

**Spin-up of  
biogeochemical  
models**

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# A semi-analytical solution to accelerate spin-up of a coupled carbon and nitrogen land model to steady state

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## Abstract

The spin-up of land models to steady state of coupled carbon-nitrogen processes is computationally so costly that it becomes a bottleneck issue for global analysis. In this study, we introduced a semi-analytical solution (SAS) for the spin-up issue. SAS is fundamentally based on the analytic solution to a set of equations that describe carbon transfers within ecosystems over time. SAS is implemented by three steps: (1) having an initial spin-up with prior pool-size values until net primary productivity (NPP) reaches steady state, (2) calculating quasi steady-state pool sizes by letting fluxes of the equations equal zero, and (3) having a final spin-up to meet the criterion of steady state. Step 2 is enabled by averaged time-varying variables over one period of repeated driving forcings. SAS was applied to both site-level and global scale spin-up of the Australian Community Atmosphere Biosphere Land Exchange (CABLE) model. For the carbon-cycle-only simulations, SAS saved 95.7 % and 92.4 % of computational time for site-level and global spin-up, respectively, in comparison with the traditional method. For the carbon-nitrogen-coupled simulations, SAS reduced computational cost by 84.5 % and 86.6 % for site-level and global spin-up, respectively. The estimated steady-state pool sizes represent the ecosystem carbon storage capacity, which was 12.1 kgCm<sup>-2</sup> with the coupled carbon-nitrogen global model, 14.6 % lower than that with the carbon-only model. The nitrogen down-regulation in modeled carbon storage is partly due to the 4.6 % decrease in carbon influx (i.e., net primary productivity) and partly due to the 10.5 % reduction in residence times. This steady-state analysis accelerated by the SAS method can facilitate comparative studies of structural differences in determining the ecosystem carbon storage capacity among biogeochemical models. Overall, the computational efficiency of SAS potentially permits many global analyses that are impossible with the traditional spin-up methods, such as ensemble analysis of land models against parameter variations.

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## 1 Introduction

Modeling ecosystem biogeochemical cycles is an initial value problem. It requires setting up initial values of all state variables (e.g., carbon and nitrogen pools sizes) in any biogeochemical models before scientists can use the models for any analysis.

5 The initial values are either estimated from observations (D'Odorico et al., 2004; Luo and Reynolds, 1999) or assumed to be at steady state. The latter is usually achieved by performing long model simulations (i.e., spin-up) until no trend of change in pool sizes over many periods of the repeated climate forcing even though the pool sizes vary seasonally and inter-annually within one period of the repeated forcing (Johns et al., 1997; McGuire et al., 1997; Thornton et al., 2002; Yang et al., 1995). Spinning  
10 biogeochemical models to steady state is computationally expensive, especially so when simulations are performed with global biogeochemical models of coupled C and N cycles (Thornton and Rosenbloom, 2005). In general, a fully coupled earth system model with biogeochemical cycles needs to be spun up in several sequential steps, and each step can take several thousands of simulation years (Doney et al., 2006). Spin-  
15 up of a fully coupled earth system model with a relatively coarse resolution ( $\approx 3.75^\circ$ ) atmospheric model is estimated to take hundreds of real-world days of computation using the present National Center for Atmosphere Research (NCAR) supercomputers (Jochum et al., 2010). As a consequence, spin-up has become a serious constraint on  
20 global modeling analysis of biogeochemical cycles.

Thornton and Rosenbloom (2005) has explored a few spin-up methods for achieving steady states of a coupled carbon-nitrogen ecosystem model. The spin-up of their model begins with initial values of no soil organic matter (SOM) and very small plant pools (Thornton et al., 2002). During the spin-up, the accumulation rate of SOM  
25 strongly depends on the nitrogen addition rate in their model. They periodically increase the mineral nitrogen supply during the early stage of the spin-up to accelerate the spin-up. Each nitrogen addition period is followed by a period with reducing nitrogen input linearly to the normal level (Thornton and Rosenbloom, 2005; Thornton

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et al., 2002). The efficiency of this punctuated nitrogen addition method is low without an optimal combination of the nitrogen addition and reduction periods. A prior analysis for searching such optimal combinations is needed when this method is applied to a new model. An accelerated decomposition method is based on an assumption of linear scaling between decomposition rates and litter/soil pool sizes. A linearity test between decomposition rates and pool sizes is needed before performing this method for each new model (Thornton and Rosenbloom, 2005). The accelerated decomposition method can distort interactions between carbon and nitrogen cycles in CASACNP model (Wang, 2012) and ocean physics for spinning up an ocean model (Bryan and Lewis, 1979; Danabasoglu et al., 1996). Although the above methods have been used in some modeling studies (Randerson et al., 2009), most models still use the traditional spin-up method with long-term iterative simulations to achieve the steady states of variables.

Carbon processes in terrestrial ecosystems can be represented by first-order, linear differential equations (Bolker et al., 1998; Luo and Weng, 2011; Luo et al., 2012). This property renders a possibility to obtain an analytical solution of steady states for terrestrial carbon cycle models (Bolker et al., 1998; Comins, 1994; Govind et al., 2011; King, 1995; Luo et al., 2001). Ludwig et al. (1978), for example, has used a two-step approximation. They first calculated the steady-state pool sizes of the fast variables by holding the slow variables fixed, and then analyzed the slow variables with the fast variables held at corresponding steady-state pool sizes. Most of the analytical solutions (Comins, 1994; Govind et al., 2011; King, 1995; Luo et al., 2001) are obtained by using constant net primary productivity (NPP). However, temporal climate fluctuations and seasonal plant growth make it difficult to get analytical solution for state variables in the models. Some studies (Lardy et al., 2011; Martin et al., 2007) have attempted to obtain the analytical solution of steady-state pool sizes via managing the climate fluctuations. These methods have used matrix-based analysis but still need to solve several relatively complicated equations. For these reasons, no effective analytical method has been developed to save the spin-up time for global land models.

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An analytical solution is still possible to obtain steady-state pool sizes if we can overcome two obstacles. First, we need to get time-averaged approximations of the time-varying variables, such as environmental scalars and NPP. Since all spin-up uses repeated climate forcing variables to estimate steady states of pool sizes, we can estimate averages of those time-varying variables within one period of the repeated forcing variables. Second, for most carbon-nitrogen coupled models, nitrogen regulates carbon cycle via its influences on photosynthesis and decomposition. Nitrogen pool sizes are usually related with carbon cycle via carbon/nitrogen (C/N) ratios (Gerber et al., 2010; Wang et al., 2010). Nitrogen influences on photosynthesis are fast processes and can be accounted for by short, initial spin-up to reach a steady state of NPP. The C/N ratios at the end of initial spin-up for steady-state NPP can be used to estimate steady-state nitrogen pool sizes. Because the carbon-nitrogen interactions might continue to influence dynamics of the slow pools even after NPP reached steady state, some additional spin-up may be necessary to achieve steady states of all pools.

This study was intended to develop a semi-analytical solution (SAS) to accelerate spin-up of global carbon-nitrogen coupled models. We first discussed biogeochemical principles underlying SAS. SAS becomes permissible because a set of first-order ordinary differential equations can adequately describe carbon transfers within ecosystems over time and be analytically solved to obtain steady-state pool sizes. We applied SAS to the Australian Community Atmosphere Biosphere Land Exchange (CABLE) model and developed a general procedure of SAS for spin-up. There are three key steps of SAS. The first step was a short spin-up to obtain steady-state NPP, averaged environmental factors within one period of repeated forcing variables, and C/N ratios. The second step is to analytically solve the differential equation to calculate steady-state carbon and nitrogen pool sizes. The last step is to make an additional spin-up to meet the steady-state criterion for all pools. We evaluated the computational efficiency of SAS for the carbon-only and the coupled carbon-nitrogen models against the traditional spin-up, applications of SAS to other biogeochemical models, and possible model analyses enabled by SAS.

## 2 Methods

### 2.1 Biogeochemical principles for the semi-analytical solution (SAS)

The semi-analytical solution (SAS) we introduced in this study is built upon principles of biogeochemical cycles in terrestrial ecosystems. The biogeochemical cycle of carbon in an ecosystem is usually initiated with plant photosynthesis, which fixes CO<sub>2</sub> from the atmosphere into an ecosystem. The photosynthetic carbon is partitioned into leaf, root, and woody biomass. Dead biomass becomes litter to metabolic, structural, and coarse woody debris (CWD) litter pools. The litter carbon is partially released to the atmosphere as respired CO<sub>2</sub> and partially converted to soil organic matter (SOM) in fast, slow, and passive pools (Fig. 1). The mean carbon residence time varies greatly among different pools, from several months in leaves and roots to hundreds or thousands of years in woody tissues and SOM (Torn et al., 1997). The carbon processes in terrestrial ecosystem can be mathematically expressed by the following first-order ordinary differential matrix equation (Luo et al., 2003):

$$\frac{dX(t)}{dt} = \xi ACX(t) + BU(t) \quad (1)$$

where  $X(t) = (X_1(t), X_2(t), \dots, X_9(t))^T$  is a  $9 \times 1$  vector describing nine carbon pool sizes in leaf, wood, root, metabolic litter, structural litter, CWD, fast SOM, slow SOM, and passive SOM, respectively, in the Community Atmosphere-Biosphere-Land Exchange

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(CABLE) model (Wang et al., 2011).  $A$  and  $C$  are  $9 \times 9$  matrices given by

$$A = \begin{pmatrix} -1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & -1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & -1 & 0 & 0 & 0 & 0 & 0 & 0 \\ a_{41} & a_{42} & 0 & -1 & 0 & 0 & 0 & 0 & 0 \\ a_{51} & a_{52} & 0 & 0 & -1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 & -1 & 0 & 0 & 0 \\ 0 & 0 & 0 & a_{74} & a_{75} & a_{76} & -1 & 0 & 0 \\ 0 & 0 & 0 & 0 & a_{85} & a_{86} & a_{87} & -1 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & a_{97} & a_{98} & -1 \end{pmatrix} \quad (2)$$

$$C = \text{diag}(c) \quad (3)$$

where  $A$  denotes the carbon transfer matrix, in which  $a_{ij}$  represents the fraction of carbon transfer from pool  $j$  to  $i$ . The  $\text{diag}(c)$  is a  $9 \times 9$  diagonal matrix with diagonal entries given by vector  $c = (c_1, c_2, \dots, c_9)^T$ , components  $c_j$  ( $j = 1, 2, \dots, 9$ ) quantify the fraction of carbon left from pool  $X_j$  ( $j = 1, 2, \dots, 9$ ) after each time step.  $\xi$  is an environmental scalar accounting for effects of soil type, temperature and moisture on carbon decomposition.  $B = (b_1, b_2, b_3, 0, \dots, 0)^T$  represents the partitioning coefficients of the photosynthetically fixed carbon into different pools.  $U$  is the input of fixed carbon via plant photosynthesis. In general, Eq. (1) can adequately summarize  $C$  cycle processes in most land models (Cramer et al., 2001; Parton et al., 1987).

Equation (1) can not be directly solved to obtain the steady-state values of carbon pools because matrix  $A$  and  $B$ , the environmental scalar  $\xi$ , and ecosystem carbon influx  $U(t)$  vary with time and driving variables. Since carbon influx involves fast processes, its steady-state value  $U_{ss}$  can be obtained from short spin-up. All model spin-ups use repeated driving variables in long-term, iterative simulations. Thus it is possible to calculate averaged values of the environmental scalar ( $\bar{\xi}$ ), the carbon transfer ( $\bar{A}$ ) and partitioning ( $\bar{B}$ ) coefficients within one period of repeated driving variables. With  $U_{ss}$  and the mean values of the time-varying variables ( $\bar{\xi}$ ,  $\bar{A}$ , and  $\bar{B}$ ) we can analytically

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calculate the steady-state carbon pool sizes  $X_{ss}$  by letting Eq. (1) equal zero as:

$$X_{ss} = -(\bar{\xi}\bar{A}\bar{C})^{-1}\bar{B}U_{ss} \quad (4)$$

We divided  $X_{ss}$  by C/N ratio in each pool to obtain steady-state nitrogen pool sizes  $N_{ss}$ .

The C/N ratios were the values at the end of the initial spin-up for  $U_{ss}$ . The steady-state carbon and nitrogen pool sizes that are analytically calculated by Eq. (3) may need to be further adjusted with additional spin-up to meet the criterion of steady states for all carbon and nitrogen processes.

Overall, our semi-analytic solution of spin-up consists of three steps: (1) an initial spin-up to obtain steady-state carbon influx  $U_{ss}$ , temporally averaged values of the time-varying variables in Eq. (3) ( $\bar{\xi}$ ,  $\bar{A}$ , and  $\bar{B}$ ), and C/N values; (2) calculation of the steady-state carbon pool sizes  $X_{ss}$  using Eq. (3) and the steady-state N pool sizes  $N_{ss}$  from dividing  $X_{ss}$  by C/N ratios; and (3) additional spin-up to meet the steady-state criteria for all carbon and nitrogen processes.

## 2.2 Model description

We applied the semi-analytic solution of spin-up to the CABLE model, which is one of the land surface models for simulation of biophysical and biochemical processes. Kowalczyk et al. (2006) and Wang et al. (2011) have described the CABLE model in detail. The model includes 5 submodels: radiation, canopy micrometeorology, surface flux, soil and snow, and biogeochemical cycles. The CABLE model calls the radiation submodel first to compute absorption and transmission of both diffuse and direct beam radiation in the two big-leaf canopy and at soil surface (see the details in Wang and Leuning, 1998). The canopy micrometeorology submodel estimates the canopy roughness length, zero-plane displacement height and aerodynamic transfer resistance based on the theory developed by Raupach (1989a, b; 1994) and Raupach et al. (1997). The surface flux submodel uses the absorbed radiation to estimate the water extraction and ground heat flux, which are required in the soil and snow submodel. The

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biogeochemical-cycles submodel is called last to compute the respiration of non-leaf plant tissues, the soil respiration, and the net ecosystem CO<sub>2</sub> exchange.

The biogeochemical sub-model of CABLE evolves from the CASACNP model, which was developed by Wang et al. (2010). It adopted the model structure of carbon processes from the CASA' model (Randerson et al., 1997) and contains coupled carbon, nitrogen, and phosphorus cycles. In this study, phosphorus cycle and its coupling with carbon and nitrogen cycles were not activated. CASACNP model has 9 pools, which include three plant pools (leaf, wood, and root), three litter pools (metabolic litter, structural litter, and coarse woody debris), and three soil pools (microbial biomass, slow and passive soil organic matter) (Fig. 1). There is an additional pool of inorganic nitrogen (NO<sub>3</sub><sup>-</sup> + NH<sub>4</sub><sup>+</sup>) when nitrogen cycle is coupled with carbon cycle. The equations that describe changes in pool size with time have been presented by Wang et al. (2010) and can be represented by Eq. (1). In Eq. (1), parameters  $C$  are set to be constants in the CABLE model while matrix  $A$  and  $B$ , the environmental scalar  $\xi$ , and ecosystem carbon influx  $U(t)$  vary with time and driving variables. In the  $A$  matrix, the carbon transfer coefficients are determined by lignin/nitrogen ratio from plant to litter pools, lignin fraction from litter to soil pools, and soil texture among soil pools. In the  $B$  matrix, the carbon partitioning coefficients of the photosynthetically fixed carbon into plant pools are determined by availabilities of light, water and nitrogen as the carbon allocation scheme described by Friedlingstein et al. (1999). The environmental scalar ( $\xi$ ) regulates the leaf turnover rates by cold and drought stresses on leaf senescence rate, the turnover rates of litter carbon pools via limitations of soil temperature, moisture, and nitrogen availability, and SOM turnover rates by soil temperature, moisture, and texture. The soil texture is spatially fixed in the CABLE model. The soil nitrogen will limit litter decomposition if the gross mineralization is less than the immobilization (Wang et al., 2010). We span up the model for about one hundred of simulation years to obtain the steady-state carbon influx  $U_{ss}$ .

In the CABLE model, the optimal carbon decay rates (the  $C$  matrix in Eq. 1) are pre-set and vary with vegetation types. The vegetation types for each 1° × 1° grid cell in the

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To implement SAS with the CABLE model, we did the following things (Fig. 2):

1. Developing a flow diagram to link carbon pools and fluxes within ecosystems as in Fig. 1.
2. Organizing the linkage between carbon pools and fluxes into carbon transfer matrices  $A$  and  $C$ , and plant carbon partitioning coefficients into vector  $B$ . The preset values of the optimal carbon decay rates were organized into the  $C$  matrix.
3. Figuring out how each element of the time-varying variables ( $A$ ,  $B$  and  $\xi$ ) in the Eq. (1) was determined in the model.
4. Re-coding a section in the model (e.g., the biogeochemical cycle submodel in the CABLE). The re-coding of the model includes 4 steps: (1) setting up a criterion for the steady-state NPP for the initial spin-up; (2) creating new variables to store the mean values of the time-varying parameters; (3) creating equations to calculate the analytical solutions of each pool according to the structures of matrix  $A$ ,  $C$  and vector  $B$ ; and (4) setting up a criterion for the steady-state of the slowest pool for the final spin-up. More details about the re-coding can be found in Supplement.
5. Making an initial spin-up by running the model using repeated meteorological forcing until NPP (or all plant pools) reached steady state ( $U_{ss}$ ). In this study, we run the model until the mean change in NPP over each loop (8 yr) of site simulation at Harvard Forest was smaller than  $10^{-4} \text{ gCm}^{-2}$ . For the global simulation, we run the model until the mean changes in plant carbon pools over each loop (1 yr) was smaller than 0.01 % per year. Meanwhile, the mean values of all the time-varying parameters in Eq. (3) were written to those newly created variables. Those parameters are the steady-state NPP ( $U_{ss}$ ), the mean environmental scalar ( $\bar{\xi}$ ) and matrices of carbon transfer ( $\bar{A}$ ) and partitioning ( $\bar{B}$ ) coefficients within one period of repeated forcing variables, C/N ratios at the end of the initial spin-up.
6. Calculating the analytical solution of the steady states of carbon and nitrogen pools. The steady-state carbon pools were solved by letting carbon influx equal

to efflux for each pool (Eq. 3). Nitrogen pools are obtained by dividing the state-steady carbon pools by the C/N ratios at the end of the initial spin-up.

7. Making the final spin-up by using the analytically solved carbon and nitrogen pools as initial values until the steady-state criterion for the soil carbon pools was met. The steady-state criterion set in this study was that the change in the passive soil carbon pool ( $\Delta C_{\text{pass}}$ ) within each simulation year was smaller than  $0.5 \text{ g C m}^{-2} \text{ yr}^{-1}$  (as one criterion in Thornton and Rosenbloom, 2005). According to the difference in turnover rate, a slower pool needs a longer time to reach steady state during the spin-up. The final spin-up is determined by the dynamic of the slowest carbon pool when the criterion of steady-state soil carbon pools is small enough.

### 3 Results

#### 3.1 Performances of SAS at Harvard Forest

For Harvard Forest, the traditional spin-up method took 1220 and 846 loops to get the steady states of the carbon-only and coupled carbon-nitrogen simulations ( $\Delta C_{\text{pass}} < 0.5 \text{ g C m}^{-2} \text{ yr}^{-1}$ ), respectively. The first step of the SAS was to spin up the model to reach steady state NPP. It took 8 and 42 loops for the carbon-only and coupled carbon-nitrogen simulations, respectively (Fig. 3). After the semi-analytical solution of steady-state values was obtained for all carbon and nitrogen pools, it took another 45 and 89 loops of the carbon-only and coupled carbon-nitrogen simulations, respectively, for the change in the passive soil carbon pool in each loop of simulation ( $\Delta C_{\text{pass}}$ ) less than  $0.5 \text{ g C m}^{-2} \text{ yr}^{-1}$ . In comparison with the traditional spin-up method, the SAS method saved about 95.7 % and 84.5 % of computational time for getting steady states of the carbon-only and coupled carbon-nitrogen simulations, respectively. The differences in steady-state carbon pools between the SAS and traditional spin-up methods

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were small, being 0.85 % and 0.24 % of total ecosystem carbon content for carbon-only and coupled carbon-nitrogen simulations, respectively (Table 1).

The steady-state NPP in the coupled carbon-nitrogen simulation ( $1.45 \text{ gC m}^{-2} \text{ d}^{-1}$ ) was 32.24 % less than that in the carbon-only simulation ( $2.14 \text{ gC m}^{-2} \text{ d}^{-1}$ ). The steady-state value of total ecosystem carbon pool, which represents the ecosystem carbon storage capacity, in the coupled carbon-nitrogen simulation ( $28.83 \text{ kgC m}^{-2}$ ) was 31.44 % lower than that in the carbon-only simulation ( $42.05 \text{ kgC m}^{-2}$ ; Table 1). Although the passive SOM pool determined the spin-up time of CABLE, the slow SOM pool had the largest pool size (Figs. 4 and 5) because the steady-state pool size (i.e., carbon storage capacity) is jointly determined by carbon influx and residence time.

### 3.2 Application SAS to global simulations

The traditional spin-up method spent 2780 and 5099 simulation years for carbon-only and coupled carbon-nitrogen simulation, respectively, before the change in slowest carbon pool met the steady-state criterion ( $\Delta C_{\text{pass}} < 0.5 \text{ gC m}^{-2} \text{ yr}^{-1}$ ; Fig. 6). For SAS, the initial spin-up took 200 simulation years for obtaining steady states of plant carbon pools in the global carbon-only model and 201 yr for the coupled carbon-nitrogen model. With the SAS spin-up method, all carbon pools in the carbon-only model reached steady states ( $\Delta C_{\text{pass}} < 0.5 \text{ gC m}^{-2} \text{ yr}^{-1}$ ) after analytical calculation without any final spin-up (as shown in the black arrow in Fig. 6). In the coupled carbon-nitrogen model, the SAS needed another 483 simulation years to obtain the steady states of all pools (as shown in the gray arrow in Fig. 6) after the analytical calculation. The SAS method saves about 92.4 % and 86.6 % of the computational time for spin-up of the global carbon-only and coupled carbon-nitrogen models to steady states, respectively (Fig. 6).

With the traditional spin-up method, the passive SOM pool continued to decrease after it reached the steady-state criterion ( $\Delta C_{\text{pass}} < 0.5 \text{ gC m}^{-2} \text{ yr}^{-1}$ ) (Fig. 6). Additional thousands of simulation years were needed for the traditional method to reach a steady

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state of the passive SOM pool, which was analytically obtained by the SAS with no time (Fig. 6).

### 3.3 Steady-state pools and fluxes as regulated by N

The capacity of an ecosystem to store carbon is determined by ecosystem carbon in-  
flux and residence times of different pools (Luo et al., 2003). The global mean steady-  
state NPP was greater in the carbon-only ( $0.37 \text{ kg C m}^{-2} \text{ yr}^{-1}$ ) than the coupled carbon-  
nitrogen ( $0.35 \text{ kg C m}^{-2} \text{ yr}^{-1}$ ) simulation (Fig. 7a,b). A larger proportion of photosynthet-  
ically fixed carbon was partitioned to pools with long residence time (e.g., plant wood,  
slow and passive SOM; Fig. 7). The global mean of the ecosystem carbon pool sizes at  
steady state decreased from 14.1 in the carbon-only model to  $12.1 \text{ kg C m}^{-2}$  in the cou-  
pled carbon-nitrogen model (Fig. 7). The mean residence time (as dividing steady-state  
ecosystem carbon pool size by NPP) of the ecosystem carbon pool at steady state in  
the coupled carbon-nitrogen model (34.0 yr) was 10.5 % shorter than that in the carbon-  
only model (38.0 yr). In the CABLE model, large fractions of photosynthate went to plant  
pools, the CWD pool, and the slow SOM pool, but a very small fraction ( $\sim 0.1\%$ ) to the  
passive SOM pool (Fig. 7). Globally, nitrogen processes down-regulated soil carbon  
storage much more substantially at the high (e.g., temperate conifer and mixed forests)  
than the low latitudes (e.g., arid and semi-arid deserts and tropical forest) (Fig. 8).

## 4 Discussion

### 4.1 Computational efficiency

The SAS method saved 92.4% of computational time for spin-up of global carbon-  
only model and 86.6% for the global coupled carbon-nitrogen model in comparison  
with the traditional spin-up method. At the site level in Harvard Forest, SAS saved  
95.7% and 84.5% of computational time for the carbon-only model and coupled

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carbon-nitrogen model, respectively. That means the spin-up with the SAS method can be up to 20 times as fast as the traditional method. The computational efficiency with the SAS method is higher than the best method (the accelerated decomposition method) explored by Thornton and Rosenbloom (2005) for site-level spin-up in an ever-green needle-leaf forest. Lardy et al. (2011) have recently developed an iterative matrix method to accelerate spin-up with a pasture model (PaSim). They reported that their method speeds up the spin-up by up to 20 times as well.

The SAS method can be easily implemented for biogeochemical models at site, regional, and global scales. As described in Sect. 2, implementation of the SAS method involves some light recoding of original models to enable analytical calculation of steady states of carbon and nitrogen pools together with initial and final spin-ups (Text S1). The accelerated decomposition method examined by Thornton and Rosenbloom (2005) has been found difficult to be applied to an age-structured model (Lardy et al., 2011) and requires re-parameterization with a linear scaling factor for each new model. The matrix method developed by Lardy et al. (2011) has been programmed in approximately 500 lines of code, much longer coding than the SAS method in this study (less than 150 lines).

The computational cost for spinning up models strongly depends on the criterion used for steady state. The more precise the criterion (i.e., the smaller value), the longer time the spin-up needs for a model to reach the steady state. With the criterion in this study ( $\Delta C_{\text{pass}} < 0.5 \text{ g C m}^{-2} \text{ yr}^{-1}$ ), the CABLE model was spun up for thousands of simulation years with the traditional method. Additional thousands of simulation years were needed beyond the traditional spin-up to reach a steady state of the passive SOM pool, which was analytically solved by the SAS (Fig. 6). It suggests the SAS method is more efficient to estimate the steady states with high precision than the traditional spin-up method.

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## 4.2 Applications of SAS to various types of biogeochemical models

The developed SAS is generally applicable to most of the terrestrial biogeochemical models, which share a similar structure with the CABLE model in this study (Fig. 1). The CENTURY model (Parton et al., 1993), for example, has 4 plant pools for grassland/crop (shoot, root, grain, standing dead) and 8 for forests (leaf, fine roots, fine branches, dead branches, large wood, dead large wood, coarse roots, dead coarse roots). The dead plant materials go into 4 litter pools (surface structure and metabolic litter, soil structure and metabolic litter) and then 4 soil organic matter (SOM) pools (surface microbes, soil microbes, slow SOM, and passive SOM). Similar to the CABLE model, the decomposition rate of each carbon pool ( $C$  matrix in Eq. 1) in the CENTURY model is preset with an optimal value and modified by soil texture, soil temperature and moisture. The transfer coefficients ( $A$  matrix in Eq. 1) in the CENTURY model are determined by lignin to nitrogen ratio from plant to litter pools, lignin fraction from litter to soil pools, and soil texture from soil to soil pools (Bolker et al., 1998; Parton et al., 1987). If SAS is applied to CENTURY, the flow diagram as Fig. 1 would be needed separately for grassland/crop and forests systems due to different numbers of plant pools. The rest of the SAS procedure as described in Fig. 2 can be exactly applied to CENTURY for spin-up. The Rothamsted carbon (RathC) model has four active SOM pools (decomposable plant material, resistant plant material, microbial biomass, and humified organic matter) and one inert organic matter pool (Coleman and Jenkinson, 1999). Although the biogeochemical models differ on the mechanisms of ecological feedbacks (Hurtt et al., 1998), carbon transfers among these models all follow the first-order decay equations as in Eq. (1). Thus, the SAS procedure can be applied to spin-up of the RathC model.

An application of SAS to spin up other coupled carbon-nitrogen models may require additional steps as different models couple the nitrogen cycle to the carbon cycle in different ways. For example, the Princeton-Geophysical Fluid Dynamic Laboratory (GFDL) LM3V model uses one arbitrary nitrogen storage (or buffering) pool in plants to avoid short-term switches between N sufficiency and limitation in plants (Gerber et al.,

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2010). The optimum size of the nitrogen buffering pool equals to the total nitrogen losses from living pools (leaf, root, and sapwood) over one year. The rest of the model structure of LM3V for carbon and nitrogen transfers within the ecosystem is similar to CABLE. Thus, the SAS can be applied to the LM3V model with one additional step to calculate the steady state of the optimal nitrogen buffering pool from the analytically solved steady-state live biomass carbon pools. Similarly, the CLM-CN and O-CN models have nitrogen buffering pools (Thornton and Zimmermann, 2007; Zaehle et al., 2010). Therefore, it is necessary to determine how each model couples the nitrogen cycle with carbon cycle before we apply the SAS procedure to spin-up of the coupled carbon-nitrogen models.

In principle, The SAS method can be applicable to spin up ocean biogeochemical models. Most of the ocean biogeochemical models use the traditional method for spin-up for more than 10 000 yr to reach steady state (Schmittner et al., 2008). A key issue of applying the SAS method to spin up ocean biogeochemical models is whether or not the ocean carbon and nitrogen cycles can be mathematically described by a matrix form similarly as in Eq. (1). If yes, this SAS method can facilitate high resolution analysis of or ensemble simulations with ocean models in the future.

### 4.3 SAS-facilitated model analyses

Land models have been developed by the modeling community in the past two decades to predict future states of ecosystems and climate. Model intercomparison has recently become a popular method to improve our understanding of the land model performances (Friedlingstein et al., 2006; Johns et al., 2011). A common protocol for many of the model intercomparison projects is to spin up all the involved models to steady states before global change scenarios are applied to project future changes (Schwalm et al., 2010). Presently, individual models use their own methods for spin-up, often with different criteria of steady states. Standardized spin-up with a fast, easily implemented method can help reduce uncertainties in model-model and model-data intercomparison studies. The SAS method has the potential to serve those projects in such a capacity.

Accelerated spin-up reduces computational overhead for modeling analysis and make some computationally costly analysis feasible. For example, model parameters usually represent the average physiological properties of plant functional types or mean soil attributes. Most of these parameters in the model are assigned values based on relatively few field and/or laboratory observations (Stockli et al., 2008). More and more databases have been developed to indicate that key plant physiological properties, such as leaf traits (GLOPNET; Reich et al., 2007), carboxylation capacity (V<sub>cmax</sub>; Kattge et al., 2009), and biomass allocation (Poorter et al., 2012), greatly vary among plants of different species at different locations. Similarly, properties of soil processes, such as temperature sensitivity of soil respiration (Peng et al., 2009), also greatly vary over time and space. The natural variations in key plant and soil properties can be adequately represented only by probability distributions of parameters, which would propagate in the land models to generate uncertainties in model projections (Weng and Luo, 2011; Xu et al., 2006). The model projection uncertainties can be quantified through ensemble analysis. However, such ensemble analysis of land models against parameter variations is computationally not feasible because each ensemble element requires at least spin-up once up to thousand and even million times for one ensemble analysis. Without the ensemble analysis against parameter variations at regional or global scales, uncertainties in model projections can not be fully assessed. Fast spin-up methods, such as SAS, should enable the ensemble analysis that is impossible with traditional methods.

The SAS method not only accelerates spin-up with high computational efficiency but also possibly offers an analytical framework for comparative study of structural differences in modeled ecosystem carbon storage capacity. The sum of steady-state carbon pool sizes within one ecosystem obtained from analytical solution of Eq. (3) represents the ecosystem carbon storage capacity (Luo et al., 2003). At the site of Harvard Forest the carbon storage capacity was simulated to be 42.05 kgC m<sup>-2</sup> with the carbon-only model. The capacity was down-regulated by nitrogen processes to be 28.83 kgC m<sup>-2</sup> in the coupled carbon-nitrogen model. As indicated by Eq. (3), the carbon storage capac-

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ity is determined by carbon influx ( $U_{ss}$ ) and residence time [ $(\bar{\xi}\bar{A}C)^{-1}\bar{B}$ ]. The 45.86 % decrease in carbon storage largely resulted from the NPP decrease in the coupled carbon-nitrogen model NPP ( $1.45 \text{ gCm}^{-2} \text{ d}^{-1}$ ) in comparison with that in the carbon-only model ( $2.14 \text{ gCm}^{-2} \text{ d}^{-1}$ ). Indeed, the steady-state solution of Eq. (3) can be used to analyze the determinants of ecosystem carbon storage capacity. The steady state carbon pool in the biogeochemical model is determined by the carbon influx and residence times (reverses of turnover times). The temporal and spatial variations of the residence times are controlled by environmental scalars to describe effects of temperature, moisture, and soil types on carbon transfer processes. The environmental scalars usually are substantially different among models. For example, the temperature scalar on carbon decay rates is a generalized Poisson function in the CENTURY model (Parton et al., 1994) while a simple exponential equation in the Terrestrial Ecosystem Model (TEM) model (McGuire et al., 1997). Analyses of carbon cycles in an analytical framework, such as Eq. (3), can help compare structural differences in determining ecosystem carbon storage capacity among biogeochemical models.

## 5 Conclusions

We developed a new method – semi-analytical solution (SAS) – to accelerate the spin-up of a process-oriented biogeochemical model to steady states. The SAS described in this study mainly contains 3 steps: (1) making an initial spin-up to get steady-state values of photosynthetic carbon input and plant pools; (2) calculating the semi-analytical solution of the steady-state pool sizes; (3) having a final spin-up to meet the criterion of steady states for slowest pools (Fig. 2). For spin-up of the carbon-only model, the SAS method can save about 95.7 % and 92.4 % computational time for site-level and global simulations, respectively. The efficiency of the SAS method decreased for coupled carbon-nitrogen simulations, but it still resulted in 84.5 % and 86.6 % reduction in computational cost for site-level and global simulation, respectively. We suggest that the SAS described in this study would be a candidate for solving the “spin-up problem”

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in global models. This method enables many modeling analyses, such as ensemble analysis with regard to parameter variability, which otherwise are impossible with computationally costly spin-up methods.

**Supplementary material related to this article is available online at:**

**<http://www.geosci-model-dev-discuss.net/5/803/2012/gmdd-5-803-2012-supplement.pdf>.**

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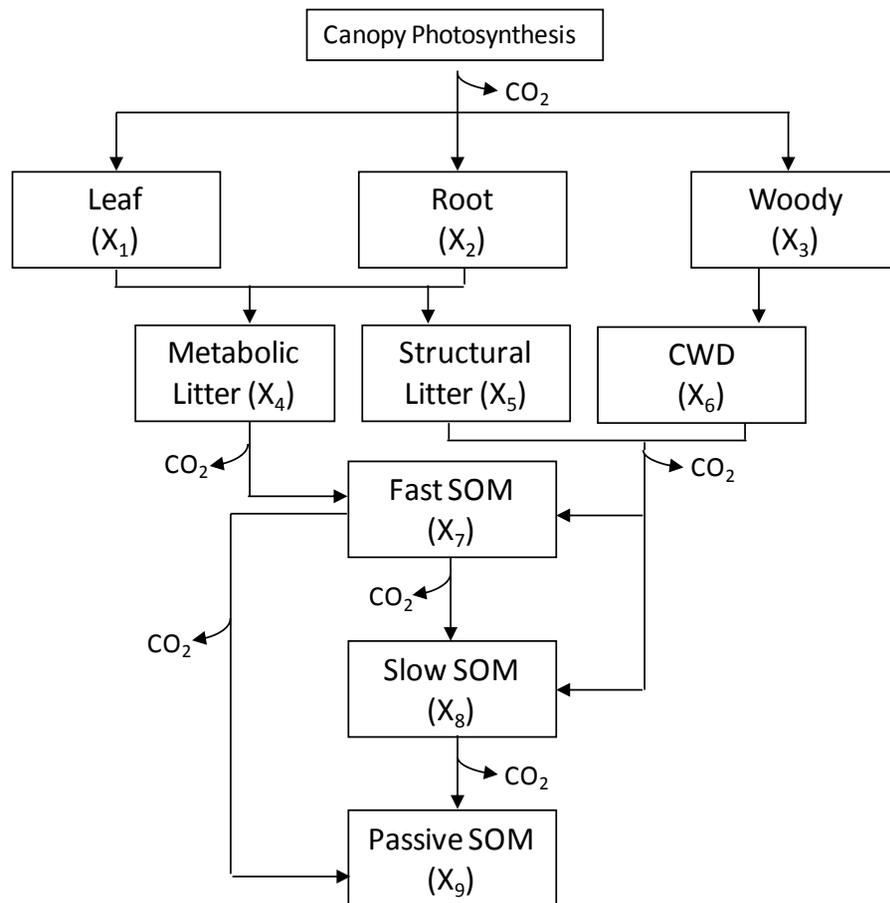
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**Table 1.** Means steady-state values ( $\text{kg C m}^{-2}$ ) of all pools and their total value ( $C_{\text{tot}}$ ) from spin-up with traditional and the SAS methods, and the corresponding relative differences ( $\Delta C$ ; %) for multiple carbon pools. Mlitter, metabolic litter pool; Slitter, structure litter pool; FSOM, fast SOM; SSOM, slow SOM; PSOM, passive SOM.

|                         | Leaf | Woody | Root | Mlitter | Slitter | CWD  | FSOM | SSOM  | PSOM | $C_{\text{tot}}$ |
|-------------------------|------|-------|------|---------|---------|------|------|-------|------|------------------|
| Carbon-only             |      |       |      |         |         |      |      |       |      |                  |
| Traditional             | 0.14 | 8.61  | 3.85 | 0.13    | 0.91    | 1.78 | 0.83 | 16.28 | 9.53 | 42.05            |
| SAS                     | 0.14 | 8.61  | 3.85 | 0.13    | 0.91    | 1.78 | 0.83 | 16.28 | 9.88 | 42.41            |
| $\Delta C$ (%)          | 0.00 | 0.00  | 0.00 | 0.00    | 0.00    | 0.00 | 0.00 | 0.00  | 3.67 | 0.85             |
| Coupled carbon-nitrogen |      |       |      |         |         |      |      |       |      |                  |
| Traditional             | 0.10 | 5.90  | 2.60 | 0.10    | 0.59    | 1.24 | 0.57 | 11.21 | 6.52 | 28.83            |
| SAS                     | 0.10 | 5.90  | 2.60 | 0.10    | 0.59    | 1.24 | 0.57 | 11.20 | 6.68 | 28.96            |
| $\Delta C$ (%)          | 0.00 | 0.00  | 0.00 | 0.00    | 0.00    | 0.00 | 0.00 | -0.09 | 2.45 | 0.24             |

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**Fig. 1.** Diagram of the carbon processes of CASACNP model on which model Eq. (1) is based. SOM stands for soil organic matter.

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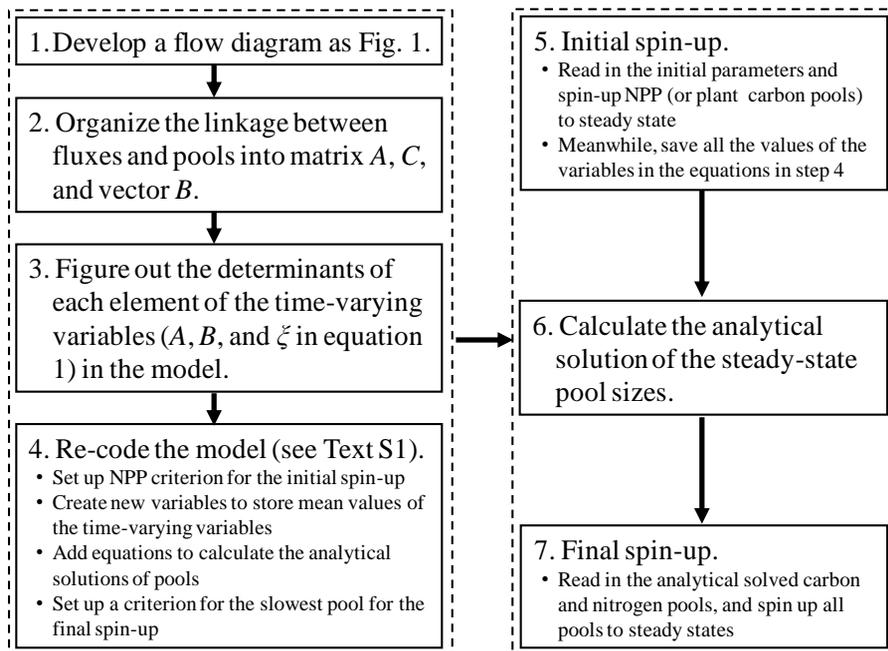
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**Fig. 2.** The spin-up strategies of the spin-up method with the semi-analytical solution (SAS) used in this study.

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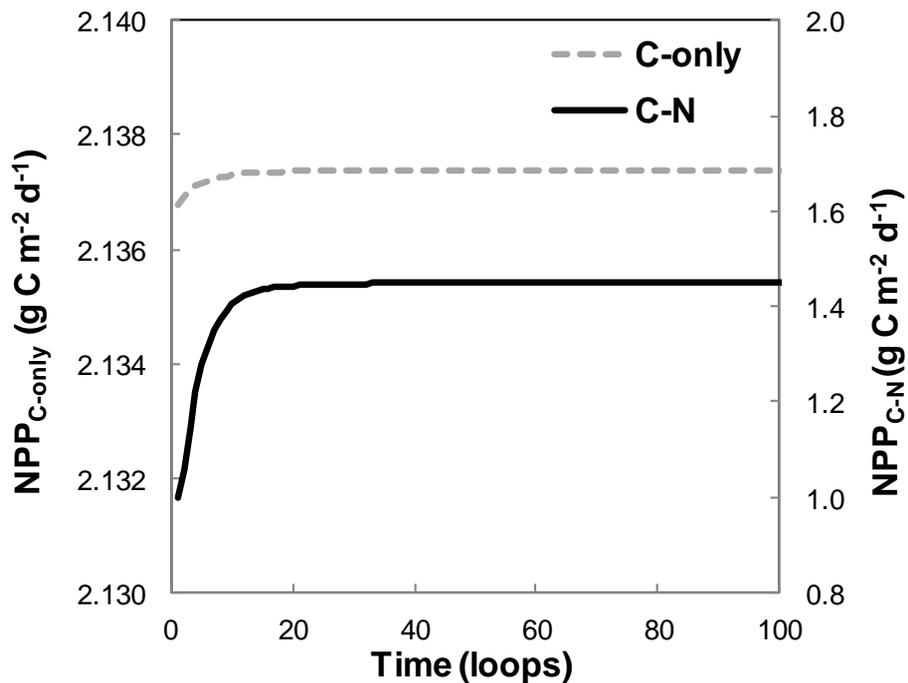
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**Fig. 3.** Dynamics of NPP in the carbon-only (C-only;  $\text{NPP}_{\text{C-only}}$ ) and coupled carbon-nitrogen (C-N;  $\text{NPP}_{\text{C-N}}$ ) simulations with Harvard Forest data.

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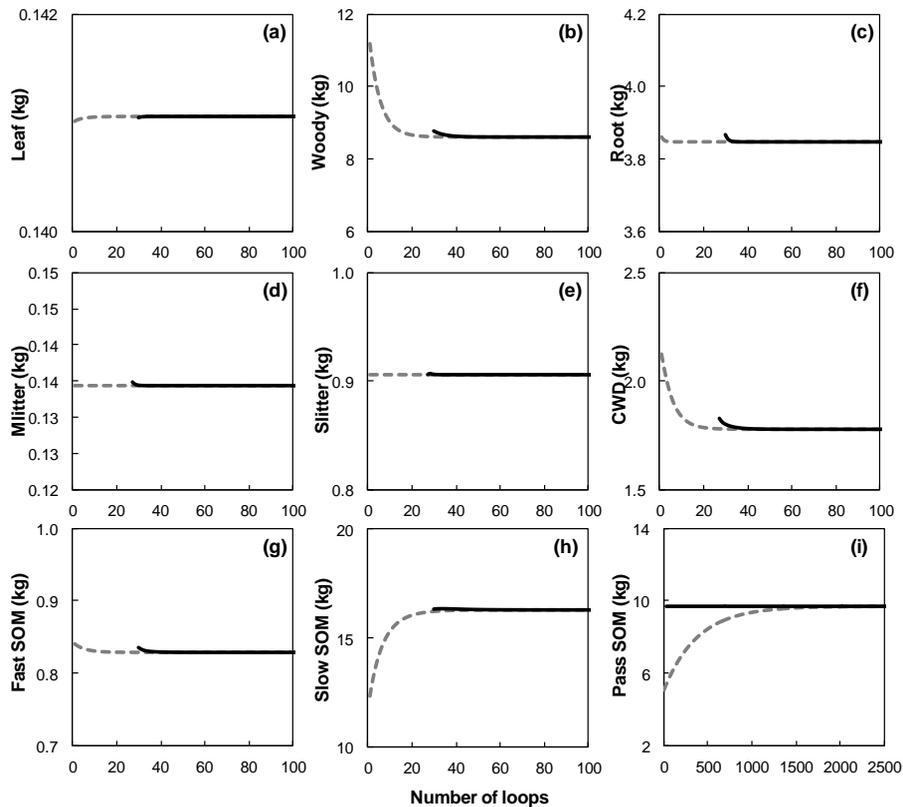
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**Fig. 4.** Carbon-only simulations: carbon state trajectories for all carbon pools at Harvard Forest site with different spin-up methods from traditional procedure (dotted grey lines) and with the SAS (straight black lines).

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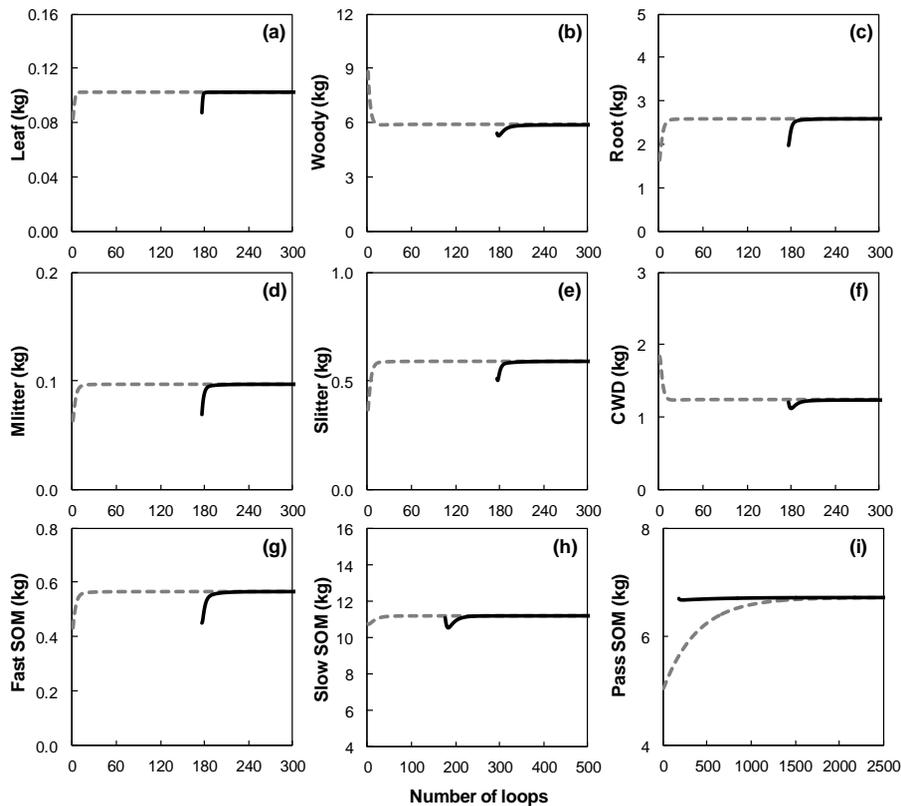
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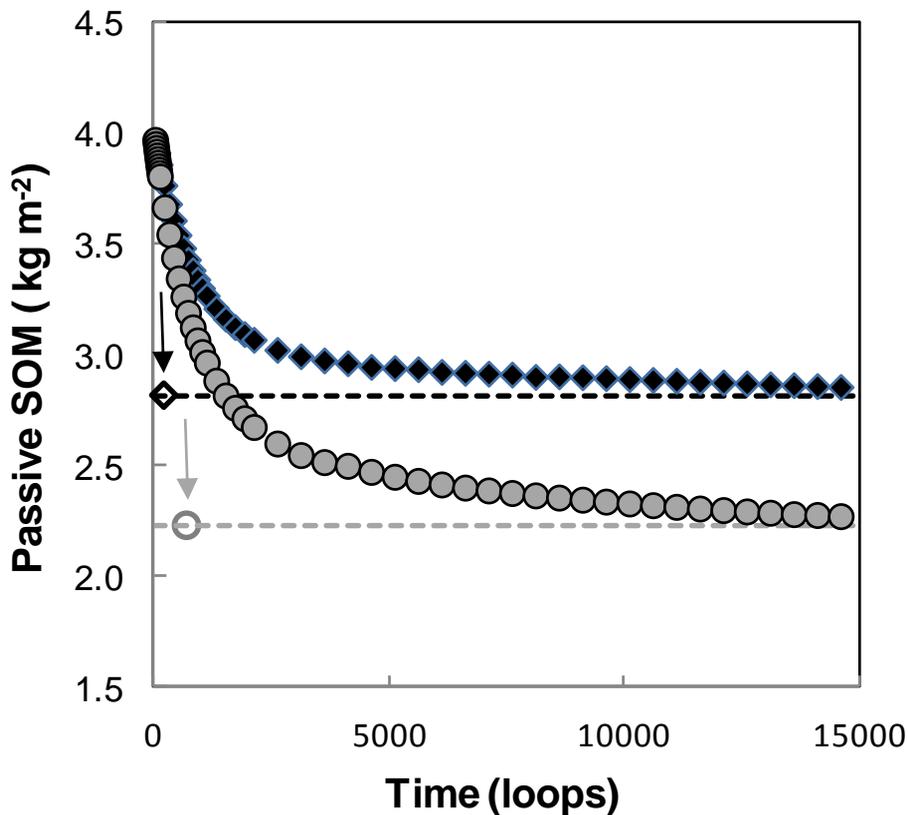
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**Fig. 5.** Coupled carbon-nitrogen simulations: carbon state trajectories for all carbon pools at Harvard Forest site with different spin-up methods from traditional procedure (dotted grey lines) and with the SAS (straight black lines).



**Fig. 6.** Dynamics of global mean passive soil carbon pool in carbon-only (filled black diamond) and coupled carbon-nitrogen (filled gray circle) simulations with traditional method and the SAS framework (dotted lines). The arrows and blank symbols shows the times and values estimated with the SAS framework for carbon-only (black) and coupled carbon-nitrogen coupled (gray) simulations.

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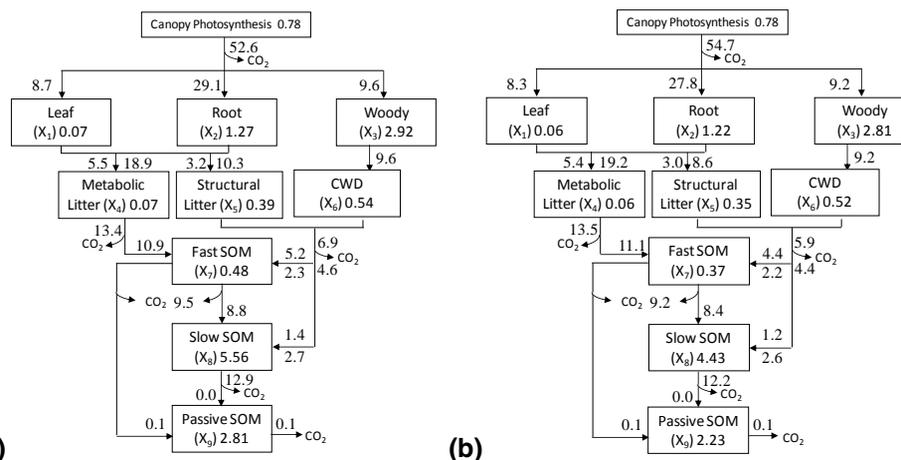
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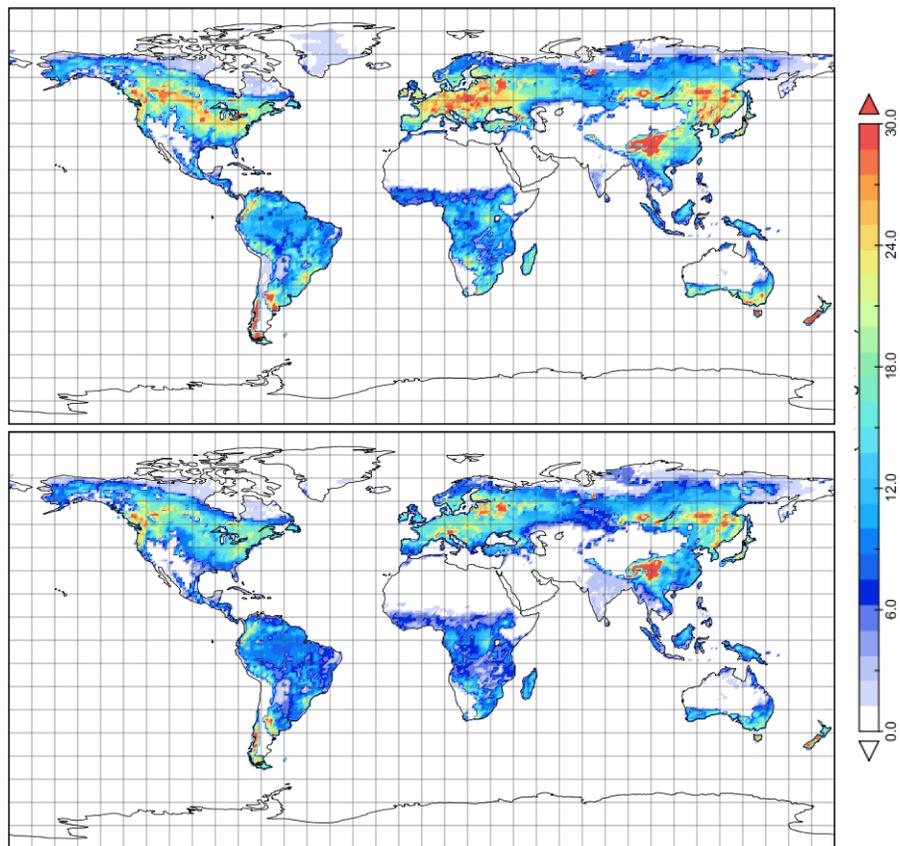
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**Fig. 7.** Structure of CASACNP model for **(a)** carbon-only and **(b)** coupled carbon-nitrogen simulations on which Eq. (1) is based. Fraction of carbon that flows through differential pathways (the numbers near the arrows) is partitioned to the 9 pools. The numbers in the boxes are steady states of ecosystem carbon influx ( $\text{kg C m}^{-2} \text{ yr}^{-1}$ ) and pool size ( $\text{kg C m}^{-2}$ ). The fraction to plant pools is determined by partitioning coefficients in the vector  $B$  in Eq. (1). The fractions to litter and soil pools are determined by the transfer coefficient matrix  $A$  in Eq. (1). The values of  $B$  and  $A$  are the global mean values at steady states. SOM stands for soil organic matter. Upper and bottom values near the arrows represent fraction from structure litter and coarse wood detritus (CWD), respectively.



**Fig. 8.** Global distributions of soil carbon density ( $\text{kg C m}^{-2}$ ) at steady state simulated by the carbon-only (upper panel) or coupled carbon-nitrogen (upper panel) models.

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