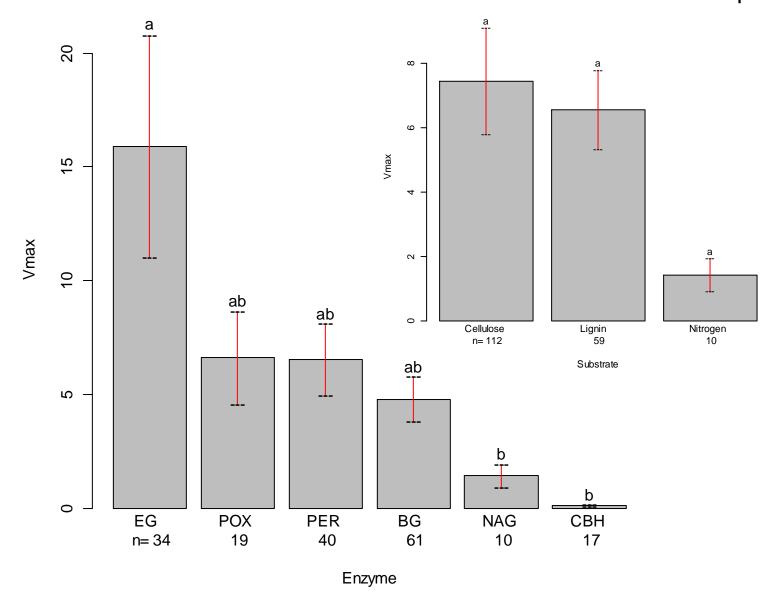


b

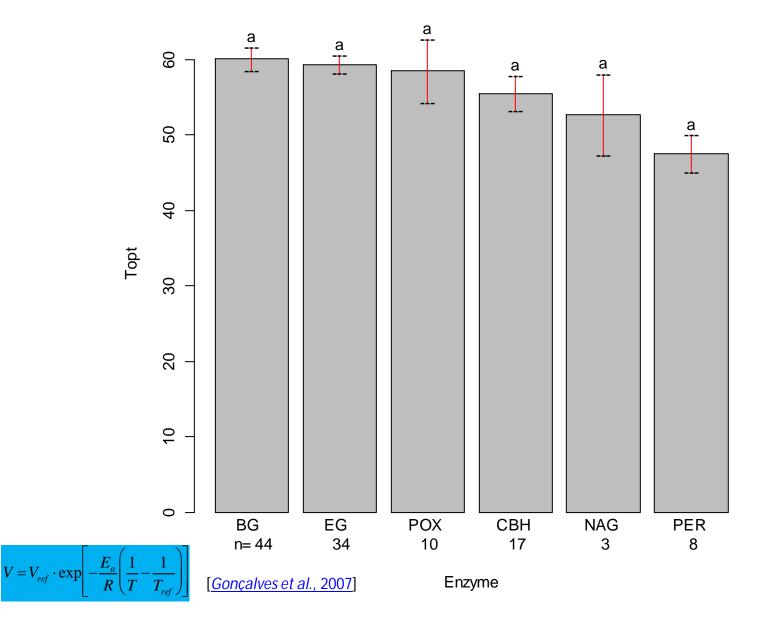
CBH

17

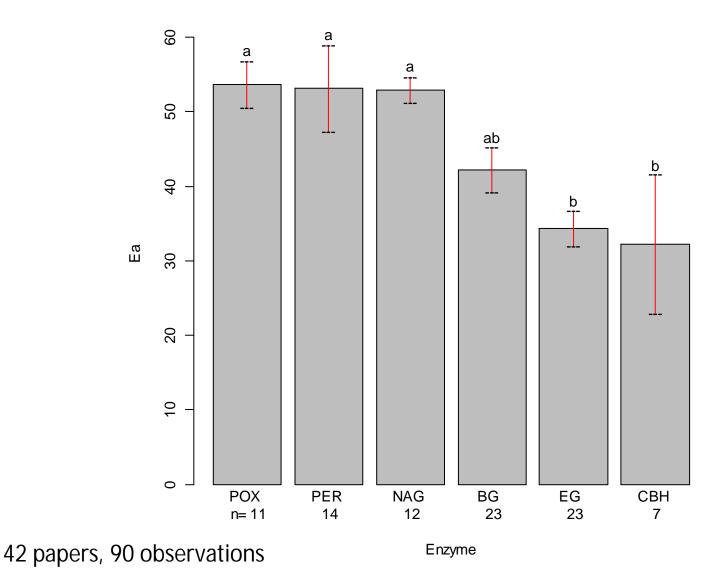
#### v<sub>m</sub> (mmol Enzyme mg<sup>-1</sup> soil h<sup>-1</sup>) at 20 °C & pH<sub>opt</sub>



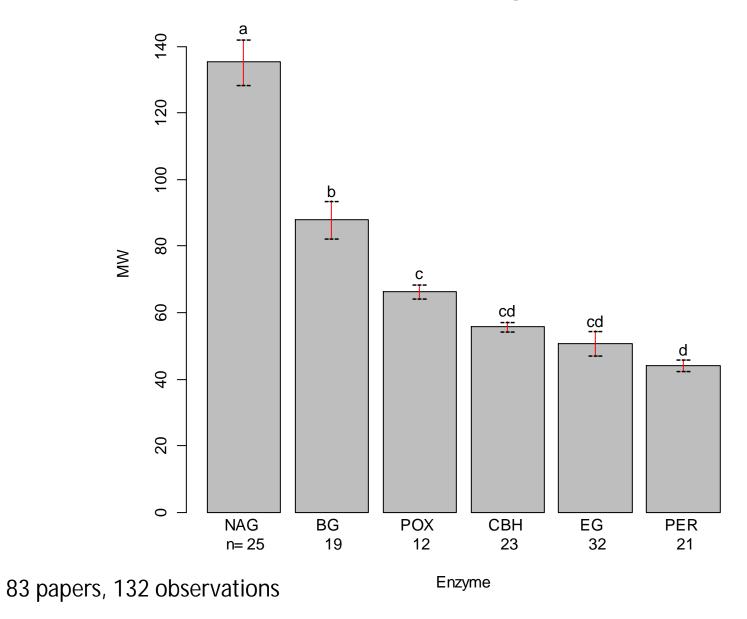
#### Optimum Temperature (°C)

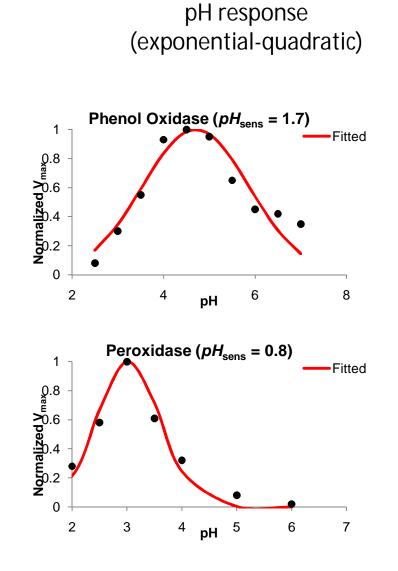


#### Activation Energy (kJ mol<sup>-1</sup>)

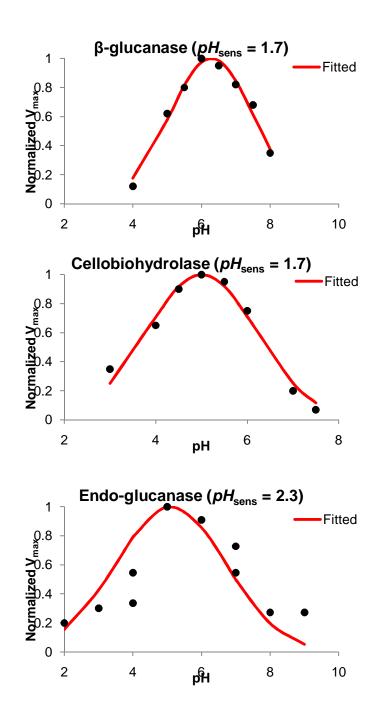


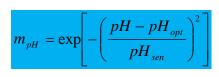
#### Molecular Weight (kDa)









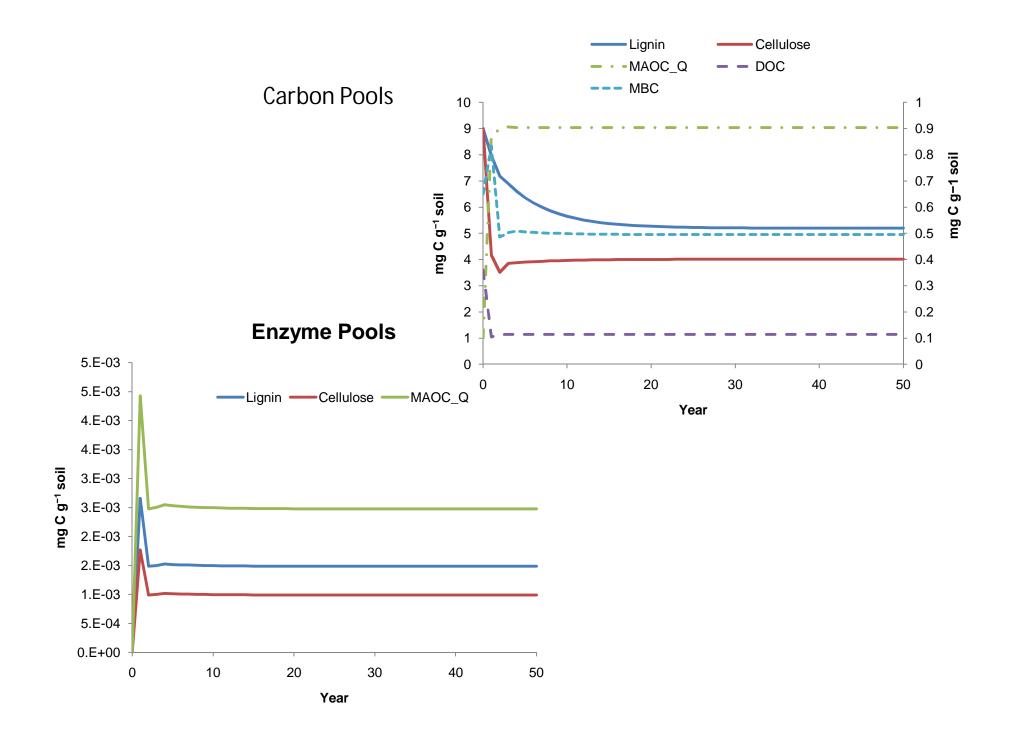


#### **Example Simulation**

Spin-up model (50 years)

- Soil: Alfisols (soc=28, Doc=0.36, MBC=0.65 mg C g<sup>-1</sup>)
- T = 20°C
- pH = 6

	Lignin	Cellulose	DOC
External input (×10 <sup>-4</sup> mg C g <sup>-1</sup> h <sup>-1</sup> )	1	5	5
V <sub>max</sub> (mg C mg <sup>-1</sup> ENZ h <sup>-1</sup> )	1.54	39.02	
Km (mg C g <sup>-1</sup> )	100	300	



#### Next Steps

- Include N equations
- Reality using long-term experimental datasets
- Currently parameters are estimated using data largely from chemostat experiments
- Estimation of soil enzyme kinetic equation parameters
  - need to be estimated from a combination of field and laboratory experiments
  - using soils and soil carbon substrates

#### Summary

- C-N interaction effects on decomposition dynamics is currently imposed in models.
- New microbial enzyme kinetics approach shows promise of addressing various observed phenomena from first principles.
- Compromises are required to limit models to manageable complesity.
- New measurement approaches are needed to estimate suitable model parameters.
- Approach offers an opportunity utilize the new information becoming available from microbial genomics.

