

CLM5 Matrix Model

Computational efficiency, diagnostics, and Improvement with Data

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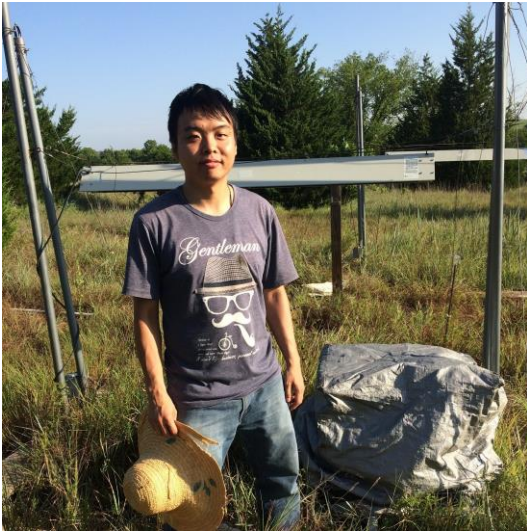
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Heroes



Chris Lu

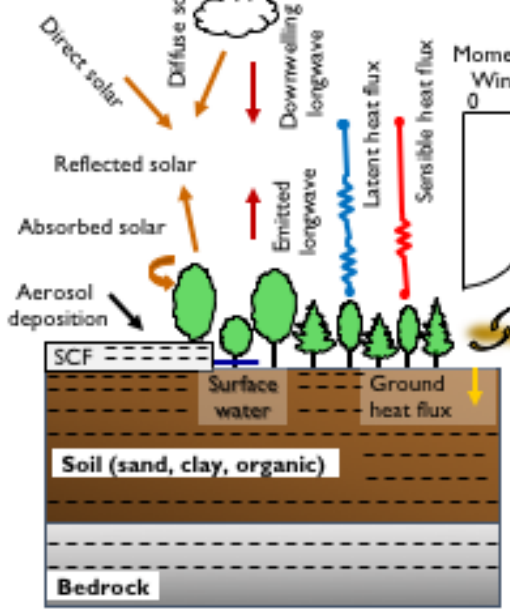


Zhenggang Du

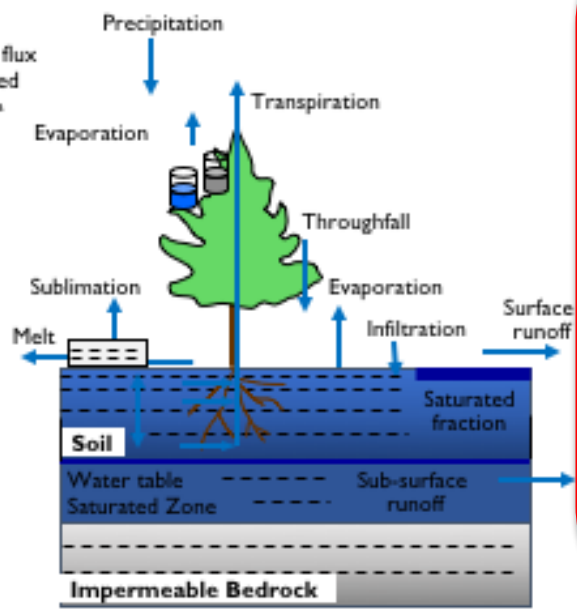


Feng Tao

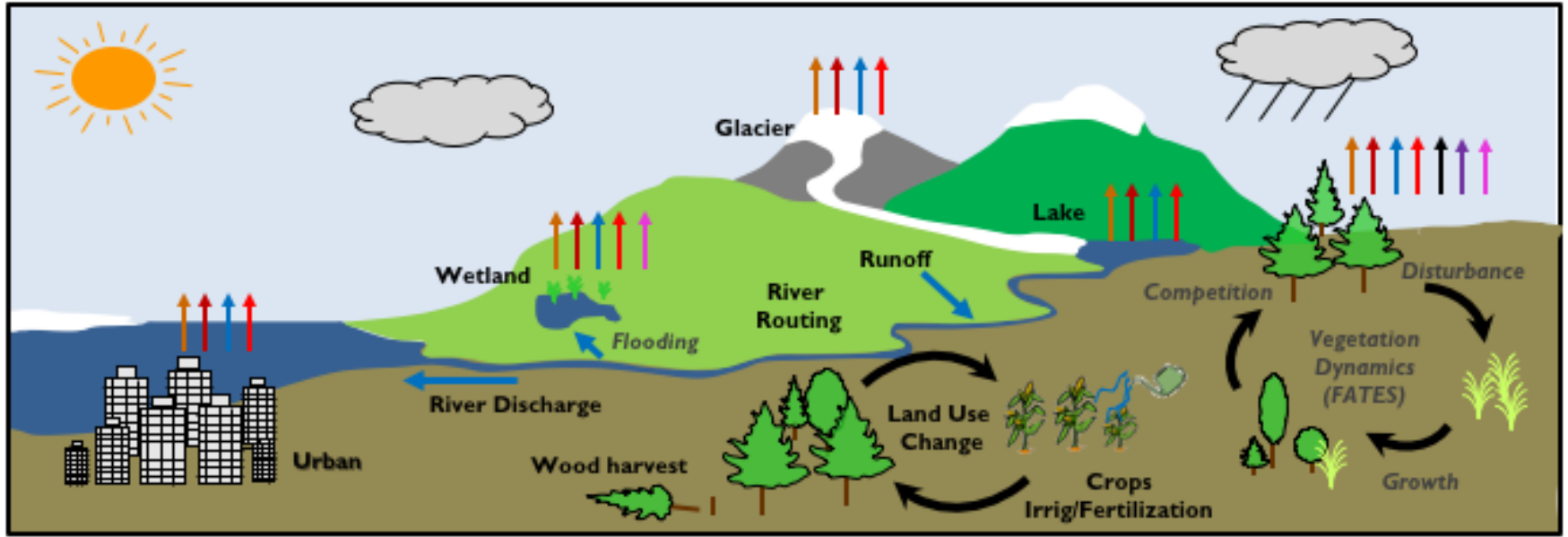
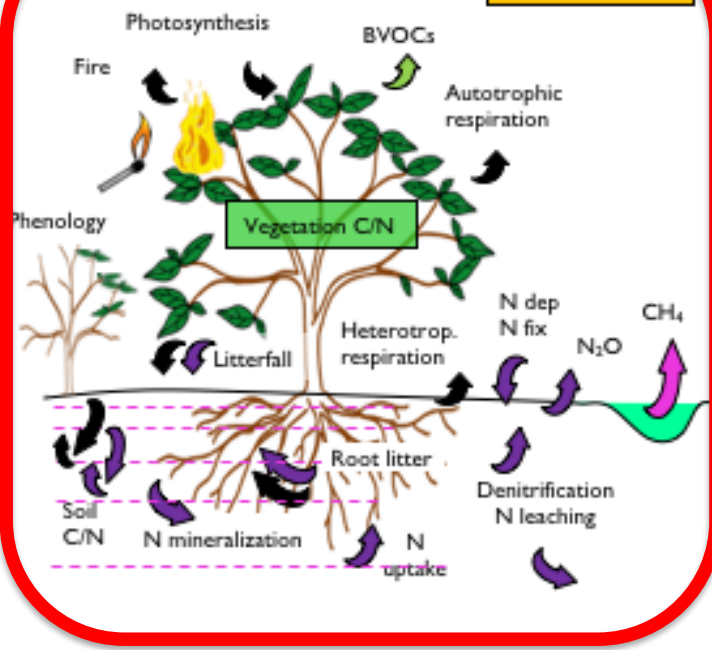
Surface energy fluxes



Hydrology

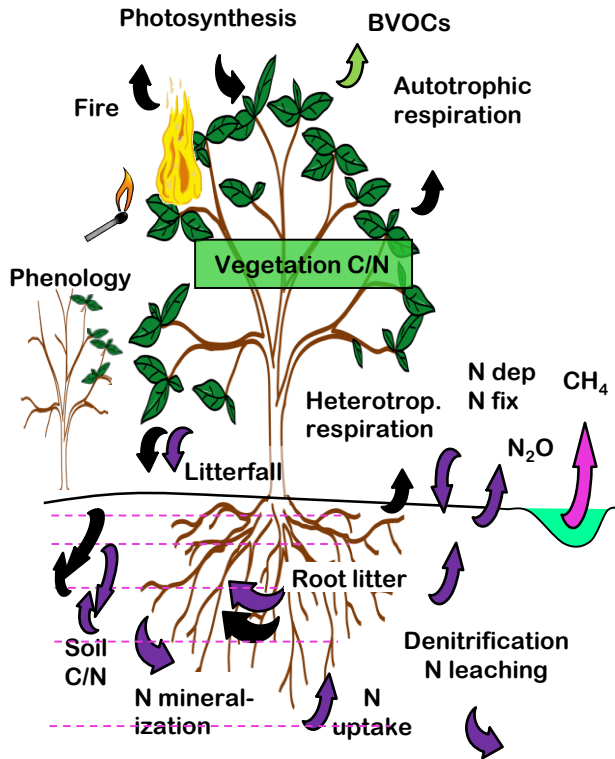


Biogeochemical cycles



CLM5.0 matrix model

Biogeochemical cycles



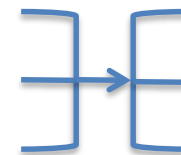
Carbon and nitrogen transfer among >892 pools
 446 pools for carbon cycle
 18 x 17 = 306 plant pools
 140 soil and litter pools over 20 layers
 446 pools for organic nitrogen
 plus Inorganic nitrogen pools

$$\frac{d}{dt} X(t) = (A_{ph}(t)K_{ph}(t) + A_{gm}(t)K_{gm}(t) + A_{fi}(t)K_{fi}(t))X(t) + B(t)F(t)$$

C transfer of phenology C transfer of gap mortality C transfer of fire pool state input
C turnover of phenology C turnover of gap mortality C turnover of fire allocation

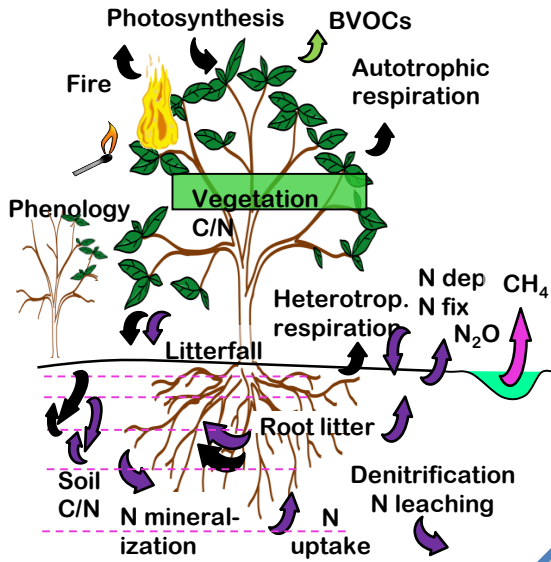
$$\frac{d}{dt} N(t) = (A_{Nph}(t)K_{Nph}(t) + A_{Ngm}(t)K_{Ngm}(t) + A_{Nfi}(t)K_{Nfi}(t))N(t) + B_N(t)F_N(t)$$

Simplicity in coding
 Diagnostic capability
 Computational efficiency

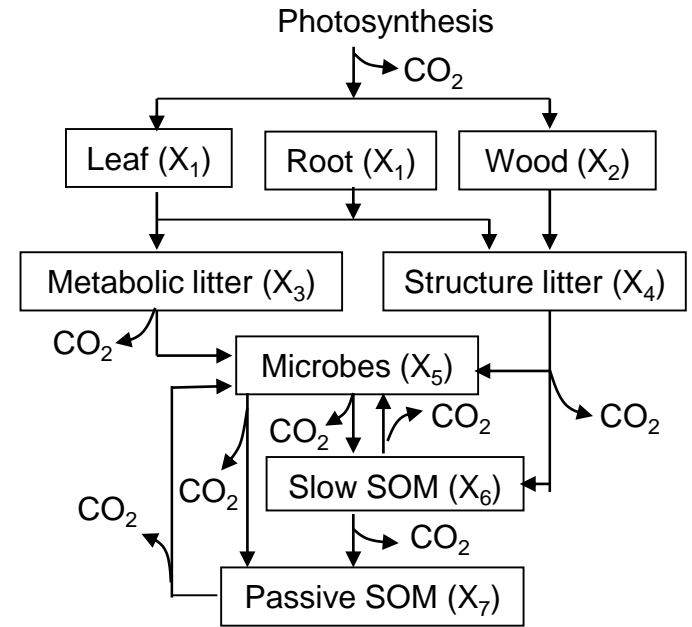


Model
 Understanding
 Evaluation
 Improvement

Matrix Approach



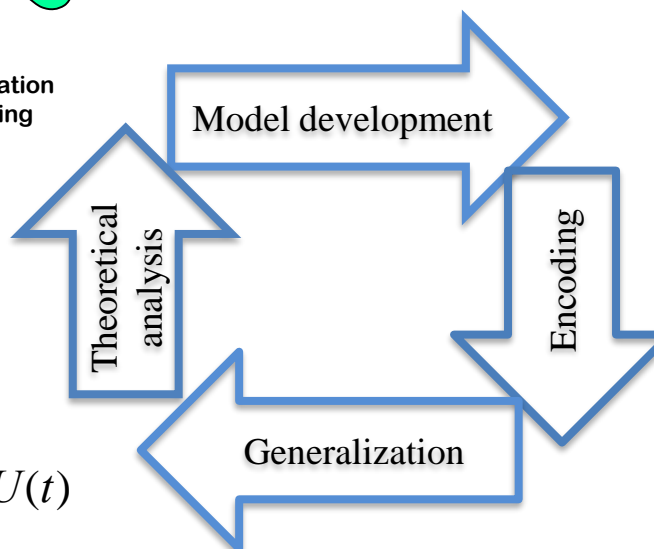
A: Basic process



B: Shared model structure

D: General model

$$\begin{cases} \dot{X}(t) = XACX(t) + BU(t) \\ X(t=0) = X_0 \end{cases}$$



C: Similar algorithm

$$\begin{cases} \text{Plant} \begin{cases} dX_1(t)/dt = b_1U(t) - \xi c_1X_1(t) \\ dX_2(t)/dt = b_2U(t) - \xi c_2X_2(t) \\ dX_3(t)/dt = b_3U(t) - \xi c_3X_3(t) \end{cases} \\ \text{Litter} \begin{cases} dX_4(t)/dt = \xi[c_1a_{41}x_1(t) + c_3a_{43}x_3(t) - c_4X_4(t)] \\ dX_5(t)/dt = \xi[c_1a_{51}x_1(t) + c_2x_2(t) + c_3a_{53}x_3(t) - c_5X_5(t)] \end{cases} \\ \text{SOM} \begin{cases} dX_6(t)/dt = \xi[c_4a_{64}x_4(t) + c_5a_{65}x_5(t) + c_7a_{67}x_7(t) + c_8a_{68}x_8(t) - c_6X_6(t)] \\ dX_7(t)/dt = \xi[c_5a_{75}x_5(t) + c_6a_{76}x_6(t) - c_7X_7(t)] \\ dX_8(t)/dt = \xi[c_6a_{86}x_6(t) + c_7a_{87}x_7(t) - c_8X_8(t)] \end{cases} \end{cases}$$

- Luo et al. 2003 GBC
- Luo and Weng 2011 TREE
- Luo et al. 2012
- Luo et al. 2015
- Luo et al. 2017

Matrix equation of CLM4.5

$$\frac{dX(t)}{dt} = B(t)I(t) - A\xi(t)KX(t) - V(t)X(t)$$

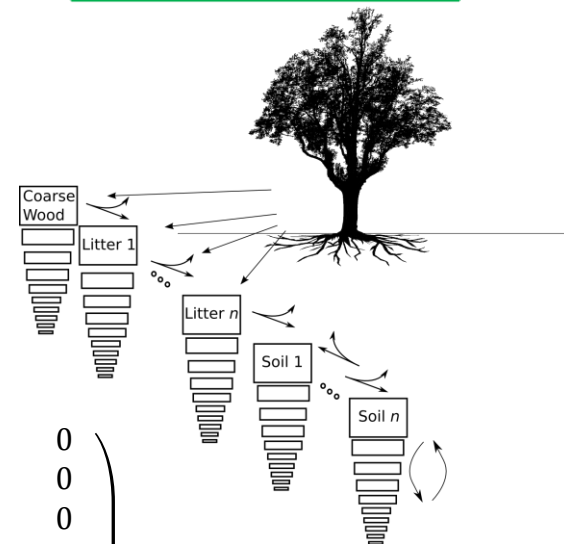
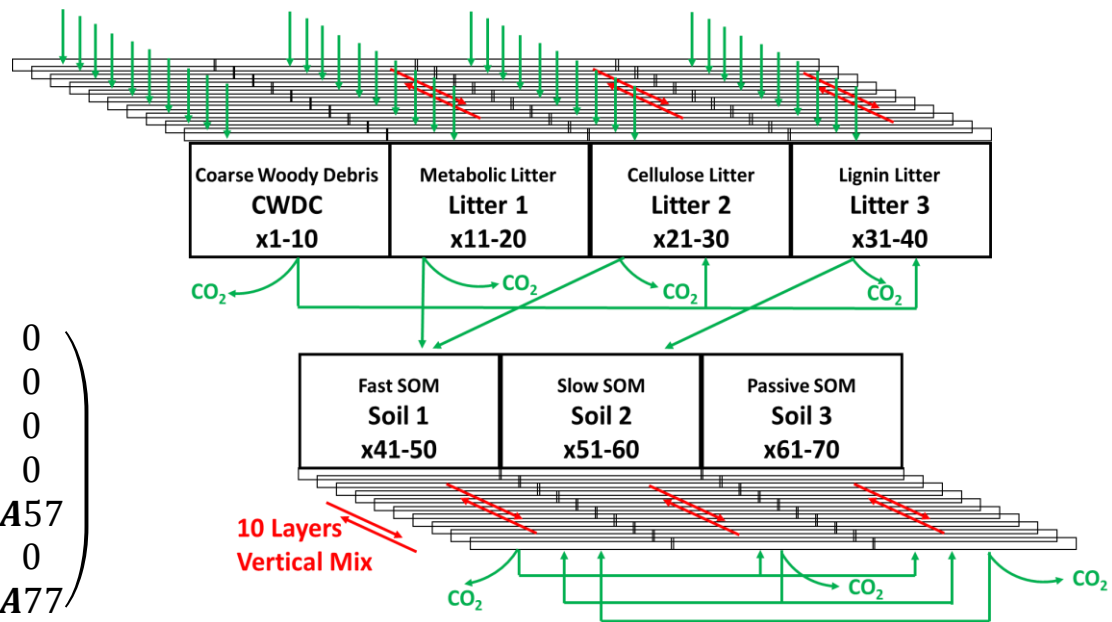
$$X(t) = (X_1(t), X_2(t), X_3(t), \dots, X_{70}(t))^T$$

$$A = \begin{pmatrix} A11 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & A22 & 0 & 0 & 0 & 0 & 0 \\ A31 & 0 & A33 & 0 & 0 & 0 & 0 \\ A41 & 0 & 0 & A44 & 0 & 0 & 0 \\ 0 & A52 & A53 & 0 & A55 & A56 & A57 \\ 0 & 0 & 0 & A64 & A65 & A66 & 0 \\ 0 & 0 & 0 & 0 & A75 & A76 & A77 \end{pmatrix}$$

$$A_{31} = \text{diag}(-f_{31}, -f_{31}, -f_{31}, -f_{31}, -f_{31}, -f_{31}, -f_{31}, -f_{31}, -f_{31}, -f_{31}, -f_{31})$$

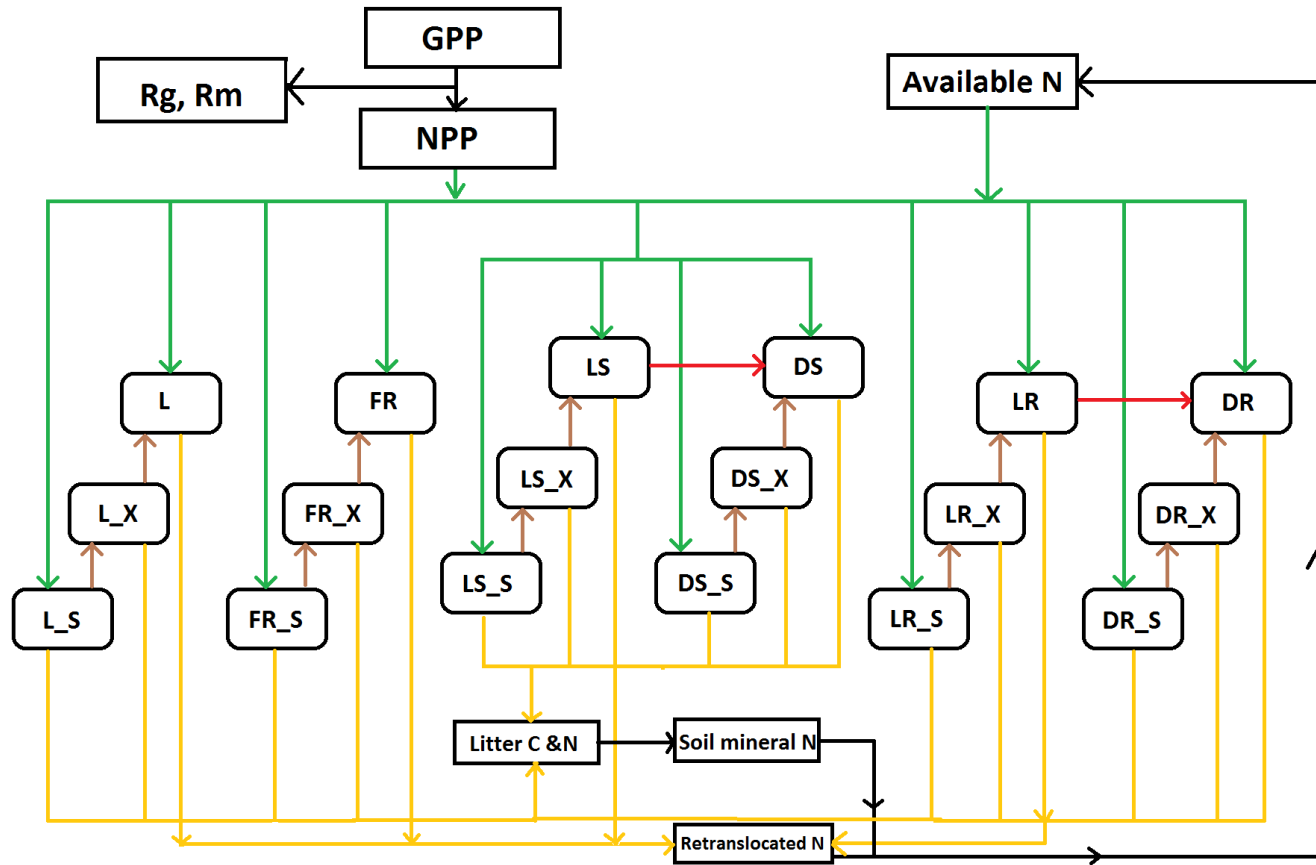
$$V(t) = \begin{pmatrix} V11 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & V22(t) & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & V33(t) & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & V44(t) & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & V55(t) & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & V66(t) & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & V77(t) \end{pmatrix}$$

$$V22 = \text{diag}(z_1, z_2, \dots, z_{10})^{-1} \begin{pmatrix} g_1 & -g_1 & 0 & 0 & \dots & 0 & 0 & 0 \\ -h_2 & h_2 + g_2 & -g_2 & 0 & \dots & 0 & 0 & 0 \\ 0 & -h_3 & h_3 + g_3 & -g_3 & \dots & 0 & 0 & 0 \\ 0 & 0 & -h_4 & h_4 + g_4 & \dots & 0 & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots & \ddots & \vdots & \vdots & \vdots \\ 0 & 0 & 0 & 0 & \dots & h_8 + g_8 & -g_8 & 0 \\ 0 & 0 & 0 & 0 & \dots & -h_9 & h_9 + g_9 & -g_9 \\ 0 & 0 & 0 & 0 & \dots & 0 & -h_{10} & h_{10} \end{pmatrix}$$



Huang et al. 2018
Global Change Biology

CLM vegetation C&N: phenology, fire etc.



—	Allocation, Phenology
—	Phenology offset, Background turnover, Gap mortality, Fire
—	Phenology, Fire
—	Phenology
	Controlling Procedure

- | | | |
|-----------------------|----------------------------------|--------------------------------|
| L: leaf; | L_X: leaf transfer; | L_S: leaf storage |
| FR: fine root; | FR_X: fine root transfer; | FR_S: fine root storage |
| LS: live stem; | LS_X: live stem transfer; | LS_S: live stem storage |
| DS: dead stem; | DS_X: dead stem transfer; | DS_S: dead stem storage |
| LR: live coarse root; | LR_X: live coarse root transfer; | LR_S: live coarse root storage |
| DR: dead coarse root; | DR_X: dead coarse root transfer; | DR_S: dead coarse root storage |

Matrix equation of vegetation carbon dynamics

$$\frac{d}{dt}X(t) = (A_{ph}(t)K_{ph}(t) + A_{gm}(t)K_{gm}(t) + A_{fi}(t)K_{fi}(t))X(t) + B(t)F(t)$$

Diagram illustrating the components of the matrix equation for vegetation carbon dynamics:

- $A_{ph}(t)$: C transfer of phenology
- $K_{ph}(t)$: C turnover of phenology
- $A_{gm}(t)$: C transfer of gap mortality
- $K_{gm}(t)$: C turnover of gap mortality
- $A_{fi}(t)$: C transfer of fire
- $K_{fi}(t)$: C turnover of fire
- $X(t)$: pool state
- $B(t)F(t)$: input
- $B(t)$: allocation

Vegetation nitrogen dynamics

$$\frac{d}{dt}N(t) = (A_{Nph}(t)K_{Nph}(t) + A_{Ngm}(t)K_{Ngm}(t) + A_{Nfi}(t)K_{Nfi}(t))N(t) + B_N(t)F_N(t)$$

Mathematical framework

Vertical profile

$$\frac{dX(t)}{dt} = (A\xi(t)K + V(t))X(t) + B(t)u(t)$$

$$\frac{dX(t)}{dt} = A\xi(t)KX(t) + Bu(t)$$

Developing models at different levels of complexity under one overarching theory

$$\frac{d}{dt}X(t) = (A_{ph}(t)K_{ph}(t) + A_{gm}(t)K_{gm}(t) + A_{fi}(t)K_{fi}(t))X(t) + B(t)F(t)$$

C transfer of phenology
C transfer of gap mortality
C transfer of fire
pool state
input

C turnover of phenology
C turnover of gap mortality
C turnover of fire
allocation

Vegetation dynamics

Unifying land carbon cycle models

Matrix models

1. CLM 3.5
2. CLM4.0
3. CLM4.5
4. CLM5.0
5. CABLE
6. LPJ-GUESS
7. ORCHIDEE
8. BEPS
9. TECO
10. JULES
11. IBIS

In progress

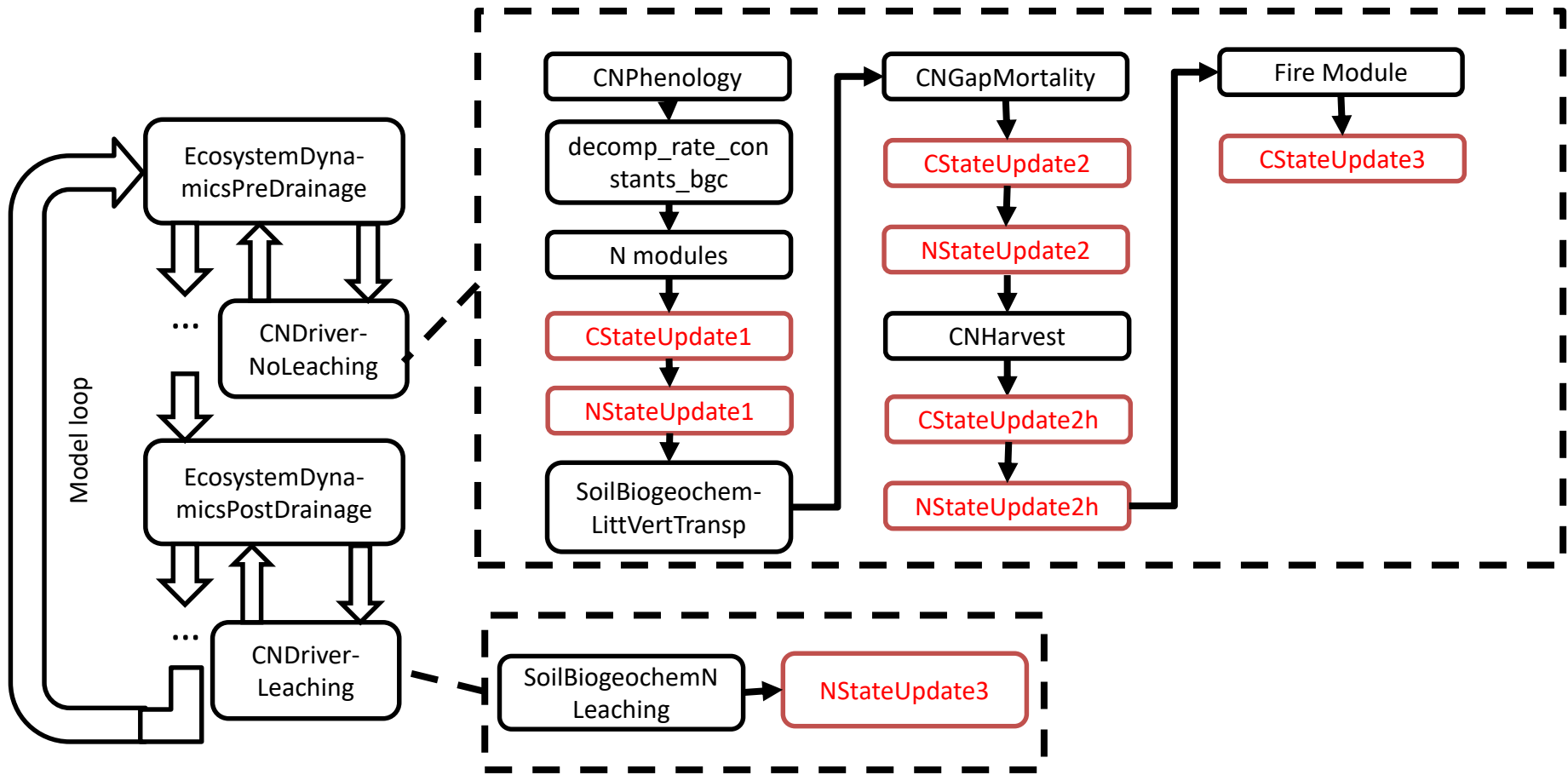
1. LM3V-N

10 nonlinear microbial models by Carlos Sierra

Luo et al. 2017

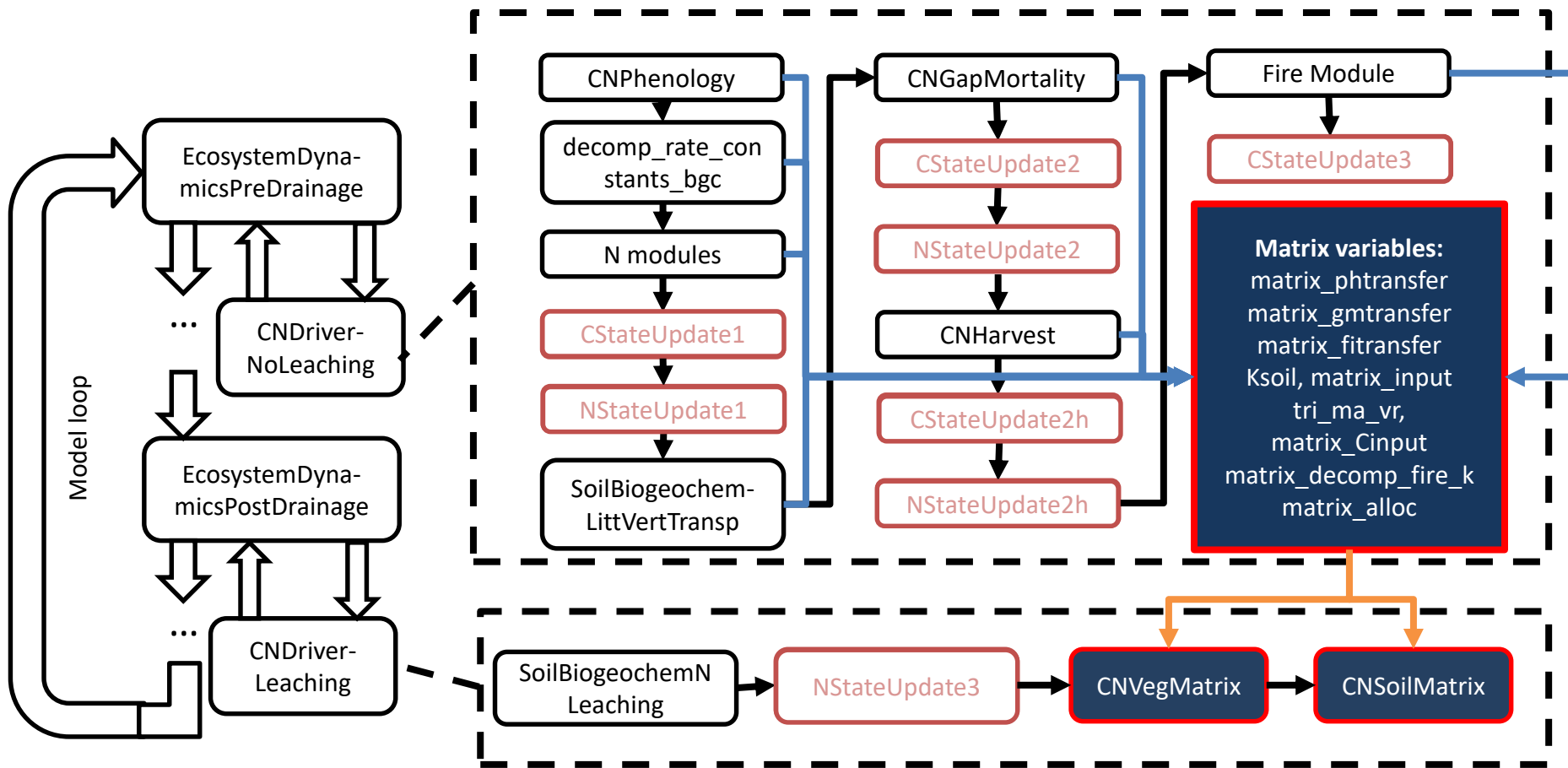
Luo et al. *to be submitted*

Programming



CLM5.0 biogeochem cycle

Programming



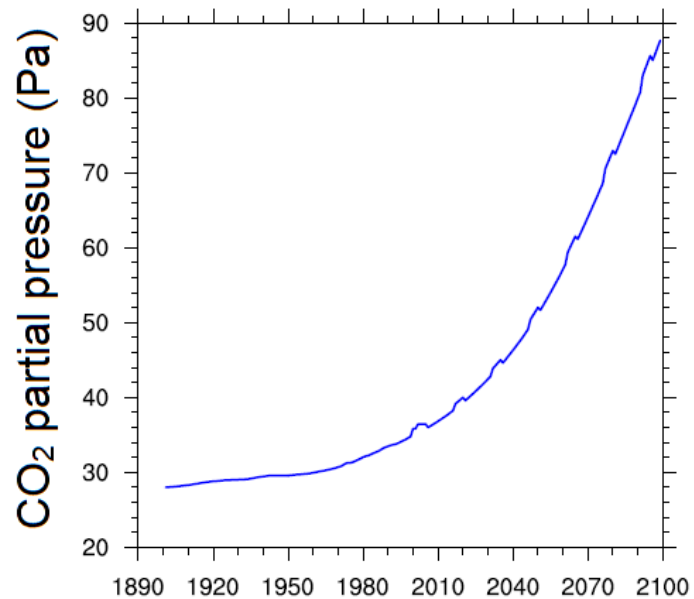
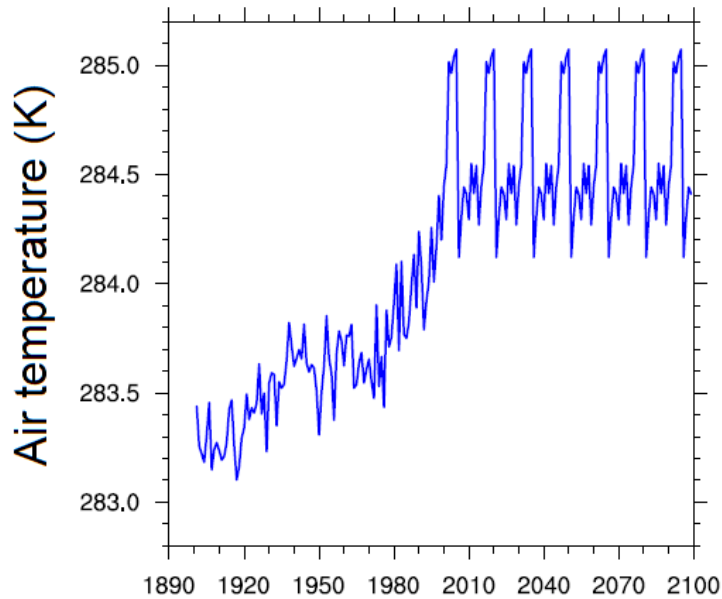
CLM5.0 biogeochem cycle with matrix

Technique test

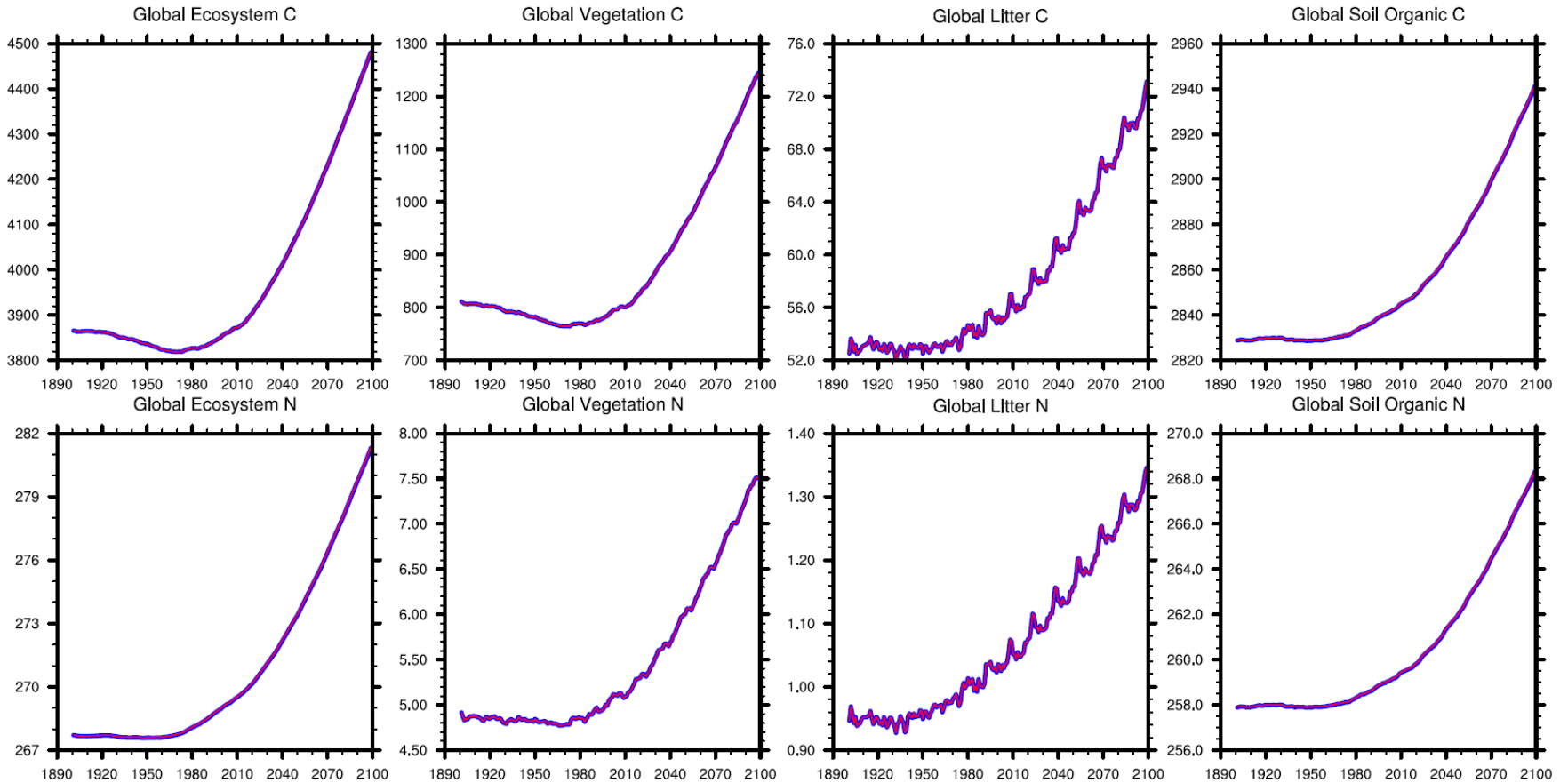
- Use automated tools (CESM test suites, use “aux_clm” testlists) to test CLM5.0 code both with matrix on and matrix off.
- Code with matrix on:
 - In total 159 tests, 149 tests are passed, 2 tests are expected fails, 6 tests are due to restart inconsistency, 1 is from C balance issue, 1 is from wrong initial file address pointed to old address.
- Code with matrix off:
 - In total 159 tests, 156 tests are passed, 2 tests are expected fails, 1 is from wrong initial file address pointed to old address..

Scientific test: Simulations

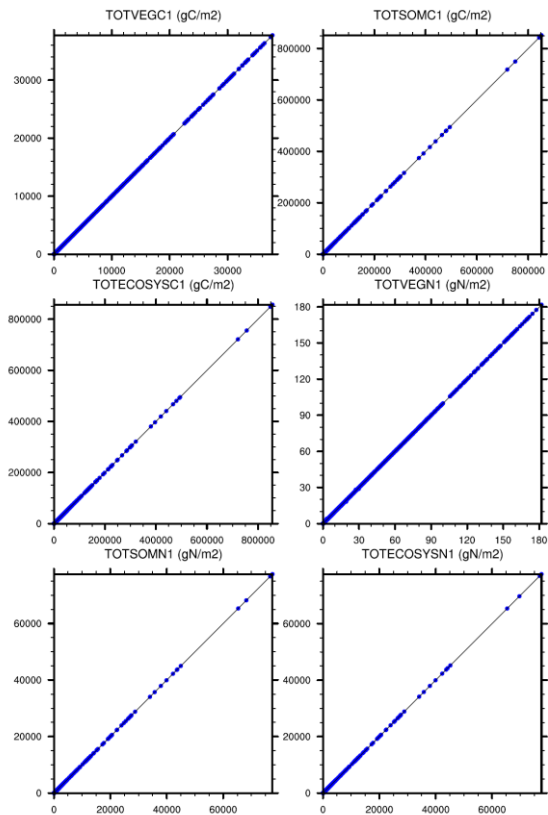
Time	Resolution	Compset	Climate	[CO ₂]
1901-2099	4x5	IHistCIm50Bgc	GSWP3	CMIP5



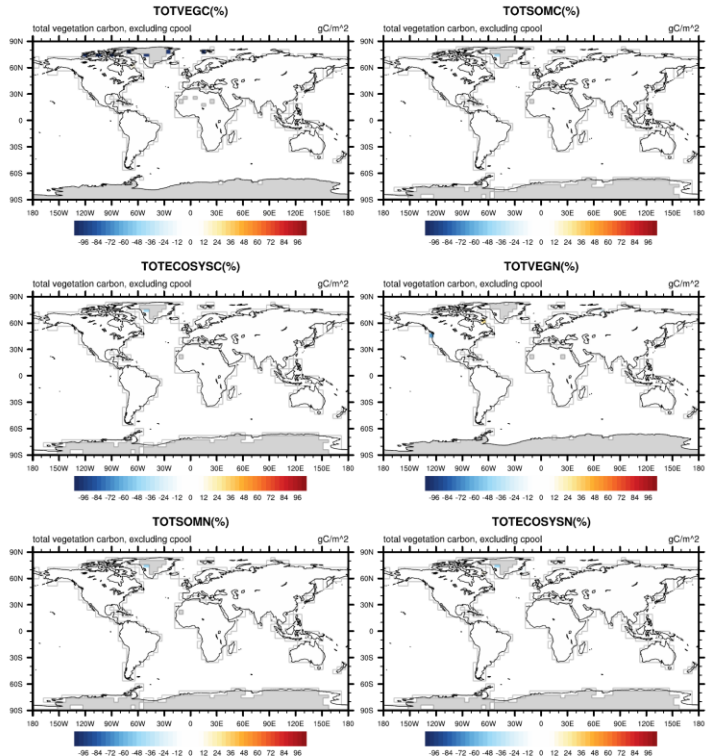
Matrix Vs default RCP 8.5 (global-level)



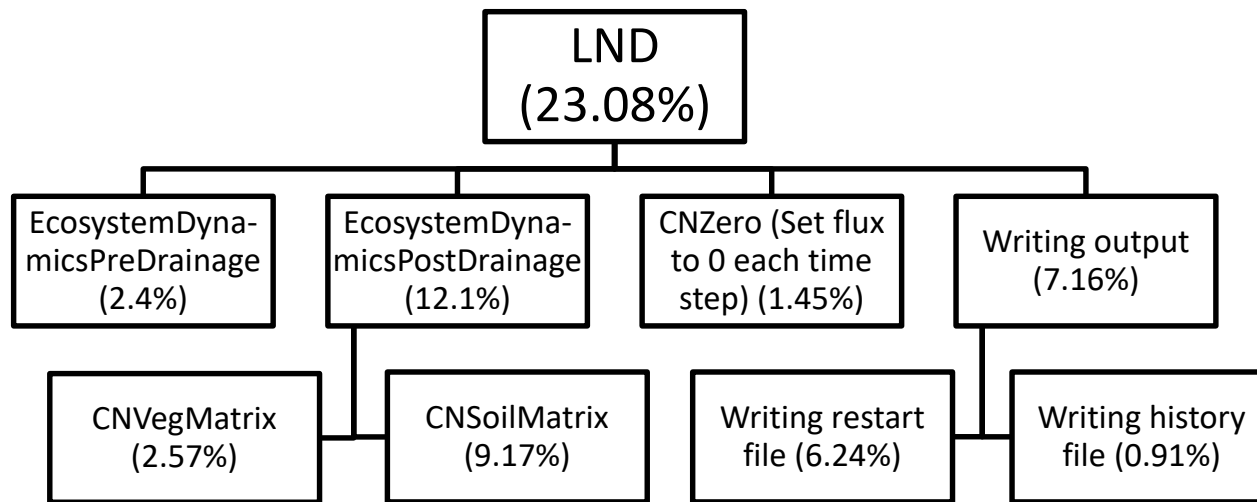
Matrix Vs default (Global)



(Matrix – Default)/Default * 100%



Computational efficiency for forward modeling



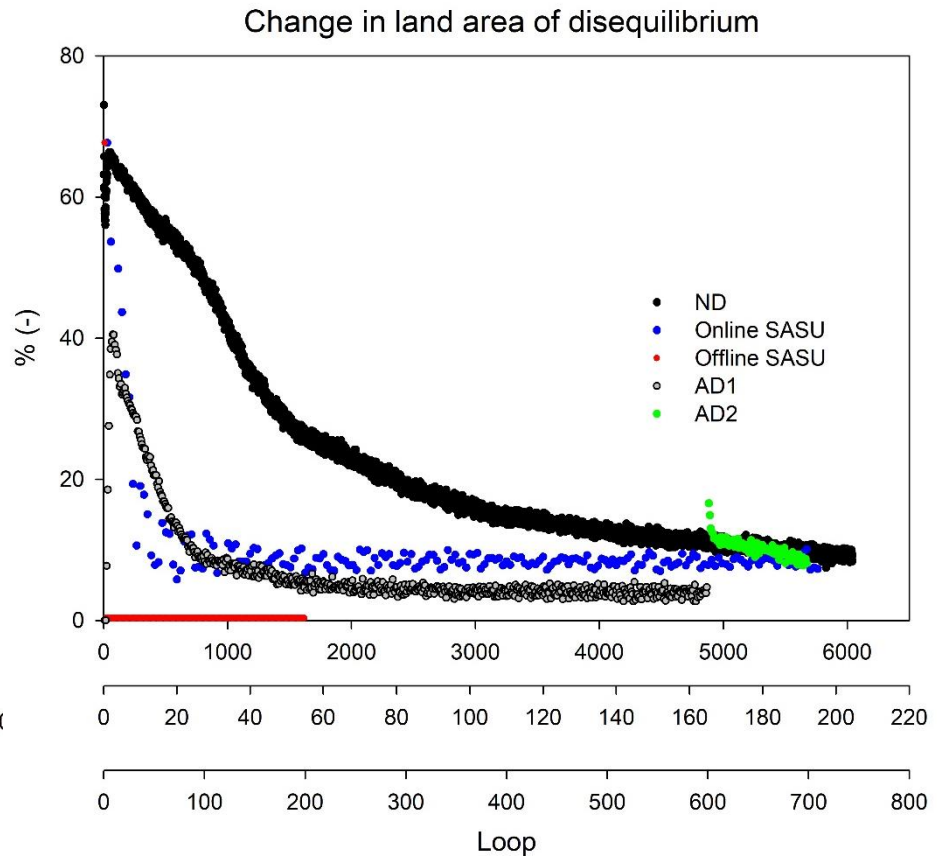
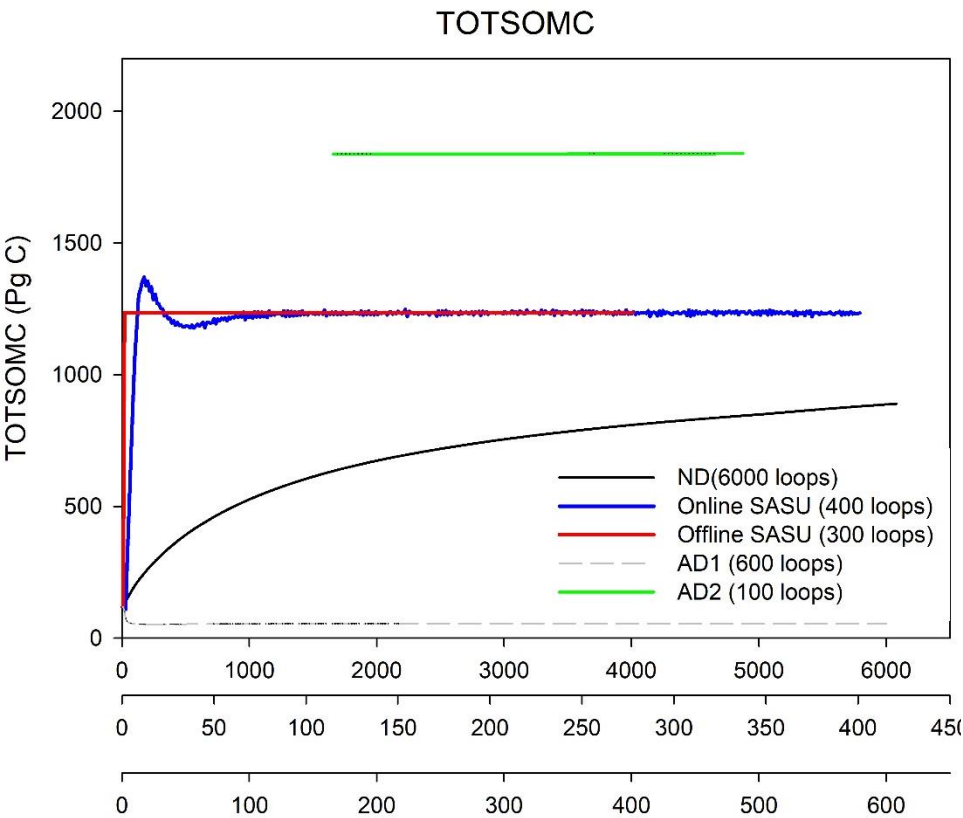
- The fraction in the bracket is a relative increase of computational time:

$$(T_{i, \text{matrixon}} - T_{i, \text{matrixoff}}) / T_{\text{total, matrixoff}} * 100\%$$

$T_{i, \text{matrixon}}$ is the computational time consumed by subroutine i , in matrix simulation. $T_{i, \text{matrixoff}}$ is the computational time consumed by subroutine i , in control simulation, $T_{\text{total, matrixoff}}$ is total computational time in control simulation.

- Comparison is between one-month global simulation ($2^{\circ} \times 2^{\circ}$) with matrix on and off.

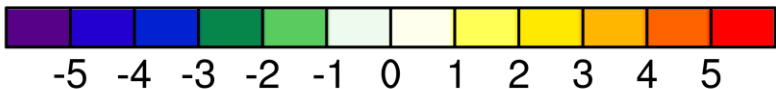
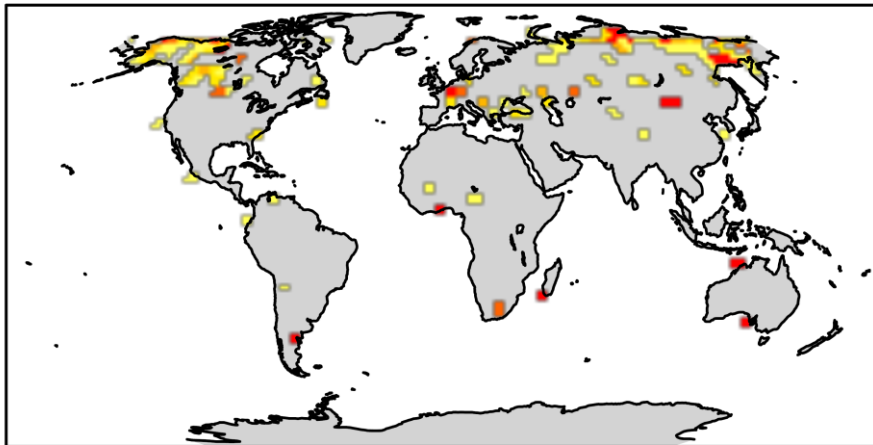
Computational efficiency for Spin-up



Forcing data: One-year (1911) GSWP3V

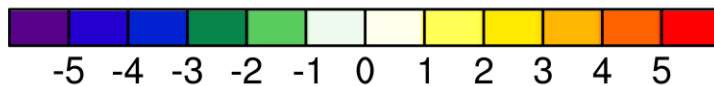
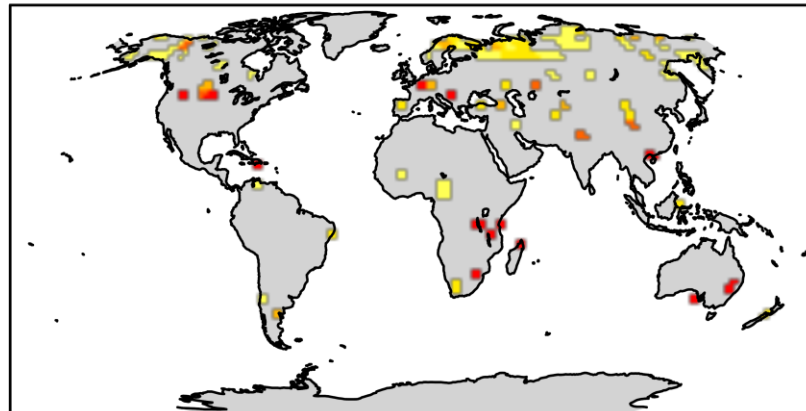
ND

TOTSOMC(gC m⁻²) loops: 6000 - 5999



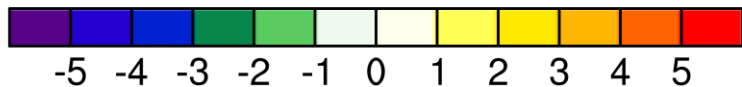
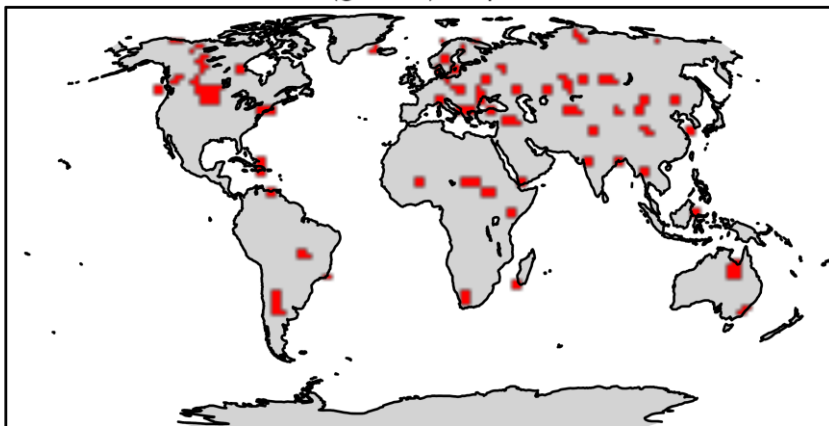
AD

TOTSOMC(gC m⁻²) loops: 700 - 699



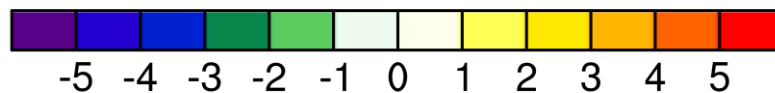
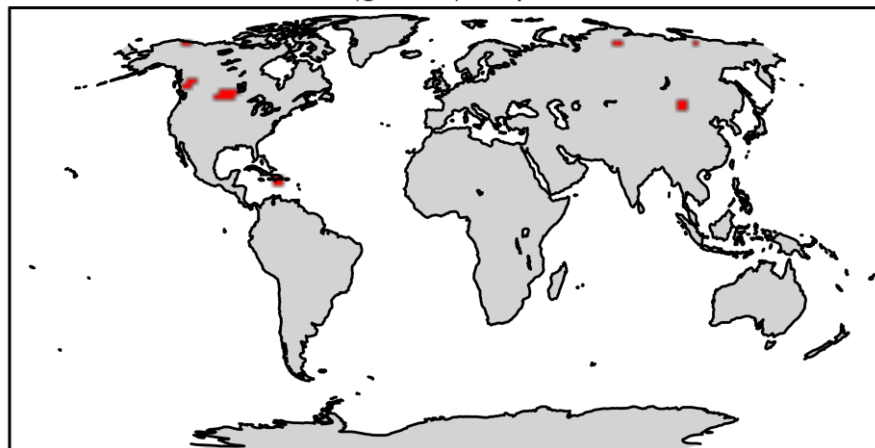
Online SASU

TOTSOMC(gC m⁻²) loops: 200 - 199

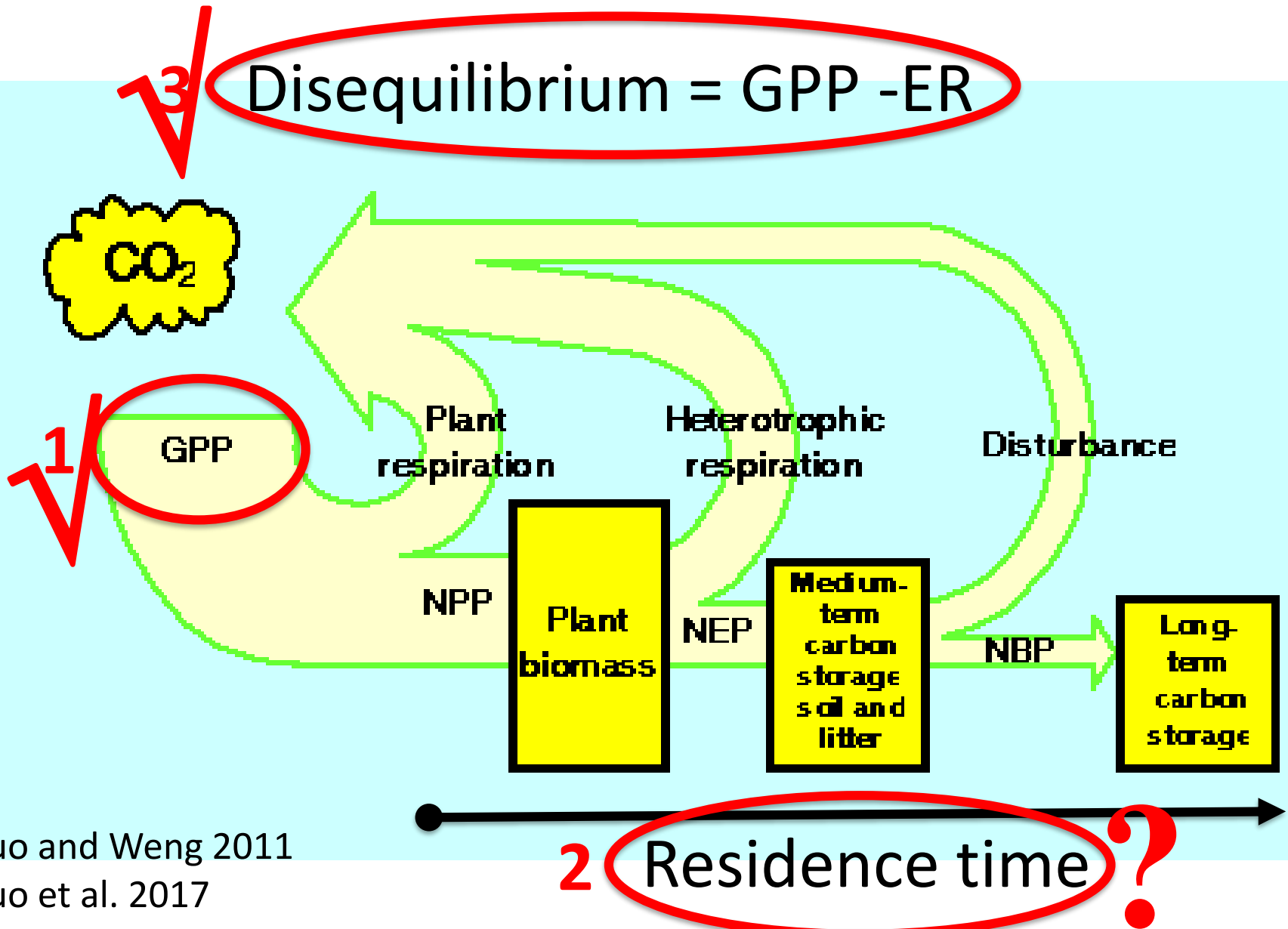


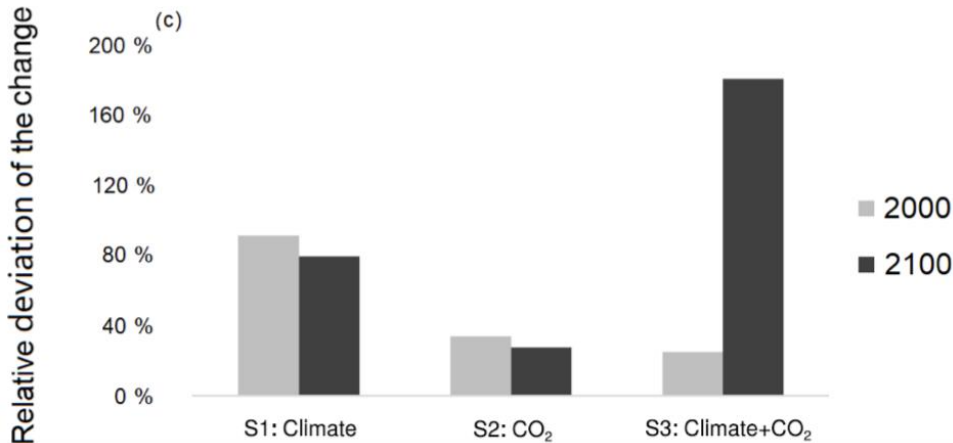
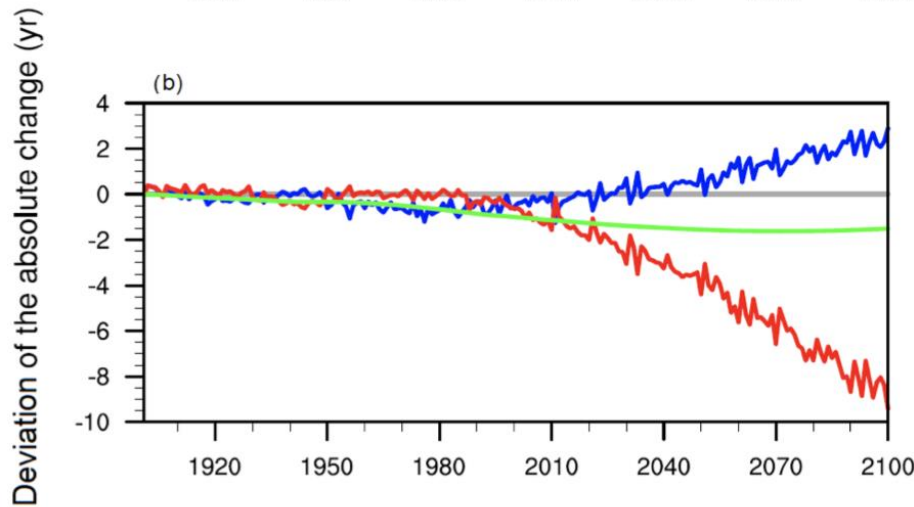
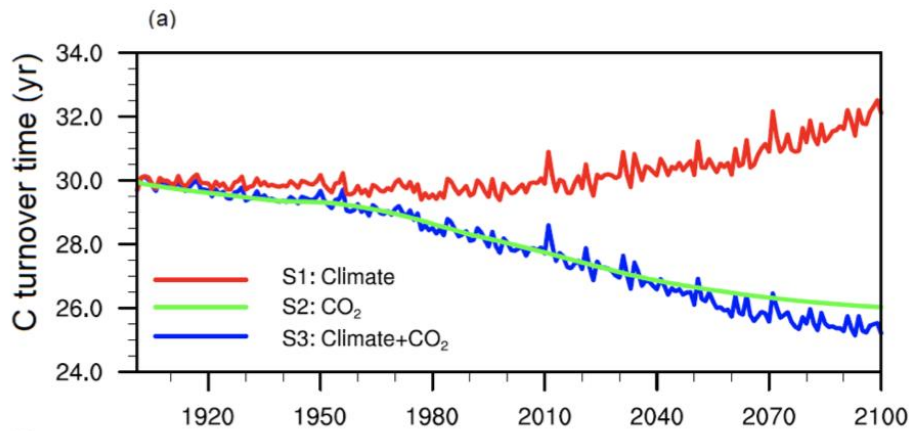
Offline SASU

TOTSOMC(gC m⁻²) loops: 200 - 199



Diagnostics

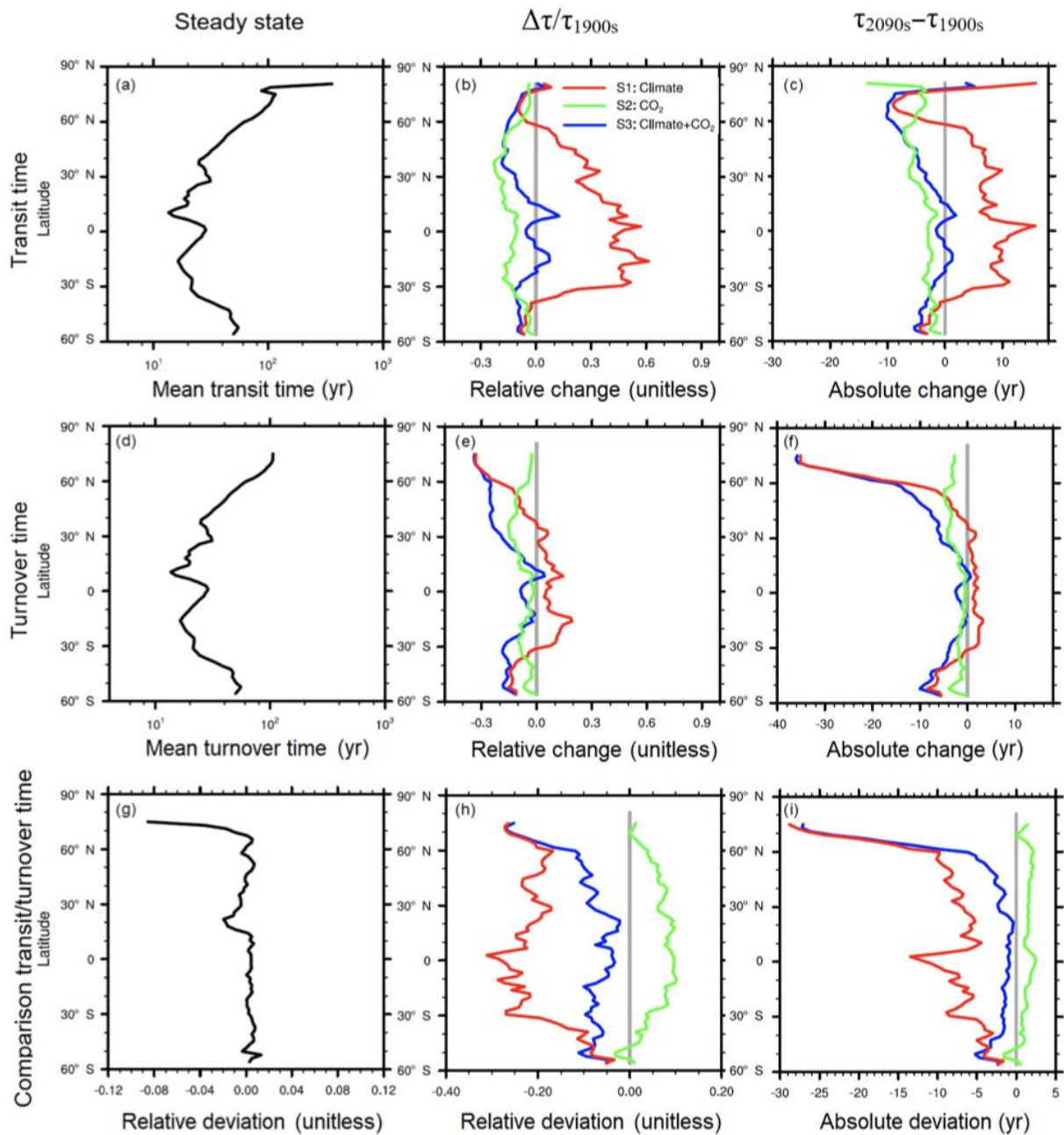




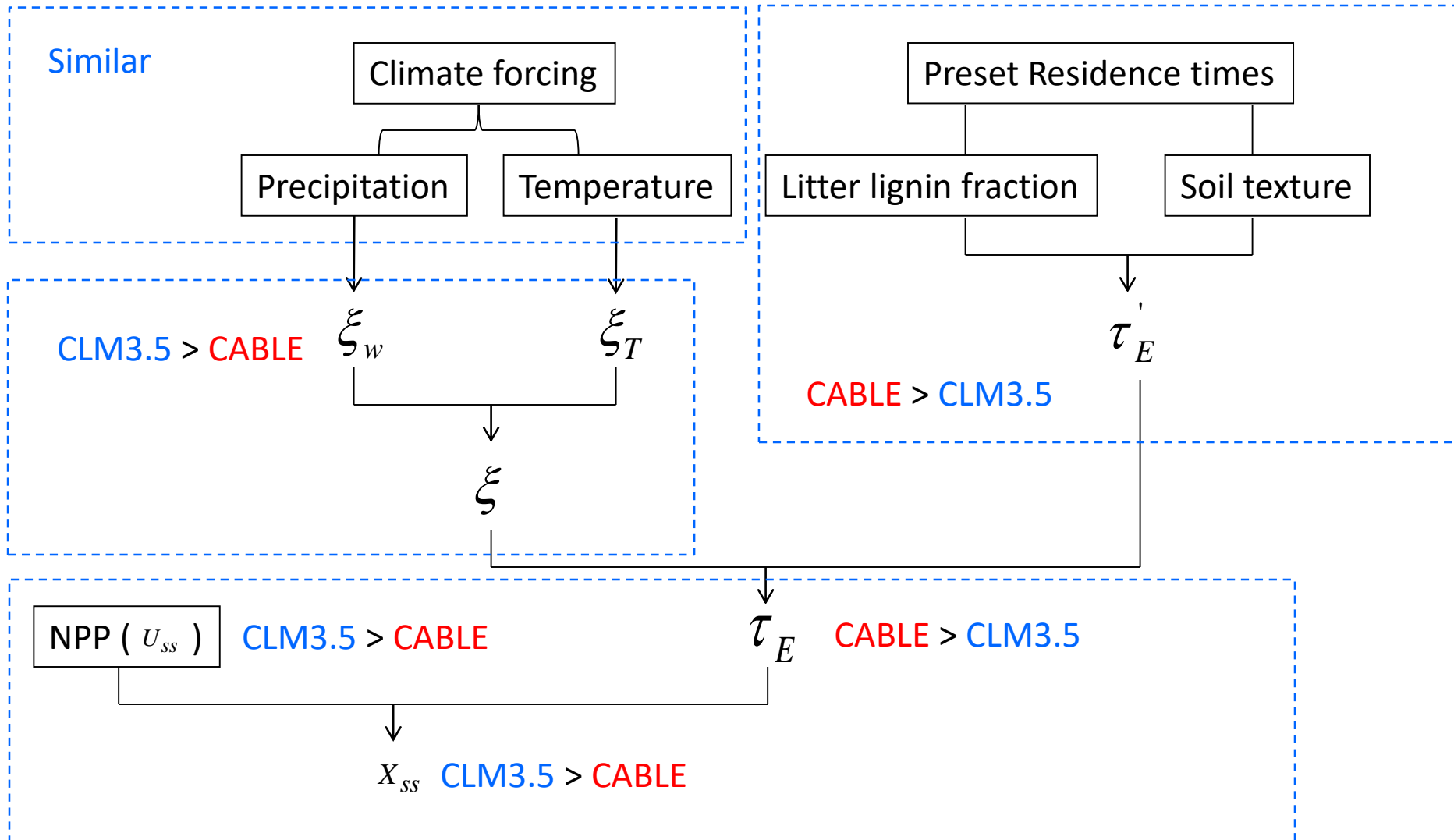
Residence (or transit) time =
time spent by carbon from
entry to exit in a network of
multiple pools

Turnover time = pool/flux

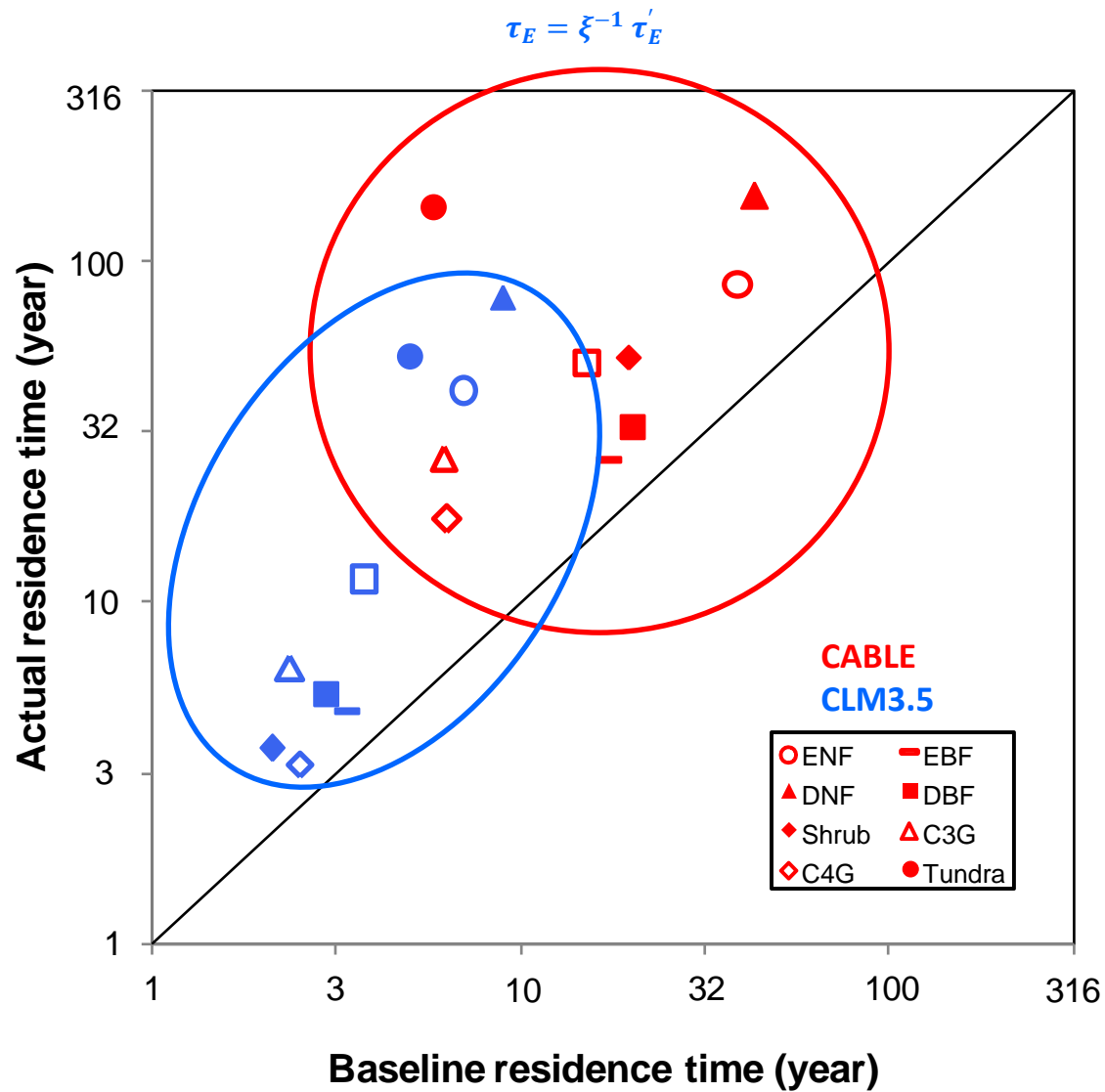
Lu et al. 2018 Biogeosciences



Traceability analysis for model Intercomparison



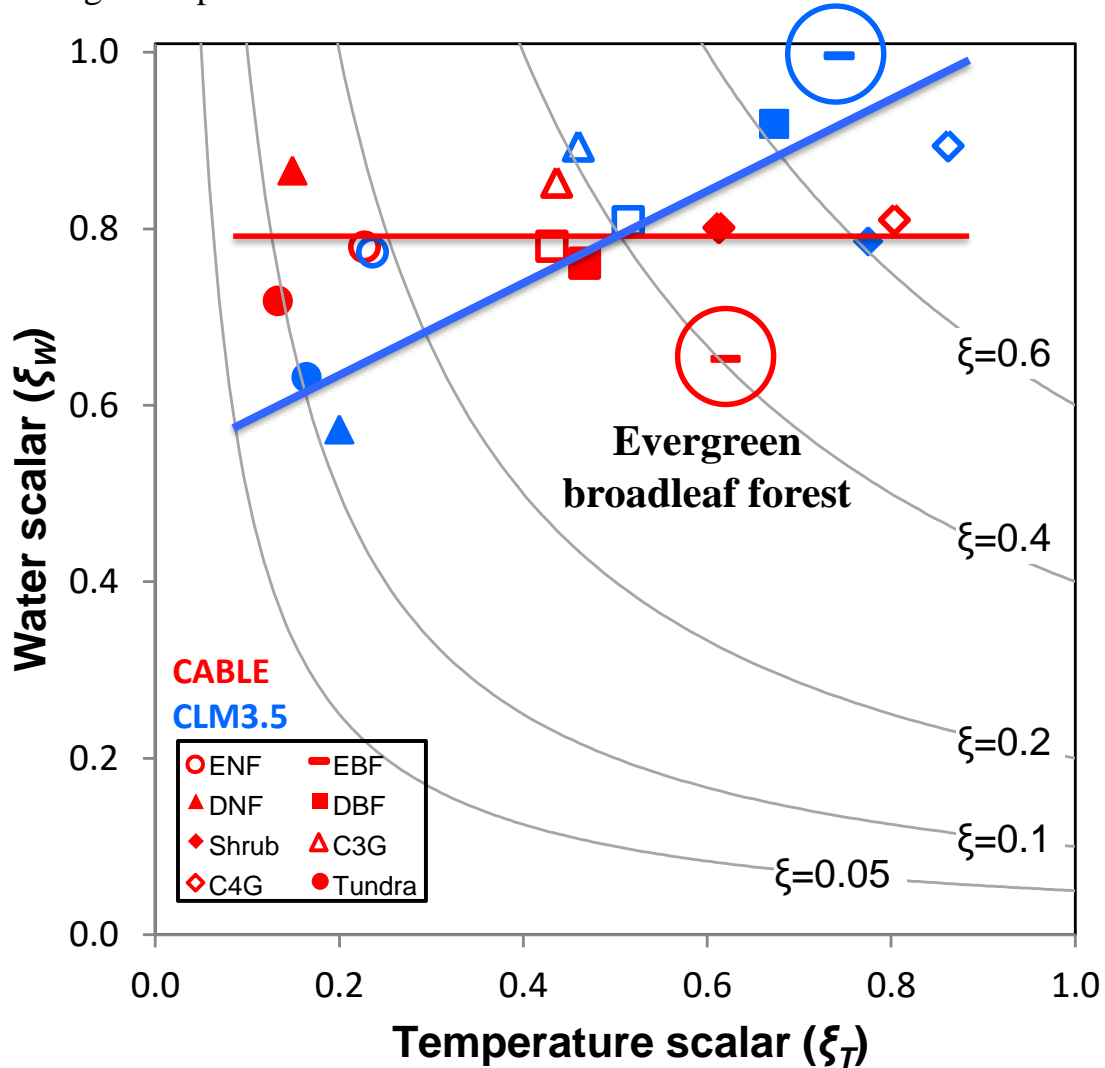
Longer residence times in CABLE than CLM3.5



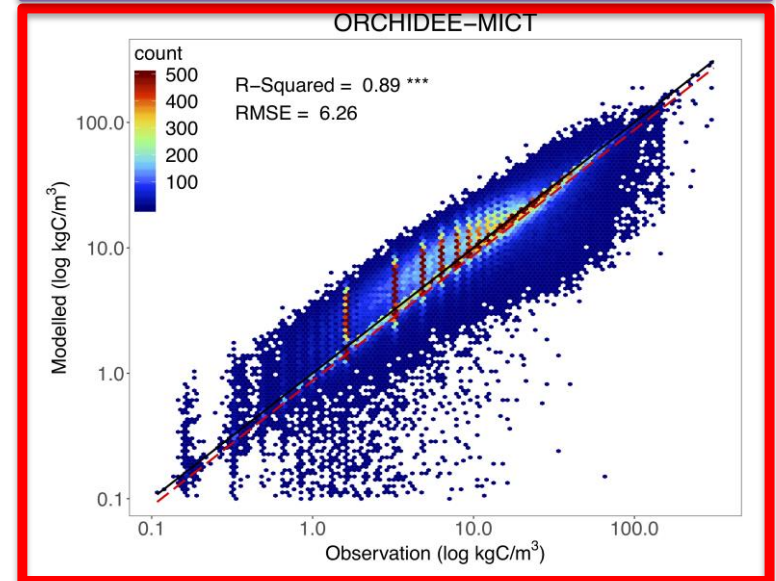
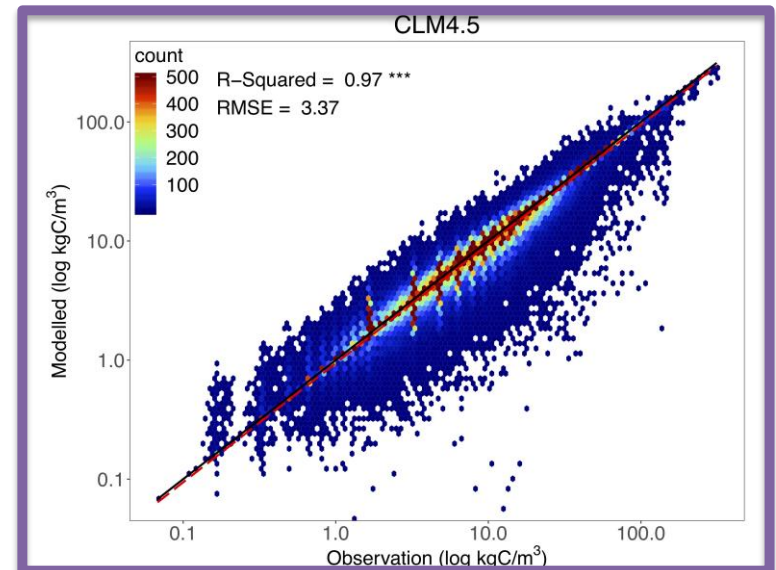
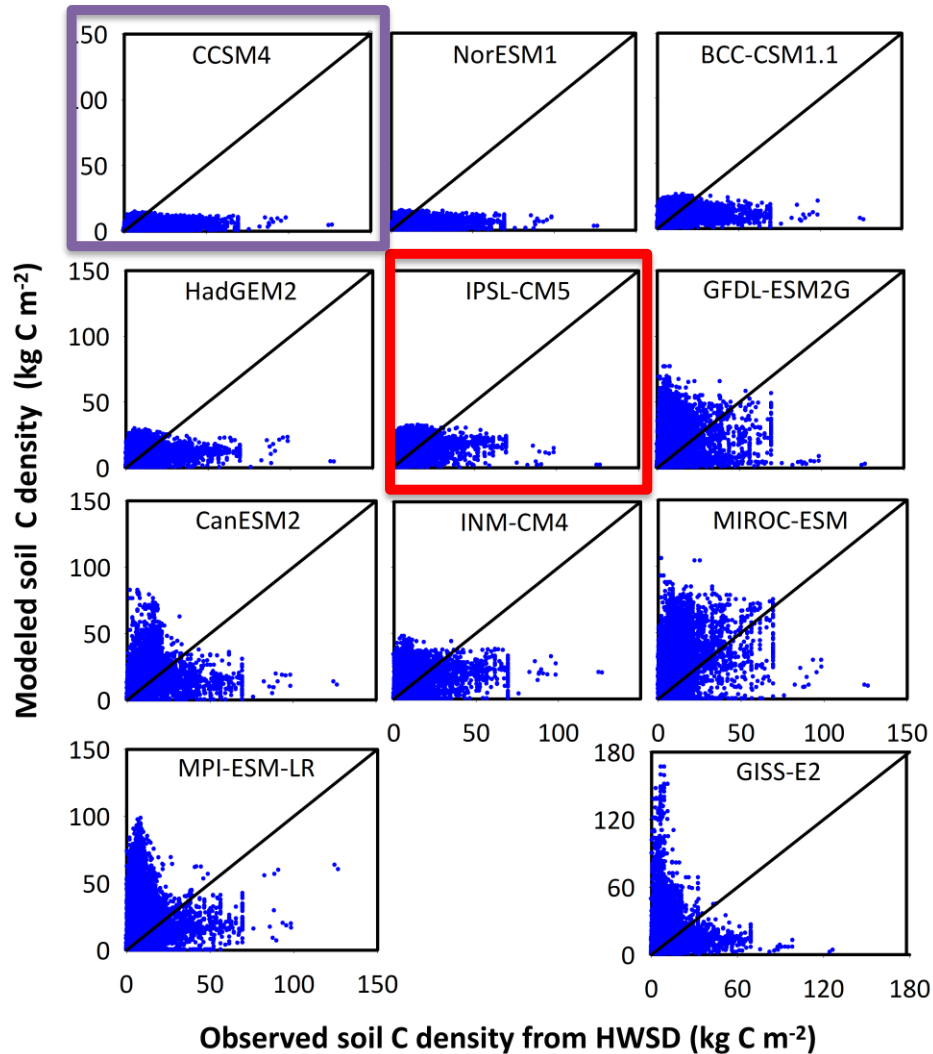
Temperature and water scalars

$$\xi = \xi_T \xi_W$$

- Larger temperature and water limitations on most biomes in CABLE



Data-model comparison



Luo et al. 2015 *Global Change Biology*

Tao et al. *In prep.*

Parameterization with data from ~30,000 soil profiles

Method	Original Model	Site by Site	Random Sampling	Neural Networking	One Batch (MPI)
R-Squared	0.57	0.97	0.59	0.69	0.64
RMSE	15.86	3.37	12.39	11.23	11.33

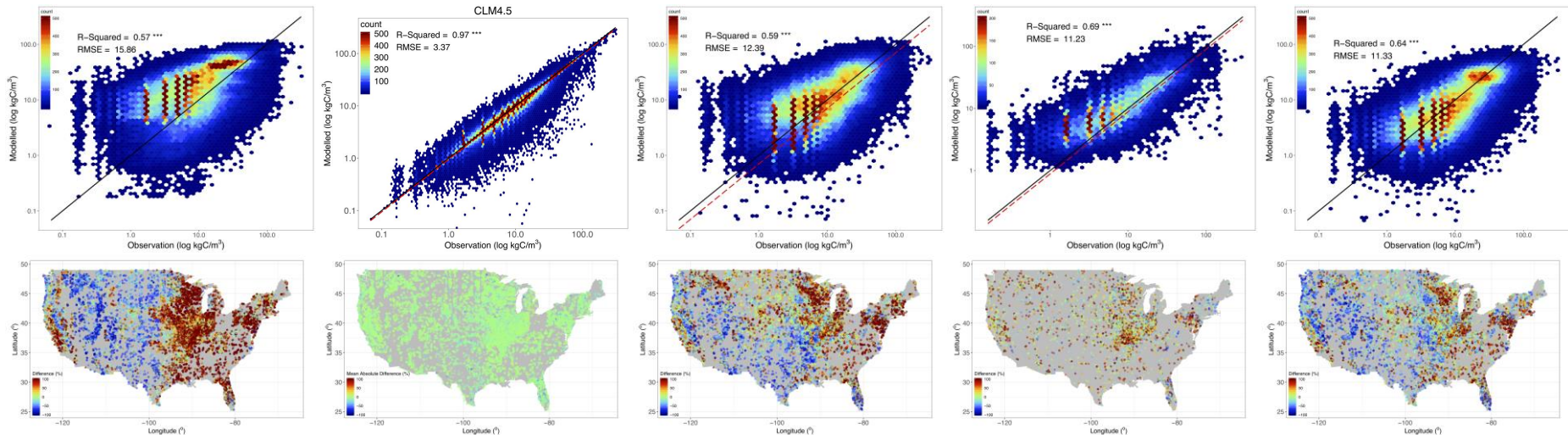
Original Model

Site by Site

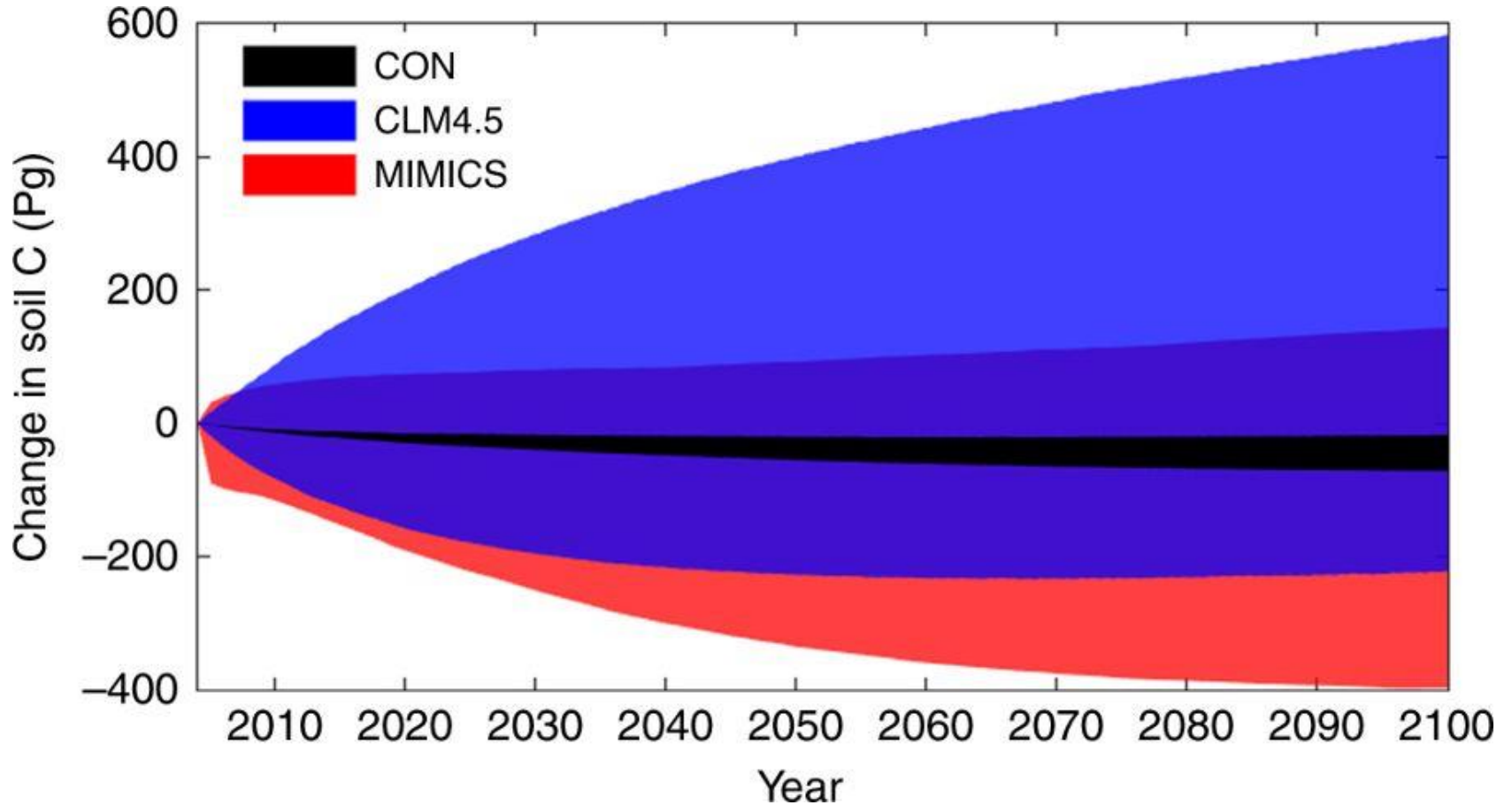
Random Sampling

Neural Networking

One Batch

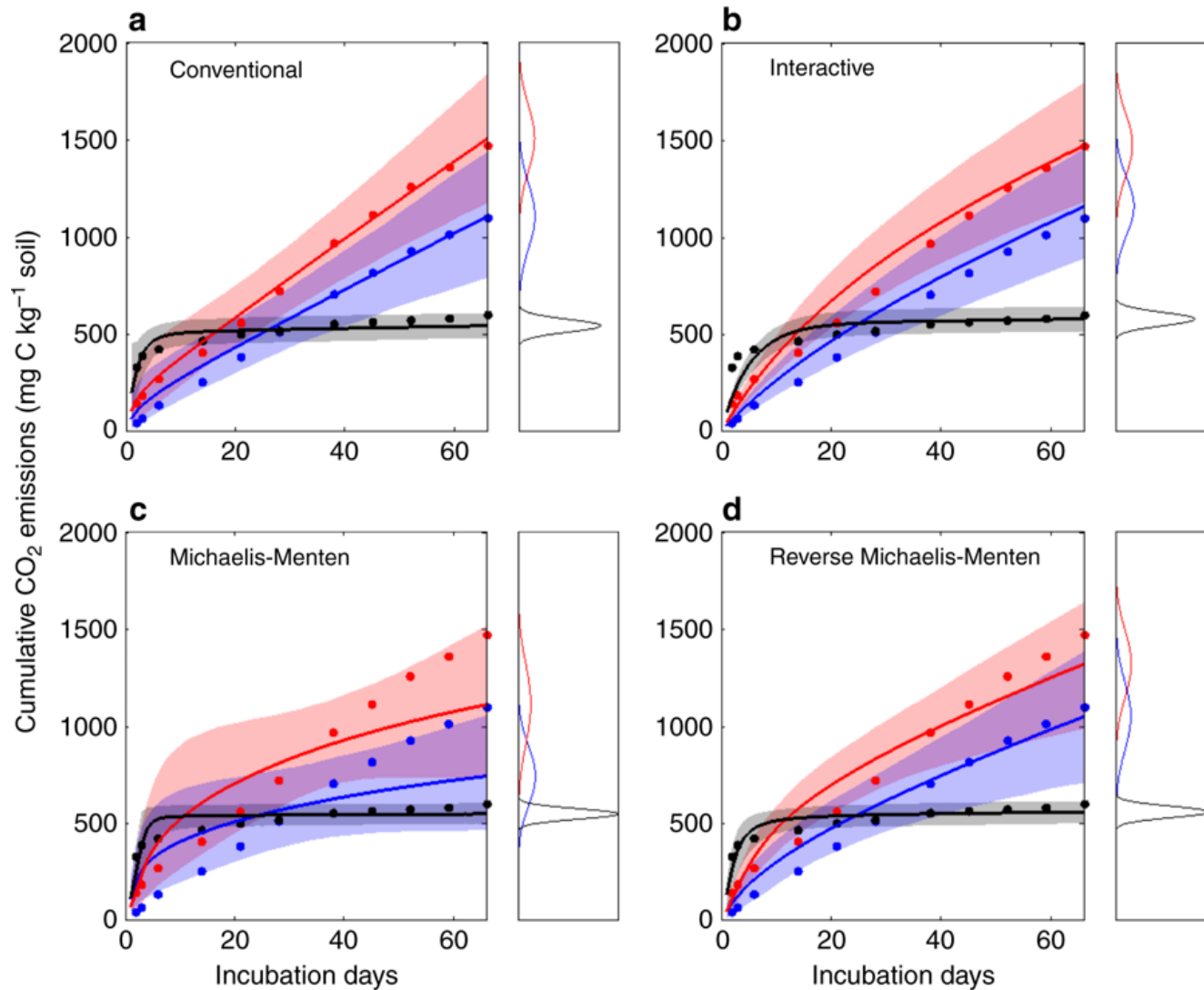


Uncertainty in projections



Shi et al. 2018 *Nat. Comm.*

Selection of model structure for priming



2nd Training Course on
New Advances in Land Carbon Cycle Modeling
Flagstaff, AZ, USA, May 13-24, 2019

- **Modelers** to gain simplicity in coding, diagnostic capability, computational efficiency for your models
- **Empiricists** to use your data to constrain models toward ecological forecasting
- **Scientists** who want to learn modeling, data assimilation, and ecological forecasting

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Lifen Jiang, lifen.jiang@nau.edu

Discussion points

1. How much details do we need in a model?
2. How can we develop an efficient model development-evaluation-improvement continuum?