

1 **Competition alters predicted forest carbon cycle responses to nitrogen availability and**
2 **elevated CO₂: simulations using an explicitly competitive, game-theoretic vegetation**
3 **demographic model**

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14 **Key words:** Allocation; Biome Ecological strategy simulator (BiomeE); Competitively-optimal
15 strategy; Game theory; Nitrogen cycle

16

17 **Abstract:** Competition is a major driver of carbon allocation to different plant tissues (e.g.
18 wood, leaves, fine roots), and allocation, in turn, shapes vegetation structure. To improve their
19 modeling of the terrestrial carbon cycle, many Earth system models now incorporate vegetation
20 demographic models (VDMs) that explicitly simulate the processes of individual-based
21 competition for light and soil resources. Here, in order to understand how these competition
22 processes affect predictions of the terrestrial carbon cycle, we simulate forest responses to
23 elevated CO₂ along a nitrogen availability gradient using a VDM that allows us to compare fixed
24 allocation strategies versus competitively-optimal allocation strategies. Our results show that
25 competitive and fixed strategies predict opposite fractional allocation to fine roots and wood,
26 though they predict similar changes in total NPP along the nitrogen gradient. The competitively-
27 optimal allocation strategy predicts decreasing fine root and increasing wood allocation with
28 increasing nitrogen, whereas the fixed allocation strategy predicts the opposite. Although
29 simulated plant biomass at equilibrium increases with nitrogen due to increases in photosynthesis
30 for both allocation strategies, the increase in biomass with nitrogen is much steeper for
31 competitively-optimal allocation due to its increased allocation to wood. The qualitatively
32 opposite fractional allocation to fine roots and wood of the two strategies also impacts the effects
33 of elevated [CO₂] on plant biomass. Whereas the fixed allocation strategy predicts an increase in
34 plant biomass under elevated [CO₂] that is approximately independent of nitrogen availability,
35 competition leads to higher plant biomass response to elevated [CO₂] with increasing nitrogen
36 availability. Our results indicate that the VDMs that explicitly include the effects of competition
37 for light and soil resources on allocation may generate significantly different ecosystem-level
38 predictions of carbon storage than those that use fixed strategies.

39

40 **1 Introduction**

41 Allocation of assimilated carbon to different plant tissues is a fundamental aspect of plant growth
42 and profoundly affects terrestrial ecosystem biogeochemical cycles (Cannell and Dewar, 1994;
43 Lacoïnte, 2000). Ecologically, allocation represents an evolutionarily-honed “strategy” of plants
44 that use limited resources and compete with other individuals and consequently drives
45 successional dynamics and vegetation structure (De Kauwe et al., 2014; DeAngelis et al., 2012;
46 Haverd et al., 2016; Tilman, 1988). Biogeochemically, allocation links plant physiological
47 processes, such as photosynthesis and respiration, to biogeochemical cycles and carbon storage
48 of ecosystems (Bloom et al., 2016; De Kauwe et al., 2014). Thus, correctly modeling allocation
49 patterns is critical for correctly predicting terrestrial carbon cycles and Earth system dynamics.

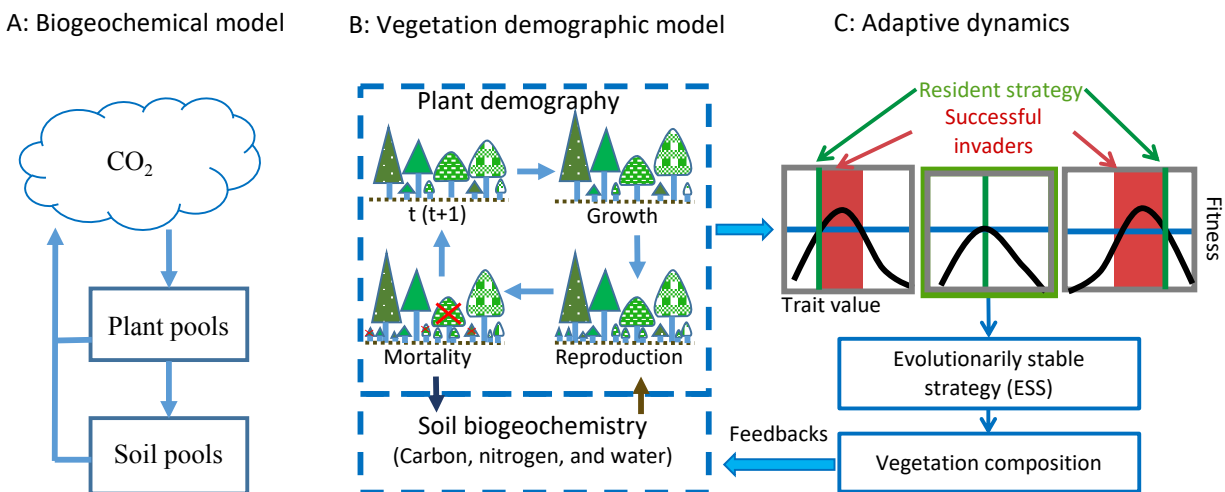
50 In current Earth System Models (ESMs), the terrestrial carbon cycle is usually simulated by
51 pool-based compartment models that simulate ecosystem biogeochemical cycles as lumped pools
52 and fluxes of plant tissues and soil organic matter (Fig. 1: A) (Emanuel and Killough, 1984;
53 Eriksson, 1971; Parton et al., 1987; Randerson et al., 1997; Sitch et al., 2003). In these models,
54 the dynamics of carbon can be described by a linear system of equations (Koven et al., 2015;
55 Luo et al., 2001; Luo and Weng, 2011; Sierra and Mueller, 2015; Xia et al., 2013):

$$56 \frac{dx}{dt} = AX + BU \quad (\text{Eq. 1})$$

57 where X is a vector of ecosystem carbon pools, U is carbon input (i.e., Gross Primary Production,
58 GPP), B is the vector of allocation parameters to autotrophic respiration and plant carbon pools
59 (e.g., leaves, stems, and fine roots), and A is a matrix of carbon transfer and turnover. In this
60 system, carbon dynamics are defined by carbon input (U), allocation (B), and residence time and
61 transfer coefficients (A). The allocation schemes (B) are thus embedded in a linear system, or

62 quasi-linear system if the allocation parameters in B are a function of carbon input (U) or plant
 63 carbon pools (X).

64 The modeling of allocation in this system (i.e., the parameters in vector B) is usually based
 65 on plant allometry, biomass partitioning, and resource limitation (De Kauwe et al., 2014;
 66 Montané et al., 2017). The allocation parameters are either fixed ratios to leaves, stems, and
 67 roots, which may vary among plant functional types (e.g., CENTURY, Parton et al., 1987; TEM,
 68 Raich et al., 1991; CASA, Randerson et al., 1997) or are responsive to climate and soil
 69 conditions as a way to phenomenologically mimic the shifts in allocation that are empirically
 70 observed or hypothesized (e.g., CTEM, Arora and Boer, 2005; ORCHIDEE, Krinner et al., 2005;
 71 LPJ, Sitch et al., 2003). These modeling approaches either assume that vegetation is equilibrated
 72 (fixed ratios) or average the responses of plant types to changes in environmental conditions as a
 73 collective behavior. Thus, the carbon dynamics in these models can be constrained by selecting
 74 appropriate parameters of allocation, turnover rates, and transfer coefficients to fit the
 75 observations (Friend et al., 2007; Hoffman et al., 2017; Keenan et al., 2013).



76
 77 **Figure 1 Hierarchical structure of vegetation models**

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79 To predict transient changes in vegetation structure and composition in response to climate
80 change, vegetation demographic models (VDMs) that are able to simulate transient population
81 dynamics are incorporated into ESMs (Fisher et al., 2018; Scheiter and Higgins, 2009).
82 Generally, VDMs explicitly simulate demographic processes, such as plant reproduction, growth,
83 and mortality, to generate the dynamics of populations (Fig. 1: B). To speed computations and
84 minimize complexity, groups of individuals are usually modeled as cohorts. With multiple
85 cohorts and PFTs, VDMs can bring plant functional diversity and adaptive dynamics into ESMs
86 when explicitly simulating individual-based competition for different resources and vegetation
87 succession and thus predict dominant plant traits changes with environmental conditions and
88 ecosystem development (Scheiter et al., 2013; Scheiter and Higgins, 2009; Weng et al., 2015).

89 The combinations of plant traits represent the competition strategies at different stages of
90 ecosystem development. Evolutionarily, a strategy that can outcompete all other strategies in the
91 environment created by itself will be dominant. This strategy is called an evolutionarily stable
92 strategy or a competitively-optimal strategy (McGill and Brown, 2007). In VDMs,
93 competitively-optimal strategies can therefore be reasonably predicted based on the costs and
94 benefits of different strategies (i.e., combinations of plant traits) through their effects on
95 demographic processes (i.e., fitness) and ecosystem biogeochemical cycles (Fig. 1:C) (e.g.,
96 Farrior et al., 2015; Weng et al., 2015).

97 The dynamics of plant traits can substantially change predictions of ecosystem
98 biogeochemical dynamics since they change the key parameters of vegetation physiological
99 processes and soil organic matter decomposition (e.g., Dybzinski et al., 2015; Farrior et al.,
100 2015; Weng et al., 2017). Therefore, the key parameters that are used to estimate carbon
101 dynamics in the linear system model (Eq. 1), such as allocation (B) and residence times in

102 different carbon pools (matrix A , which includes coefficients of carbon transfer and turnover
103 time) become functions of competition strategies that vary with environment and carbon input. In
104 addition, the turnover of vegetation carbon pools becomes a function of allocation, leaf
105 longevity, fine root turnover, and tree mortality rates, which change with vegetation succession
106 and the most competitive plant traits. These changes make the system nonlinear and can lead to
107 large biases within the framework of the compartmental pool-based models as represented by Eq.
108 (1) (Sierra et al., 2017; Sierra and Mueller, 2015). Because of the high complexity associated
109 with demographic and competition processes, the model predictions are usually sensitive to the
110 parameters in these processes and are of high uncertainty (e.g., Pappas et al., 2016).

111 In contrast to their implementation in the more complicated VDMs discussed above,
112 models of competitively-dominant plant strategies using much simpler model structures and
113 assumptions can sometimes be solved analytically (Dybzinski et al., 2011, 2015; FARRIOR et al.,
114 2013, 2015). Although simplified, such models can pin-point the key processes that improve the
115 predictive power of simulation models (Dybzinski et al., 2011; FARRIOR et al., 2013, 2015),
116 allowing them to help researchers formulate model processes and understand the simulated
117 ecosystem dynamics in ESMs. For example, the analytical model derived by FARRIOR et al. (2013)
118 that links interactions between ecosystem carbon storage, allocation, and water stress at elevated
119 CO₂ sheds light on the otherwise inscrutable processes leading to varied soil water dynamics in a
120 land model coupled with an VDM (Weng et al., 2015). Recognizing the benefit, Weng et al.
121 (2017) included both a simplified analytical model and a more complicated VDM to understand
122 competitively optimal leaf mass per area, competition between evergreen and deciduous plant
123 functional types, and the resulting successional patterns.

124 In this study, we use a stand-alone simulator derived from the LM3-PPA model (Weng et
125 al., 2017, 2015) to show how forests respond to elevated CO₂ and nitrogen availability via
126 different competitively-optimal allocation strategies. The model is an individual-based
127 vegetation demographic model, whose vegetation demographic scheme has been coupled into the
128 land model of the Geophysical Fluid Dynamical Laboratory's Earth System Model (Shevliakova
129 et al., 2009; Weng et al., 2015) and NASA Goddard Institute for Space Study's Earth system
130 model, ModelE (Schmidt et al., 2014). Using this model, we simulate the shifts in competitively
131 optimal allocation strategies in response to elevated CO₂ at different nitrogen levels based on
132 insights from the analytical model derived by Dybzinski et al. (2015). Dybzinski et al.'s (2015)
133 model predicts that increases in carbon storage at elevated CO₂ relative to storage at ambient
134 CO₂ are largely independent of total nitrogen because of an increasing shift in carbon allocation
135 from long-lived, low-nitrogen wood to short-lived, high-nitrogen fine roots under elevated CO₂
136 with increasing nitrogen availability. Here, we analyze the simulated ecosystem carbon cycle
137 variables (gross and net primary production, allocation, and biomass) of separate mono- and
138 polyculture model runs. In the monoculture runs, ecosystem properties are the result of the
139 prescribed allocation strategies of a given PFT, analogous to the fixed allocation schemes of
140 most VDMs (see above). In the polyculture runs, competition between the different allocation
141 strategies results in succession and the eventual dominance of the most competitive allocation
142 strategy for a given nitrogen availability and CO₂ level. Since everything else in the model is
143 identical, we are able to compare the predictions of single **fixed strategies** with **competitively-**
144 **optimal allocation strategies** by comparing the ecosystem properties of these two types of runs.

145 2 Methods and Materials

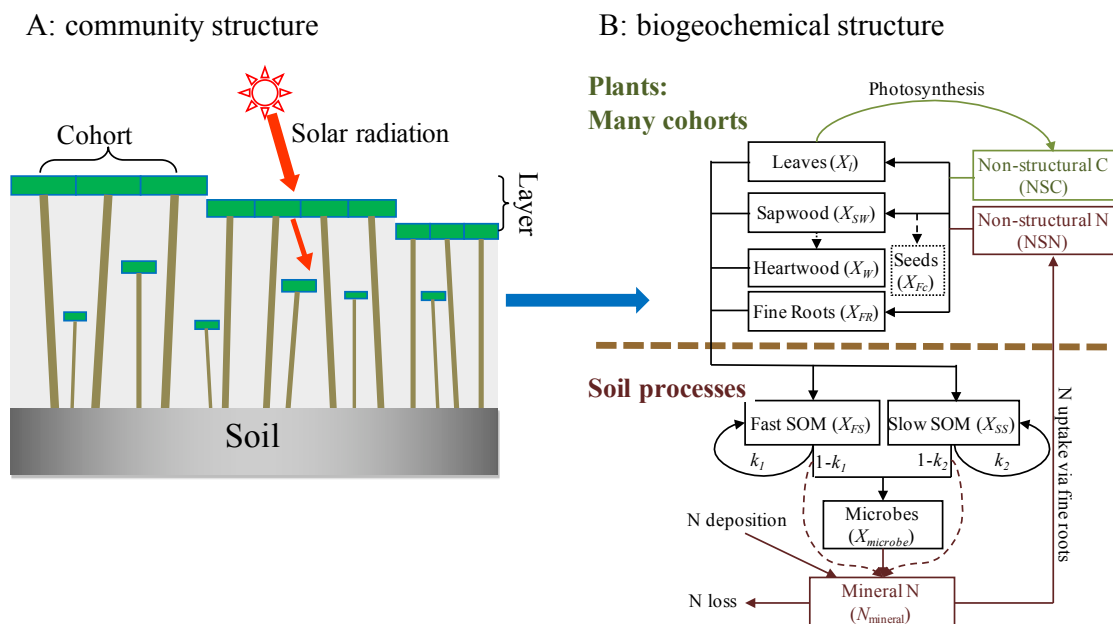
146 2.1 BiomeE model overview

147 We used a stand-alone ecosystem simulator (Biome Ecological strategy simulator,
148 BiomeE) to conduct simulation experiments. BiomeE is derived from the version of LM3-PPA
149 used in Weng *et al.* (2017). In this version, we simplified the processes of energy transfer and
150 soil water dynamics of LM3-PPA (Weng et al., 2015) but still retained the key features of plant
151 physiology and individual-based competition for light, soil water, and, via the decomposition of
152 soil organic matter, nitrogen (Fig. 2). In this model, individual trees are represented as sets of
153 *cohorts* of similar size trees and are arranged in different vertical canopy layers according to
154 their height and crown area following the rules of the Perfect Plasticity Approximation (PPA)
155 model (Strigul et al., 2008). Sunlight is partitioned into these canopy layers according to Beer's
156 law. Thus, a key parameter for light competition, critical height, is defined; all the trees above
157 this context-dependent height get full sunlight and all trees below this height are shaded by the
158 upper layer trees.

159 Each tree consists of seven pools: leaves, fine roots, sapwood, heartwood, fecundity
160 (seeds), and non-structural carbohydrates and nitrogen (NSC and NSN, respectively) (Fig. 2: b).
161 The carbon and nitrogen in plant pools enter the soil pools with the mortality of individual trees
162 and the turnover of leaves and fine roots. There are three soil organic matter (SOM) pools for
163 carbon and nitrogen: fast-turnover, slow-turnover, and microbial pools, along with a mineral
164 nitrogen pool for mineralized nitrogen in soil. The simulation of SOM decomposition and
165 nitrogen mineralization is based on the models of Gerber *et al.* (2010) and Manzoni *et al.* (2010)
166 and described in detail in Weng et al. (2017). The decomposition rate of a SOM pool is
167 determined by the basal turnover rate together with soil temperature and moisture. The nitrogen

168 mineralization rate is a function of decomposition rate and the C:N ratio of the SOM. Microbes
 169 must consume more carbon in the high C:N ratio SOM pool to get enough nitrogen and must
 170 release excessive nitrogen in the low C:N ratio SOM pool to get enough carbon for energy
 171 (Weng *et al.* 2017).

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Figure 2. Model structure of BiomeE

175 Panel A: vegetation structure: trees organize their crowns into canopy layers according to both
 176 their height and their crown area following the rules of the PPA model, which mechanistically
 177 models light competition. Panel B: Biogeochemical structure and compartmental pools. The
 178 green, brown, and black lines are the flows of carbon, nitrogen, and coupled carbon and nitrogen,
 179 respectively. The green box is for carbon only. The brown boxes are N pools. The black boxes
 180 are for both carbon and nitrogen pools, where X can be C (carbon) and N (nitrogen). The C:N
 181 ratios of leaves, fine roots, seeds, and microbes are fixed. The C:N ratios of woody tissues, fast
 182 soil organic matter (SOM), and slow SOM are flexible. Only one tree's C and N pools are shown
 183 in this figure. The model can have multiple cohorts of trees, which share the same pool structure.
 184 The dashed line separates the plant and soil processes.

185

Table 1 Model parameters

Symbol	Definition	Unit	Default value
α_Z	Parameter of tree height	$\text{m m}^{-0.5}$	36
θ_Z	Diameter exponent of tree height	-	0.5
Λ	Taper factor	-	0.75
ρ_W	Wood density	kgC m^{-3}	300
α_C	Parameter of crown area	$\text{m m}^{-1.5}$	150
θ_C	Diameter exponent of crown area	-	1.5
l^*	Target crown leaf area layers (crown leaf area index)	$\text{m}^2 \text{m}^{-2}$	3.5
σ	Leaf mass per unit area	kgC m^{-2}	0.14
γ	Specific root area, calculated from root radius and density	$\text{m}^2 \text{kgC}^{-1}$	34.5
φ_{RL}	Ratio of target fine root area to target leaf area	$\text{m}^2 \text{m}^{-2}$	Vary with PFTs
α_{CSA}	ratio of target sapwood cross-sectional area to target leaf area	$\text{m}^2 \text{m}^{-2}$	0.2E-4
$f_{U,max}$	Maximum mineral N absorption rate	hour^{-1}	0.5
K_{FR}	Root biomass at which the N-uptake rate is half of the maximum	kgC m^{-2}	0.3
$CN_{L,0}$	Target C:N ratio of leaves	kgC kgN^{-1}	76.5(Function of LMA)
$CN_{FR,0}$	Target C:N ratio of fine roots	kgC kgN^{-1}	60
$CN_{W,0}$	Target C:N ratio of wood	kgC kgN^{-1}	350
$CN_{F,0}$	Target C:N ratio of seeds	kgC kgN^{-1}	20
f_1	Supply rate of NSC and NSN	-	1/(3*365)
f_2	Maximum fraction of NSC and NSN used for growth in a day	-	0.02
$f_{LFR,max}$	Maximum fraction of available carbon allocated to leaves and fine roots	-	0.85
v	Fraction of carbon converted to seeds	-	0.1
$r_{D/S}$	Nitrogen-limiting factor	-	Solve by the model (Eqs 9 and 10)

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Plant growth and reproduction are driven by the carbon assimilation of leaves via

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photosynthesis, which is in turn dependent on water and nitrogen uptake by fine roots. The

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photosynthesis model is identical to that of LM3-PPA (Weng et al., 2015), which is a simplified

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version of Leuning model (Leuning et al., 1995). This model first calculates photosynthesis rate,

192 stomatal conductance, and water demand of the leaves of each tree (cohort) in the absence of soil
 193 water limitation. Then, it calculates available water supply and reduces the demand-based
 194 assimilation and stomatal conductance accordingly if water supply is less than water demand.
 195 Assimilated carbon enters into the NSC pool and is subsequently used for respiration, growth,
 196 and reproduction. (Please see Supplementary Information I for details of this model).

197 Empirical allometric equations relate woody biomass (including coarse roots, bole, and
 198 branches), crown area, and stem diameter. The individual-level dimensions of a tree, *i.e.*, height
 199 (Z), biomass (S), and crown area (A_{CR}) are given by empirical allometries (Dybzinski et al.,
 200 2011; Farrior et al., 2013):

$$\begin{aligned} Z(D) &= \alpha_Z D^{\theta_Z} \\ S(D) &= 0.25\pi\lambda\rho_W\alpha_Z D^{2+\theta_Z} \\ A_{CR}(D) &= \alpha_c D^{\theta_c} \end{aligned} \tag{Eq. 2}$$

201 where Z is tree height, D is tree diameter, S is total woody biomass carbon (including bole,
 202 coarse roots, and branches) of a tree, α_c and α_Z are PFT-specific constants, $\theta_c=1.5$ and $\theta_Z=0.5$
 203 (Farrior et al., 2013) (although they could be made PFT-specific if necessary), π is the circular
 204 constant, λ is a PFT-specific taper constant, and ρ_W is PFT-specific wood density (kg C m^{-3})
 205 (Table 1).

206 We set *targets* for leaf (L^*), fine root (FR^*), and sapwood cross-sectional area (A_{SW}^*) that
 207 govern plant allocation of non-structural carbon and nitrogen during growth. These *targets* are
 208 related by the following equations based on the assumption of the pipe model (Shinozaki,
 209 Kichiro et al., 1964):

$$\begin{aligned}
L^*(D, p) &= l^* \cdot A_{CR}(D) \cdot \sigma \cdot p(t) \\
FR^*(D) &= \varphi_{RL} \cdot l^* \cdot \frac{A_{CR}(D)}{\gamma} \\
A_{SW}^*(D) &= \alpha_{CSA} \cdot l^* \cdot A_{CR}(D)
\end{aligned}
\tag{Eq. 3}$$

210 where $L^*(D, p)$, $FR^*(D)$, and $A_{SW}^*(D)$ are the targets of leaf mass (kg C/tree), fine root biomass
211 (kg C/tree), and sapwood cross sectional area (m²/tree), respectively, at tree diameter D ; l^* is the
212 target leaf area per unit crown area of a given PFT; $A_{CR}(D)$ is the crown area of a tree with
213 diameter D ; σ is PFT-specific leaf mass per unit area (LMA); and $p(t)$ is a PFT-specific function
214 ranging from zero to one that governs leaf phenology (Weng et al., 2015); φ_{RL} is the target ratio
215 of total root surface area to the total leaf area; γ is specific root area; and α_{CSA} is an empirical
216 constant (the ratio of sapwood cross-sectional area to target leaf area). The phenology function
217 $p(t)$ takes values 0 (non-growing season) or 1 (growing season) following the phenology model
218 of LM3-PPA (Weng et al., 2015). The onset of a growing season is controlled by two variables,
219 growing degree days (GDD), and a weighted mean daily temperature (T_{pheno}), while the end of a
220 growing season is controlled by T_{pheno} . (Please see Supplementary Information I for details of the
221 phenology model)

222 Nitrogen uptake

223 The rate of nitrogen uptake (U , g N m⁻² hour⁻¹) from the soil mineral nitrogen pool is an
224 asymptotically increasing function of fine root biomass density ($C_{FR, total}$, kg C m⁻²), following
225 McMurtrie *et al.* (2012)

$$U = f_{U, max} \cdot N_{mineral} \cdot \frac{C_{FR, total}}{C_{FR, total} + K_{FR}}, \tag{Eq. 4}$$

226 where, $N_{mineral}$ is the mineral N in soil (g N m⁻²), $f_{U, max}$ is the maximum rate of nitrogen
227 absorption per hour when $C_{FR, total}$ approaches infinity, K_{FR} is a shape parameter (kg C m⁻²) at

228 which the nitrogen uptake rate is half of the parameter $f_{U,max}$. The nitrogen uptake rate of an
 229 individual tree (U_{tree} , kg N hour⁻¹ tree⁻¹) is calculated as follows:

$$U_{tree} = U \cdot \frac{C_{FR,tree}}{C_{FR,total}}, \quad (\text{Eq. 5})$$

230 where, $C_{FR,tree}$ is the fine root biomass of a tree (kgC tree⁻¹). The nitrogen absorbed by roots
 231 enters into the NSN pool and then is allocated to plant tissues through plant growth.

232 **Allocation and plant growth**

233 The partitioning of carbon and nitrogen into the plant pools (*i.e.*, leaves, fine roots, and
 234 sapwood) is limited by the allometric equations, targets of leaves, fine roots, and sapwood cross-
 235 sectional area, and the stoichiometry (*i.e.*, C:N ratios) of these plant tissues. At a daily time step,
 236 the model calculates the amount of carbon and nitrogen that are available for growth according
 237 to the total NSC and NSN and current leaf and fine root biomass. Basically, the available NSC
 238 (G_C) is the summation of a small fraction (f_1) of the total NSC in an individual plant and the
 239 differences between the targets of leaf and fine roots and their current biomass capped by a larger
 240 fraction (f_2) of NSC (Eq. 6.1). The available NSN (G_N) is analogous to that of the NSC and
 241 meets approximately the stoichiometrical requirement of plant tissues (Eq. 6.2).

$$G_C = \min (f_1 NSC + L^* + FR^* - L - FR, f_2 NSC) \quad (\text{Eq. 6.1})$$

$$G_N = \min (f_1 NSN + N_L^* + N_{FR}^* - N_L - N_{FR}, f_2 NSN,) \quad (\text{Eq. 6.2})$$

242 where L^* and FR^* are the targets of leaves and fine roots, respectively (see Eq. 3); L and FR are
 243 current leaf and fine roots biomass, respectively; N_L^* and N_{FR}^* are nitrogen of leaves and fine
 244 roots at their targets according to their target C:N ratios. The parameter f_2 gives the daily
 245 availability of NSC during periods of leaf flush at the beginning of a growing season and f_1

246 normal growth of stems after plant leaves and fine roots approach their targets. Usually,
 247 parameter f_1 is much greater than f_2 . We let $f_1=0.02$ and $f_2=1/(365 \times 3)$ in this study.

248 The allocation of the available NSC (i.e., G_C) to wood (G_W), leaves (G_L), fine roots (G_{FR}),
 249 and seeds (G_F) follows the equations below (Eq. 7). These equations describe the mass growth of
 250 plant tissues with nitrogen effects on the carbon allocation between high-nitrogen tissues and
 251 low-nitrogen tissues (wood) for maximizing leaves and fine roots growth (G_L and G_{FR} ,
 252 respectively), optimizing carbon usage at given nitrogen supply (G_N), and keeping the tissues at
 253 their target C:N ratios.

$$G_C \geq G_W + G_L + G_{FR} + G_F \quad (\text{Eq. 7.1})$$

$$G_N \geq \frac{G_L}{CN_{L,0}} + \frac{G_{FR}}{CN_{FR,0}} + \frac{G_F}{CN_{F,0}} + \frac{G_W}{CN_{W,0}} \quad (\text{Eq. 7.2})$$

$$\frac{(FR+G_{FR})\gamma}{(L+G_L)/\sigma} = \varphi_{RL} \quad (\text{Eq. 7.3})$$

$$G_L + G_{FR} = \text{Min} \left(\frac{L^* + FR^* - L - FR,}{f_{LFR,max} G_C} \right) \cdot r_{S/D} \quad (\text{Eq. 7.4})$$

$$G_F = \left[G_C - \text{Min} \left(\frac{L^* + FR^* - L - FR,}{f_{LFR,max} G_C} \right) r_{S/D} \right] \cdot v \cdot r_{S/D} \quad (\text{Eq. 7.5})$$

$$G_W = \left[G_C - \text{Min} \left(\frac{L^* + FR^* - L - FR,}{f_{LFR,max} G_C} \right) r_{S/D} \right] \cdot (1 - v \cdot r_{S/D}) \quad (\text{Eq. 7.6})$$

254 where, $CN_{L,0}$, $CN_{FR,0}$, $CN_{F,0}$, and $CN_{W,0}$ are the target C:N ratios of leaves, fine roots, seeds, and
 255 sapwood, respectively; γ is specific root area ($\text{m}^2 \text{kgC}^{-1}$); σ is leaf mass per unit area (kg C m^{-2});
 256 $f_{LFR,max}$ is the maximum fraction of G_C for leaves and fine roots (0.85 in this study); v is the
 257 fraction of left carbon for seeds (0.1 in this study); $r_{S/D}$ is a nitrogen-limiting factor ranging from
 258 0 (no nitrogen for leaves, fine roots, and seeds) to 1 (nitrogen available for full growth of leaves,
 259 fine roots, and seeds). The parameter $r_{S/D}$ controls the allocation of G_C and G_N to the four plant
 260 pools (Eq. 7.1). It can be analytically solved (Eqs. 8 and 9).

$$r_{S/D} = \text{Min} \left[1, \text{Max} \left(0, \frac{G_N - G_C / CN_W}{N_{\text{demand}} - G_C / CN_W} \right) \right], \quad (\text{Eq. 8})$$

261 where, N_{demand} is the potential nitrogen demand for plant growth at $r_{S/D}=1$ (i.e., no nitrogen
262 limitation).

$$N_{\text{demand}} = \frac{\gamma \sigma \left[FR + \text{Min} \left(\frac{L^* + FR^* - L - FR}{f_{LFR, \text{max}} G_C} \right) \right] - \varphi_{RL} L}{(\gamma \sigma + \varphi_{RL}) CN_L} +$$

$$\frac{\varphi_{RL} \left[L + \text{Min} \left(\frac{L^* + FR^* - L - FR}{f_{LFR, \text{max}} G_C} \right) \right] - \gamma \sigma L}{(\gamma \sigma + \varphi_{RL}) CN_F} + \frac{v \left[G_C - \text{Min} \left(\frac{L^* + FR^* - L - FR}{f_{LFR, \text{max}} G_C} \right) \right]}{CN_F} +$$

$$\frac{(1-v) \left[G_C - \text{Min} \left(\frac{L^* + FR^* - L - FR}{f_{LFR, \text{max}} G_C} \right) \right]}{CN_W}. \quad (\text{Eq. 9})$$

263 When $G_N \geq N_{\text{demand}}$ ($r_{S/D} = 1$), there is no nitrogen limitation, and all the G_C will be used for plant
264 growth and the allocation follows the rules of the carbon only model (Eqs 7.4~7.6 as $r_{S/D} = 1$).
265 The excessive nitrogen ($G_N - N_{\text{demand}}$) will be returned to the NSN pool. When $G_C / CN_{W,0} < G_N <$
266 N_{demand} (i.e., $0 < r_{S/D} < 1$), all G_C and G_N will be used in new tissue growth; however, the leaves
267 and fine roots cannot reach their targets at this step. When $G_N \leq G_C / CN_{W,0}$ ($r_{S/D} = 0$), all the G_N
268 will be allocated to sapwood and the excessive carbon ($G_C - G_N CN_{W,0}$) will be returned to NSC
269 pool. This is a very rare case since a low G_N leads to low leaf growth, reducing G_C before the
270 case $G_N < G_C / CN_{W,0}$ happens. Therefore, in most cases, Eq. 7.1 is: $G_C = G_W + G_L + G_{FR} + G_F$.

271 Allocation to wood tissues (G_W) drives the growth of tree diameter, height, and crown
272 area and thus increases the targets of leaves and fine roots (Eq. 3). By differentiating the stem
273 biomass allometry in Eq. 2 with respect to time, using the fact that dS/dt equals the carbon
274 allocated for wood growth (G_W), we have the diameter growth:

$$\frac{dD}{dt} = \frac{G_W}{0.25 \pi \Delta \rho_w \alpha_z (2 + \theta_z) D^{1 + \theta_z}} \quad (\text{Eq. 10})$$

275 This equation transforms the mass growth to structural changes in tree architecture. With an
276 updated tree diameter, we can calculate the new tree height and crown area using allometry
277 equations (Eq. 2) and targets of leaf and fine root biomass (Eq. 3) for the next growth step.

278 Overall, this is a flexible allocation scheme and still follows the major assumptions in the
279 previous version of LM3-PPA (Weng, et al., 2015, 2017). This allocation scheme prioritizes the
280 allocation to leaves and fine roots, maintains a minimum growth rate of stems, and keeps the
281 constant area ratio of fine roots to leaves. Based on these allocation rules, the average allocation
282 of carbon and nitrogen to leaves, fine roots, and wood over a growing season are governed by the
283 targets for the leaf area per unit crown area (i.e., crown leaf area index, l^*) and fine root area per
284 unit leaf area (φ_{RL}). Since the crown leaf area index, l^* , is fixed in this study, φ_{RL} is the key
285 parameter determining the relative allocation of carbon to fine roots and stems. A high φ_{RL}
286 means a high relative allocation to fine roots and therefore low relative allocation to stems, and
287 *vice versa*. Note, here φ_{RL} is fixed for each PFT and will remain so for all the model runs.

288 The process of choosing a context-dependent competitively dominant φ_{RL} will take place
289 after finding the fitness of each φ_{RL} in monoculture and in competition with other PFTs (i.e.,
290 different values of φ_{RL}). The competitively optimal strategy is the one that can successfully
291 exclude all others in the processes of competition and succession, but it is not necessarily the one
292 that maximizes production in monoculture. For example, each φ_{RL} creates an environment of
293 light profile and soil nitrogen in its monoculture. Other φ_{RL} PFTs may have higher fitness in this
294 environment than the one that creates it. Only the competitively dominant strategy has the
295 highest fitness in the environment it creates (Fig. 1: C).

296 **2.2 Site and Data**

297 Data pertaining to vegetation, climate, and soil at Harvard Forest (Aber et al., 1993; Hibbs, 1983;
298 Urbanski et al., 2007) were used to design the plant functional types (PFTs) and ecosystem
299 nitrogen levels used in the simulation experiments, to drive the model, and to calibrate model
300 parameters. Harvard Forest is located in Massachusetts, USA (42.54°, -72.17°). The climate of
301 Harvard Forest is cool temperate with annual precipitation 1050 mm, distributed fairly evenly
302 throughout the year. The annual mean temperature is 8.5 °C with a high monthly mean
303 temperature of 20°C in July and a low of -7°C in January. The soils are mainly sandy loam with
304 average depth around 1 m and are moderately well drained in most areas. In forest sites, soil
305 carbon is around 8 kg C m⁻² and nitrogen 300 g N m⁻² (Compton and Boone, 2000). The
306 vegetation is deciduous broadleaf/mixed forest with major species red oak (*Quercus rubra*), red
307 maple (*Acer rubrum*), black birch (*Betula lenta*), white pine (*Pinus strobus*), and hemlock (*Tsuga*
308 *canadensis*) (Compton and Boone, 2000; Savage et al., 2013). The data used to drive our model
309 runs are gap-filled hourly meteorological data at Harvard Forest from 1991 to 2006, obtained
310 from North American Carbon Program (NACP) Site-Level Synthesis datasets (Barr et al., 2013).

311

312 **2.3 Simulation experiments**

313 We set two atmospheric CO₂ concentration ([CO₂]) levels: 380 ppm and 580 ppm, and
314 eight ecosystem total nitrogen levels (ranging from 114.5 g N m⁻² to 552 g N m⁻² at the interval
315 of 62.5 g N m⁻²) by assigning the initial content of the slow SOM pool for our simulation
316 experiments (Table 2). This range covers the soil nitrogen content at Harvard Forest (Compton
317 and Boone, 2000; Melillo et al., 2011). The nitrogen cycles through the plant and soil pools and
318 is redistributed among them via plant demographic processes, soil carbon transfers, and plant
319 uptake. In all the simulation experiments, we assume the ecosystem has no nitrogen inputs and

320 no outputs for convenience since we already have eight total nitrogen levels to represent the
 321 consequences of different nitrogen input and output processes at an equilibrium state. The PFTs
 322 were based on an evergreen needle-leaved tree PFT with different leaf to fine root area ratios,
 323 ϕ_{RL} , in the range from 1 to 8 (Table 2). Simply stated, the PFTs we investigate only differ in
 324 parameter ϕ_{RL} .

325 We define the model runs initialized with only one fixed- ϕ_{RL} PFT as “monoculture runs”
 326 although the actual allocation of carbon to different plant tissues varies with $[CO_2]$ concentration
 327 and ecosystem nitrogen availability. We define the model runs initialized with multiple PFTs as
 328 “**polyculture runs**” (eight PFTs with different ϕ_{RL} at the beginning, although many are driven to
 329 extinction during a given model run). We conducted one set of monoculture runs and two sets of
 330 polyculture runs (Table 2).

331

332

Table 2 Simulation experiments

Type	Model runs	Initial PFT(s) ϕ_{RL}	Ecosystem total nitrogen levels	CO ₂ concentration [CO ₂]
Monoculture runs	One model run per combination of PFT (ϕ_{RL}), nitrogen level, and CO ₂ concentration	One of the following PFTs: $\phi_{RL} = 1, 2, 3, 4, 5, 6, 7, \text{ or } 8$	Eight levels ranging from 114.5 g N m ⁻² to 552 g N m ⁻² at the interval of 62.5 g N m ⁻² : 114.5 g N m ⁻² , 177 g N m ⁻² , 239.5 g N m ⁻² , 302 g N m ⁻² , 364.5 g N m ⁻² , 427 g N m ⁻² , 489.5 g N m ⁻² ,	Ambient: 380 ppm Elevated: 580 ppm
Polyculture runs I	One model run per combination of nitrogen level and CO ₂ concentration	All the PFTs ($\phi_{RL} = 1 \sim 8$) used in the monoculture runs		
Polyculture runs II	One model run per combination of nitrogen level and CO ₂ concentration	Eight PFTs with ϕ_{RL} ranging from 4.5-0.5 <i>i</i> to 8.5-0.5 <i>i</i> at the interval of 0.5, where <i>i</i> denotes the eight nitrogen		

levels from 114.5 to 552 g N m ⁻² .	552 g N m ⁻²
--	-------------------------

333

334 In the monoculture runs, we run the full combinations of eight PFTs with root/leaf area
335 ratios (ϕ_{RL}) from 1 to 8, eight ecosystem total nitrogen levels, and two CO₂ concentrations [CO₂]
336 (380 ppm and 580 ppm) (Table 2). For the eight PFTs, only those with $\phi_{RL} \leq 6$ survived at
337 ambient [CO₂] (380 ppm) because the carbon consumed by fine roots exceeded what leaves
338 provided at $\phi_{RL} > 6$. The **monoculture** runs are for exploring the model predictions of gross
339 primary production (GPP), net primary production (NPP), allocation, and biomass at equilibrium
340 with fixed ϕ_{RL} and ecosystem total nitrogen levels, analogous to the functional relationship
341 schemes used in many ecosystem models (e.g., De Kauwe et al., 2014).

342 In polyculture runs I, we used the same PFTs as in the monoculture runs, where their ϕ_{RL}
343 varies from 1 to 8 at the interval of 1.0 and the ecosystem total nitrogen levels are the same as
344 those used in the monoculture runs (Table 2). This set of polyculture runs was used to explore
345 successional patterns at both ambient and elevated [CO₂] concentrations (380 ppm and 580 ppm,
346 respectively). However, this set of model runs could not show the details of equilibrium plant
347 biomass and allocation patterns along the nitrogen gradient because of the large intervals
348 between the ϕ_{RL} values.

349 To achieve greater resolution in our competition predictions, we designed the polyculture
350 runs II using a dynamic PFT combination scheme according to the ranges of ϕ_{RL} obtained from
351 the polyculture runs I that could survive at a particular nitrogen level at both CO₂ concentrations.
352 For each nitrogen level, we set eight PFTs with ϕ_{RL} that varied in a range 3.5 (e.g., $x \sim x+3.5$) at
353 the interval of 0.5, starting with the highest ϕ_{RL} of 8.0 at the lowest N level (114.5 g N m⁻²) and
354 decreasing 0.5 per level of increase in ecosystem total N. Let $i=1, 2, \dots, 8$ denote the eight N

355 levels from 114.5 to 552 g N m⁻², the ϕ_{RL} of the eight PFTs at each level are (5.0-0.5*i*, 5.5-
356 0.5*i*, ..., 8.5-0.5*i*) (Table 2). For example, at the nitrogen of 114.5 g N m⁻² (*i* = 1), the ϕ_{RL} of the
357 eight PFTs are 4.5, 5.0, ..., 8.0 and at 177 g N m⁻² (*i* = 2), they are 4.0, 4.5, ..., 7.5.

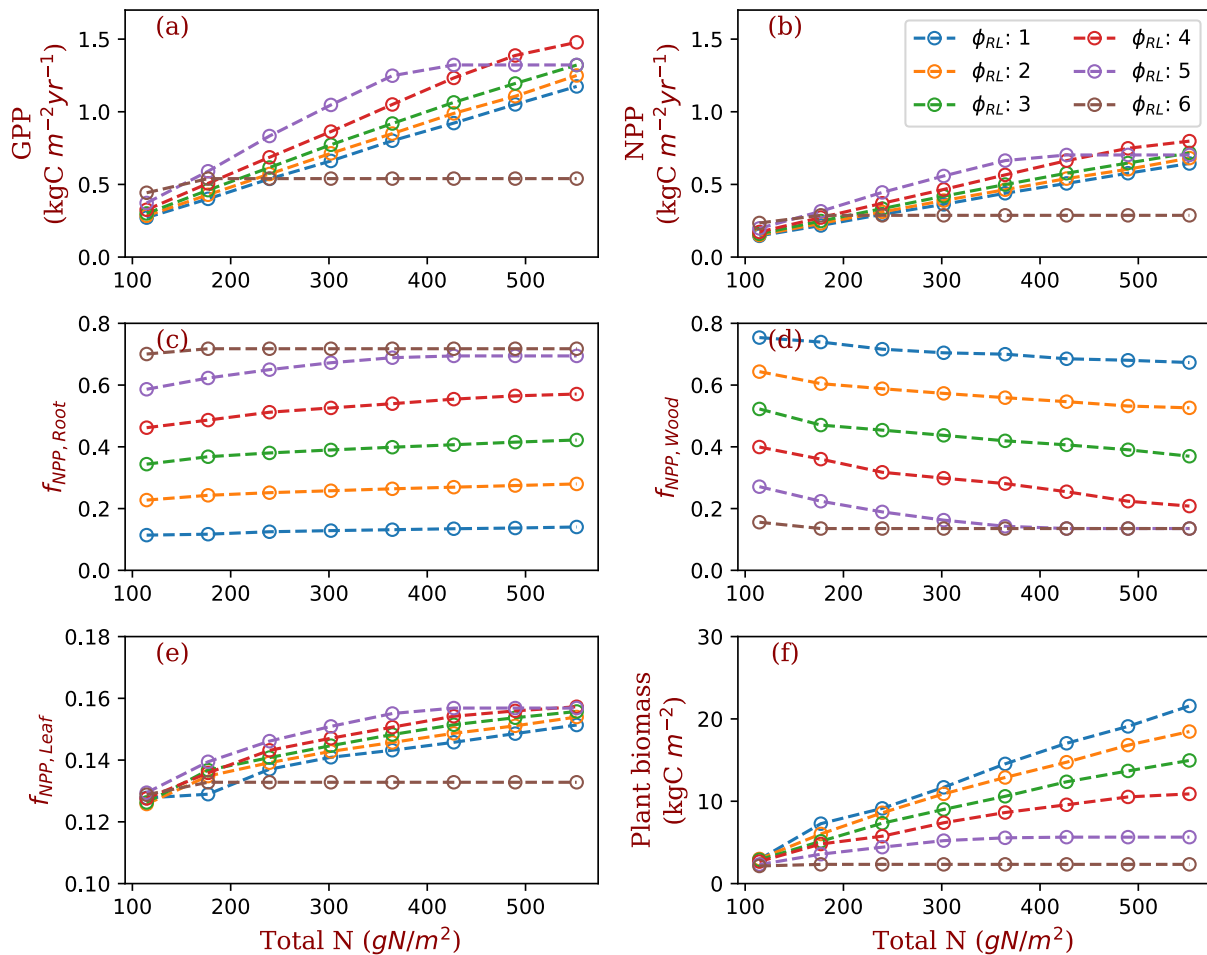
358 For both monoculture and polyculture runs, visual inspection indicated that stands had
359 reached equilibrium after ~1200 years. To be conservative, we present equilibrium data by
360 averaging model properties between years 1400 and 1800. We compared simulated equilibrium
361 gross primary production (GPP), net primary production (NPP), allocation (both absolute amount
362 of carbon and fractions of the total NPP), and plant biomass of the polyculture runs II with those
363 from the monoculture runs. We used the results from one PFT ($\phi_{RL}=4$) to highlight the
364 differences of plant responses with competitively optimal allocation strategies obtained from the
365 polyculture runs II.

366

367 **3 Results**

368 In the monoculture runs, GPP and NPP increase by a factor of three along the gradient of
369 nitrogen used in this study (114.5 - 552 g N m⁻²) at both ambient (Fig. 3) and elevated [CO₂]
370 (Figs. S1). The magnitude of differences in GPP and NPP due to differences in fixed allocation
371 within a given nitrogen level is comparable to the magnitude of differences in GPP and NPP due
372 to nitrogen level within a given fixed allocation strategy (Fig. 3: a and b) when ϕ_{RL} is in the
373 range that allows plants to grow normally (1~5 in the case of ambient [CO₂]). As prescribed by
374 the definition of ϕ_{RL} , allocation of NPP to fine roots increases with ϕ_{RL} in monoculture runs (Fig.
375 3: c). As a consequence, allocation of NPP to wood decreases as ϕ_{RL} increases (Fig. c: d).
376 Allocation to leaves does not change much with ϕ_{RL} . (Fig. 3: e, note differences in scale).
377 Correspondingly, plant biomass at equilibrium decreases with ϕ_{RL} (Fig. 3: f). The effects of

378 nitrogen on the allocation of carbon to fine roots and wood follow our allocation model
 379 assumptions because *proportionally* more carbon is allocated to low-nitrogen woody tissues in
 380 our model when nitrogen is limited. However, the amplitude of changes in GPP and NPP
 381 induced by nitrogen availability is lower than the amplitude of changes resulting from different
 382 values of ϕ_{RL} in the monoculture runs.

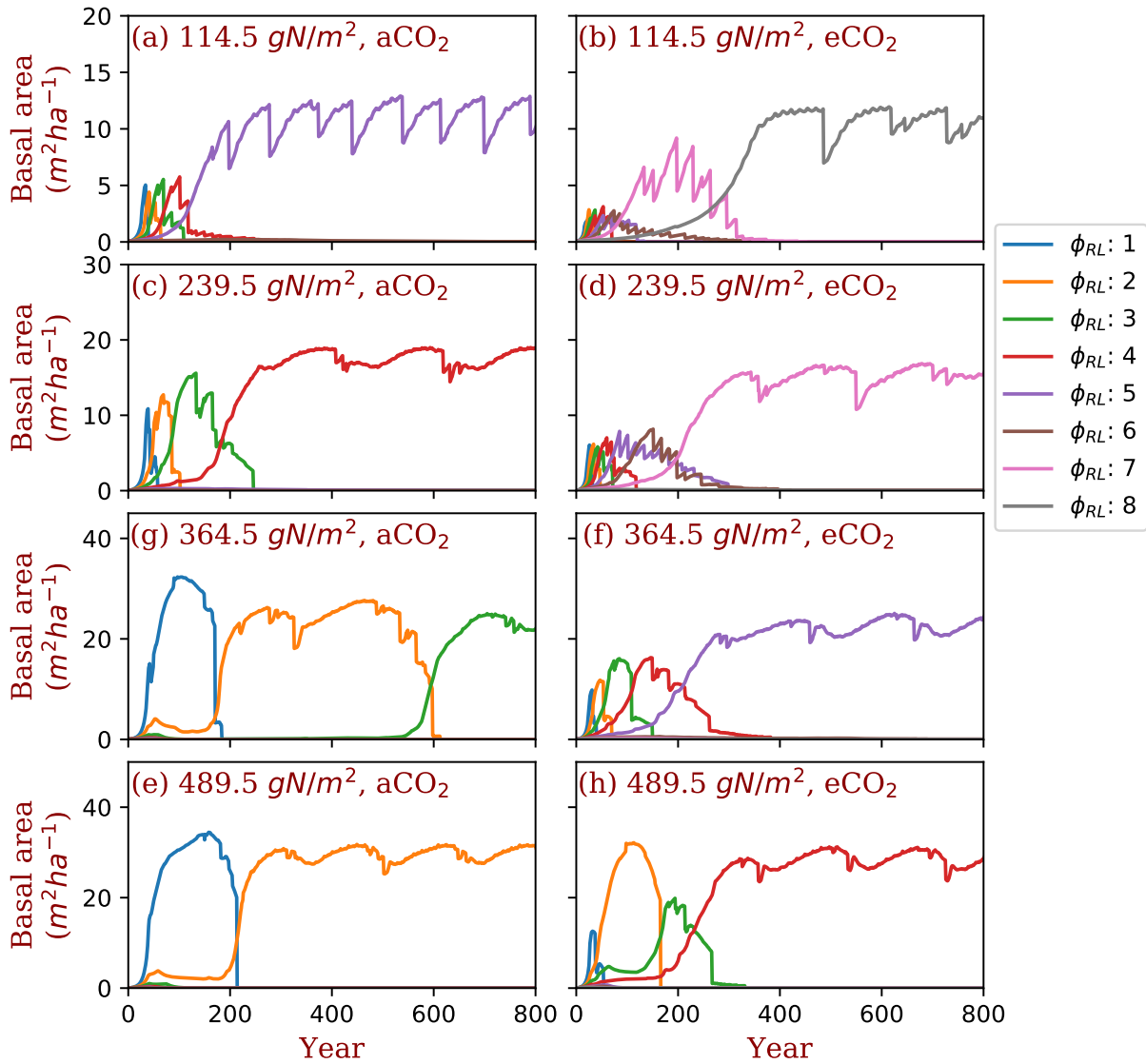


383

384 **Figure 3. GPP, NPP, Allocation and Plant biomass at equilibrium state simulated by**
 385 **monoculture runs.** GPP: Gross primary production; NPP: Net primary production; $f_{NPP,x}$: the
 386 fraction of NPP allocated to x , where x is Root (fine roots), Leaf (leaves in crown), or Wood
 387 (including tree trunk, stems, and coarse roots). The data are from the averages of the model run

388 years from 1400 and 1800. Each model run is initiated with one PFT with fixed ratio of fine root
 389 area to leaf area (ϕ_{RL}).

390



391

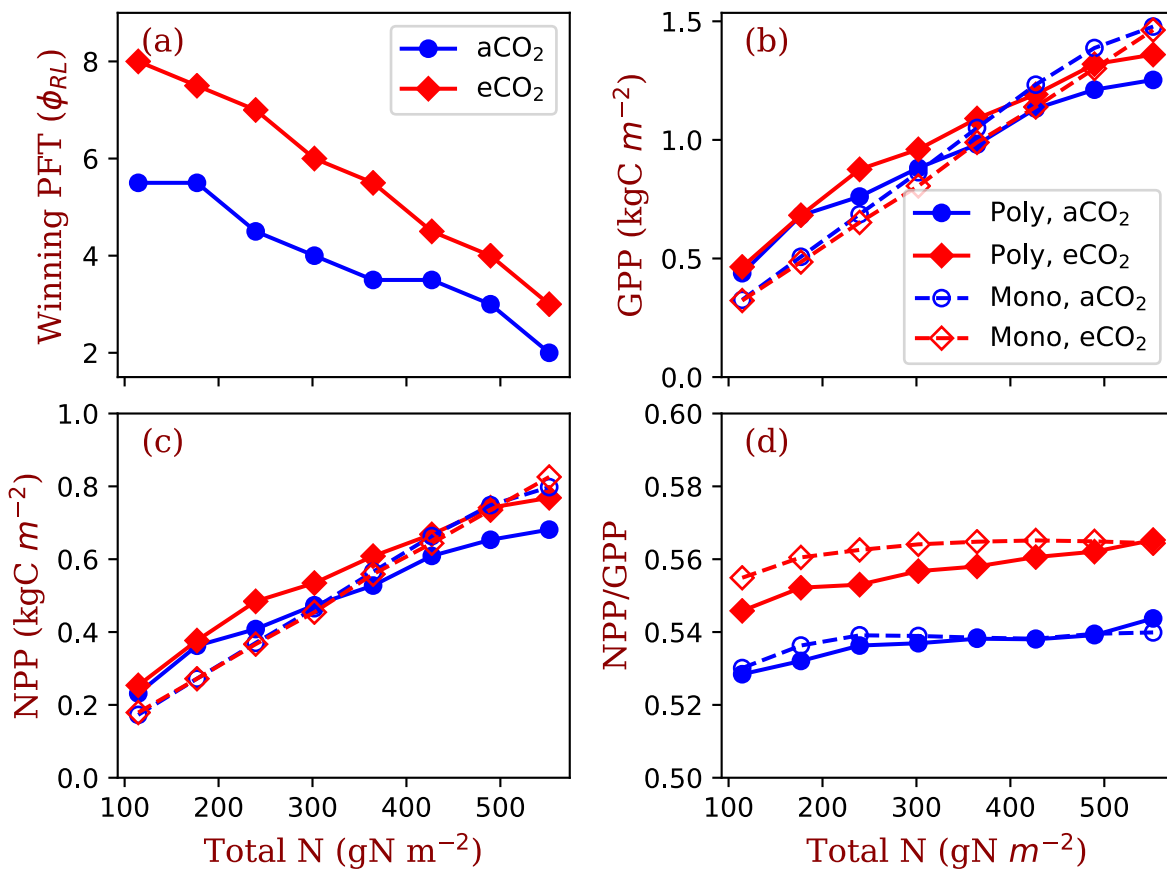
392 **Figure 4 Successional patterns of polyculture runs I at ambient and elevated CO2**
 393 **concentrations.**

394

395 We used two sets of polyculture runs to look for the ϕ_{RL} that is closest to the competitively
 396 optimal (i.e., evolutionarily stable strategy). In the polyculture runs I, where ϕ_{RL} ranges from 1 to

397 8 at all nitrogen levels, the winning strategy (ϕ_{RL}) increases from 5 to 2 as the total nitrogen
 398 increases from 114.5 g N m⁻² to 489.5 g N m⁻² at ambient CO₂ (380 ppm) (Fig. 4: a, c, g, e).
 399 Elevated CO₂ (580 ppm) shifts the winning strategy to higher (ϕ_{RL}) at all the total nitrogen
 400 levels. As shown in Fig. 4, the winning strategy shifts from $\phi_{RL}=5$ to $\phi_{RL}=8$ at 114.5 g N m⁻² and
 401 from $\phi_{RL}=2$ to $\phi_{RL}=4$ at 489.5 g N m⁻².

402



403

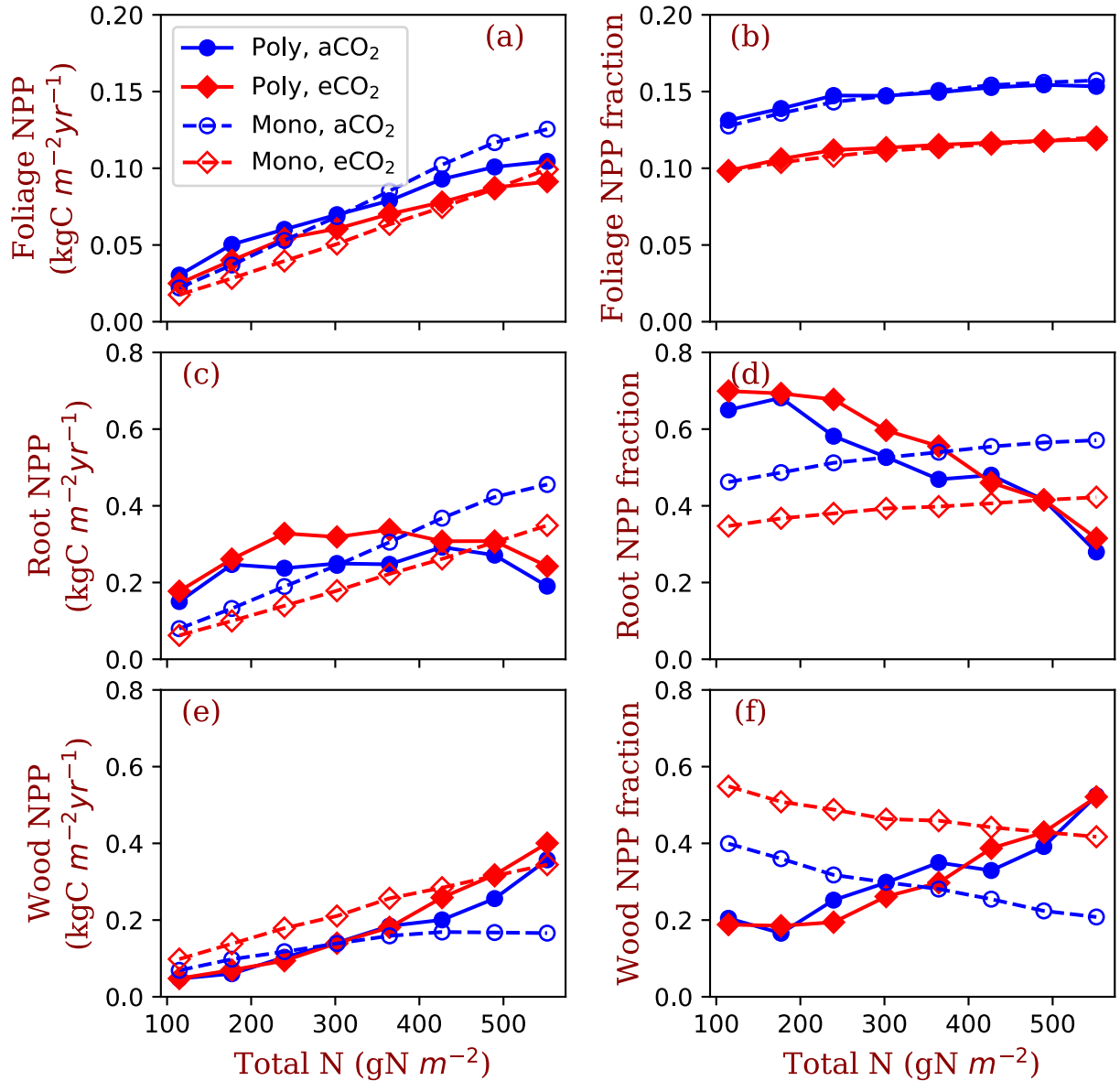
404 **Figure 5 Winning PFTs (ϕ_{RL} , a) in polyculture runs II and equilibrium Gross Primary**
 405 **Production (GPP, b), Net Primary Production (NPP, c), and Carbon Use Efficiency**
 406 **(NPP/GPP, d).** The closed symbols with solid line represent polyculture runs. The open symbols
 407 with dashed lines represent monoculture runs (only $\phi_{RL}=4$ shown in this figure).

408

409 Based on the shifts of the winning ϕ_{RL} from aCO₂ to eCO₂ at the eight nitrogen levels, we
410 designed the polyculture runs II with high resolution of ϕ_{RL} and calculated their GPP, NPP,
411 allocation, and plant biomass at equilibrium state. The of ϕ_{RL} of the winning PFTs decreases
412 from 5.5 to 2 at ambient [CO₂] and from 8.0 to 3.0 at elevated [CO₂] as total N increases from
413 114.5 gN m⁻² to 552.0 gN m⁻². The equilibrium GPP and NPP increase with total nitrogen at
414 values similar to those of the monoculture runs (Fig. 5: b and c). However, the CO₂ stimulation
415 of NPP increases with total nitrogen in the polyculture runs more than it in the monoculture runs.
416 Elevated [CO₂] increases carbon use efficiency (defined as the ratio of NPP to GPP in this study,
417 NPP/GPP) in both the monoculture and polyculture runs (Fig. 5: d). Also, the dependence of
418 NPP:GPP ratio on nitrogen is higher in the polyculture runs than it in the monoculture runs (Fig.
419 5:c).

420 Allocation of NPP to leaves increases with total nitrogen in all conditions, i.e. both
421 competition and monoculture at both ambient [CO₂] and elevated [CO₂] (Fig. 6: a). Foliage NPP
422 is similar in these four model runs when N is low. At high nitrogen (>400 g N m⁻²), polyculture
423 runs have higher foliage NPP than the monoculture runs generally. Allocation to leaves is
424 relatively stable across the nitrogen gradient at the two CO₂ concentration levels (Fig. 6: b). The
425 fraction of NPP allocated to leaves changes little with nitrogen (Fig. 6: b) and it is universally
426 higher at ambient [CO₂] than at elevated [CO₂].

427



428

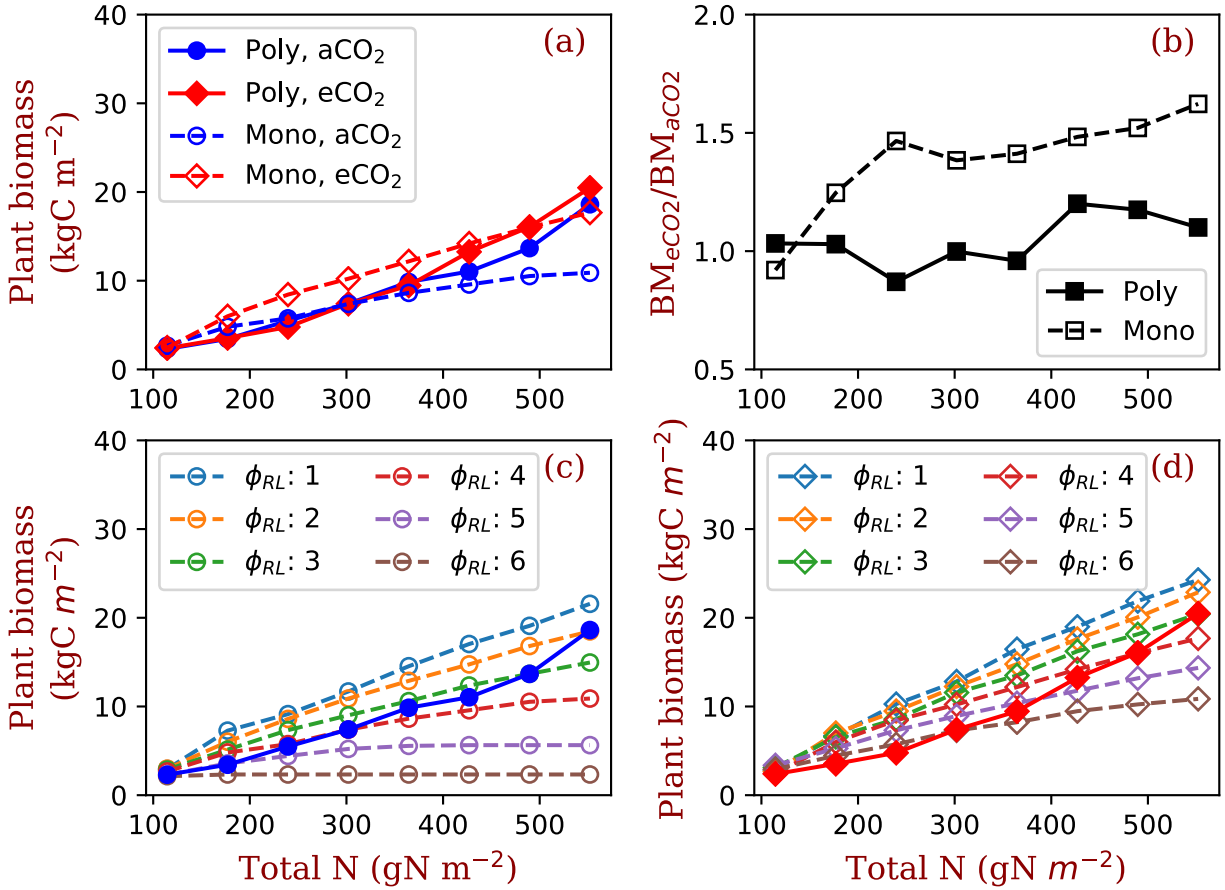
429 **Figure 6** Allocation to leaves, fine roots, and wood tissues of the competition and monoculture
 430 runs at the eight total nitrogen levels and two CO₂ concentrations. The panels a, c, and e show
 431 the NPP allocated to the tissues and the panels b, d, and f show the fractions of the allocation in
 432 total NPP. The closed symbols with solid line represent polyculture runs (poly.). The open
 433 symbols with dashed lines represent monoculture runs (only $\phi_{RL}=4$ shown in this figure).

434

435 Fine root NPP does not significantly change with total nitrogen in polyculture runs,
436 whereas it increases monotonically with increasing nitrogen in monoculture runs (Fig. 6: c).
437 Elevated [CO₂] increases fine root allocation at low nitrogen in polyculture runs but decreases
438 root allocation irrespective of nitrogen in monoculture runs (Fig. 6: c). The fraction of NPP
439 allocated to fine roots decreases with nitrogen at both CO₂ concentrations in polyculture runs but
440 it increases slightly in monoculture runs (Fig. 6: d). In monoculture runs, elevated CO₂ reduces
441 the fraction of NPP allocated to fine roots at all nitrogen levels. In polyculture runs, fractional
442 allocation to fine roots increases at elevated [CO₂] when ecosystem total nitrogen is low (e.g.,
443 114.5 - 302 g N m⁻²) and decrease at elevated [CO₂] when ecosystem total nitrogen is high (e.g.,
444 364-552 g N m⁻²).

445 In the reverse of the fine root response, NPP allocation to woody tissues increases with
446 total nitrogen in both competition and monoculture runs (Fig. 6: e). In polyculture runs, the
447 fraction of allocation to woody tissues decreases at elevated [CO₂] when ecosystem total
448 nitrogen is low (e.g., 114 – 245 g N m⁻²) and increases at elevated [CO₂] when ecosystem total
449 nitrogen is high (e.g., 302 – 552 g N m⁻²).

450



451

452

Figure 7 Plant biomass responses to elevated $[\text{CO}_2]$ and nitrogen

453

Panel a shows the equilibrium plant biomass (means of simulated plant biomass from model run

454

year 1400 to 1800) in polyculture runs and monoculture runs ($\phi_{\text{RL}}=4$). Panel b shows the ratio of

455

simulated plant biomass at elevated $[\text{CO}_2]$ to ambient $[\text{CO}_2]$ for both competition and

456

monoculture runs. Panels c and d show the comparisons with monoculture runs with ϕ_{RL}

457

increasing from 1 to 6 at ambient (c) and elevated $[\text{CO}_2]$ (d). The closed symbols with solid line

458

represent polyculture runs. The open symbols with dashed lines represent monoculture runs (ϕ_{RL}

459

ranges from 1 to 6).

460

461 As a result of the changes in competitively-optimal ϕ_{RL} , plant biomass increases
462 dramatically with ecosystem nitrogen in polyculture runs compared with that in monoculture
463 runs (Fig. 7: a). The effects of elevated $[CO_2]$ on plant biomass increase with nitrogen in
464 polyculture runs but are constant overall in monoculture runs (Fig. 7: b). Compared with the full
465 spread of monoculture runs with ϕ_{RL} ranging from 1 to 6, polyculture runs have high root
466 allocation at low nitrogen and low root allocation at high nitrogen due to changes in the
467 dominant competitive allocation strategy, which amplifies plant biomass responses to elevated
468 $[CO_2]$ with increasing nitrogen (Fig. 7: c and d).

469

470 **4 Discussion**

471 Our competitively-optimal predictions are generally consistent with observations of forest
472 ecosystem production and allocation. For example, high nitrogen environments (i.e., productive
473 environments) favor high wood allocation and low root allocation (Litton et al., 2007; Poorter et
474 al., 2012) because the woody tissues are an unlimited sink for surplus carbon. Low nitrogen
475 availability limits plant CO_2 responses (Norby et al. 2010) in the competition runs (polyculture)
476 because of high root allocation. Our model predicts increased root allocation at all nitrogen
477 levels in response to elevated $[CO_2]$ in the competition runs. Data from free air CO_2
478 enhancement (FACE) forest experiments largely agree (Drake et al., 2011; Iversen et al., 2012;
479 Jackson et al., 2009; Lukac et al., 2003; Nie et al., 2013; Pritchard et al., 2008; Smith et al.,
480 2013). However, in ORNL-FACE, the increases in root production due to elevated CO_2 increase
481 and then declined after 8 years of CO_2 enhancement (Iversen, 2010; Norby and Zak, 2011).
482 Though there are no direct data available for quantitatively validating the patterns predicted by

483 our model, especially for the long-term, competitive runs, a detailed modeling analysis can help
484 to understand the varied patterns in the experiments and shed light on the modeling of allocation.

485

486 **4.1 Modeling of allocation and competition and their effects on model predictions**

487 In our model, the allocation of carbon and nitrogen within an individual tree is based on
488 allometric scaling, functional relationships, and optimization of resource usage. Basically, the
489 allometric scaling relationships define the maximum leaf and fine root growth at a given tree size
490 and the functional relationships (pipe model) define the ratios of leaf area to sapwood cross-
491 sectional area and fine root surface area. These rules are commonly used in ecosystem models
492 (Franklin et al., 2012) and have been shown to generate reasonable predictions (De Kauwe et al.,
493 2014; Valentine and Mäkelä, 2012). Overall, these rules lead to the priority of allocation to
494 leaves and fine roots but allow for structurally-unlimited stem growth when resources (carbon
495 and nitrogen in this study) are available (i.e., the remainder goes to stems after leaf and fine root
496 growth).

497 We define a maximum leaf and fine root allocation, $f_{\text{LFR,max}}$, to limit the maximum
498 allocation to leaves and fine roots to maintain a relatively stable growth rate of wood in years of
499 low productivity. The simulated wood growth patterns agree with real wood growth in temperate
500 trees (Cuny et al., 2012; Michelot et al., 2012). Trees need to grow new wood tissues
501 continuously (especially early in the growing season) to maintain their functions (Plomion et al.,
502 2001). This parameter does not change the fact that leaves and fine roots are the priority. Since
503 allocation ratios to stems are around 0.4~0.7 in temperate forests (Curtis et al., 2002; Litton et
504 al., 2007), with a value of 0.85, $f_{\text{LFR,max}}$ only seldom affects the overall carbon allocation ratios of
505 leaves, fine roots, and stems, and still maintains wood growth in years of low productivity. If

506 $f_{LFR,max} = 1$ (i.e., the highest priority for leaf and fine root growth), simulated trunk radial growth
507 would have unreasonably high interannual variation because leaf and fine root growth would use
508 all carbon to approach to their targets, leaving nothing for stems in some years of low
509 productivity.

510 The simulation of competition for light and soil resources is based on two fundamental
511 mechanisms: 1) competition for light is based on the height of trees according to the rules of the
512 PPA model (Strigul et al., 2008); and 2) individual nitrogen uptake is linearly dependent on the
513 fine root surface area of an individual tree relative to that of its neighbors (Dybzinski et al., 2019;
514 McMurtrie et al., 2012; Weng et al., 2017). These two mechanisms define an allocational
515 tradeoff between wood and fine roots for carbon and nitrogen investment in different $[CO_2]$ and
516 nitrogen environments. Allowing competition for these resources to determine the dominant
517 traits results in very different predicted allocation patterns – and thus ecosystem level responses
518 – than those of fixed allocation strategies. For example, fractional wood allocation increases with
519 increasing nitrogen availability under competitive allocation but decreases – the opposite
520 qualitative response – under a fixed strategy (Fig. 6: f). Consequently, equilibrium plant biomass
521 is predicted to increase much more with increasing nitrogen availability under a competitive
522 strategy than under a fixed strategy (Fig. 4: c, d). In nature, the effects of competition on
523 dominant plant traits may occur through species replacement or community assembly (akin to
524 the mechanism in our model), but it may also occur through adaptive plastic responses or in-
525 place sub-population evolution of ecotypes.

526 Although the strategy that maximizes the growth rate in a fixed-allocation strategy
527 allocates very little to fine roots (Figs. 3 and S1), the competitively optimal strategy allocates
528 more carbon to fine roots to compete for nitrogen, a competitive effect termed “fine-root

529 overproliferation” (Gersani et al., 2001; McNickle and Dybzinski, 2013; O’Brien et al., 2005).
530 Elevated [CO₂] increases the carbon gain of leaves, making more carbon available for roots to
531 compete for nitrogen and thus exacerbating the fine-root overproliferation (Dybzinski et al.,
532 2015). Because most nitrogen uptake is via mass flow and diffusion (Oyewole et al., 2017) and
533 because both of these mechanisms depend on sink strength, individuals with *relatively* greater
534 fine root mass than their neighbors take a greater share of nitrogen, as was recently demonstrated
535 empirically (Dybzinski et al., 2019; Kulmatiski et al., 2017). Thus, fine roots may overproliferate
536 for competitive reasons relative to lower optimal fine root mass in the hypothetical absence of an
537 evolutionary history of competition (Craine, 2006; McNickle and Dybzinski, 2013). The
538 increased fitness (*i.e.*, reproductive success) of the *relatively* greater strategy increases the
539 *absolute* fine root mass. But again, individuals with even *relatively* greater fine root mass take a
540 greater share of nitrogen, leading to what has been termed a “tragedy of the commons” (Gersani
541 et al., 2001). This may also explain why root C:N ratio is highly variable (Dybzinski et al., 2015;
542 Luo et al., 2006; Nie et al., 2013): a high density of fine roots in soil may be more important than
543 the high absorption ability of a single root in competing for soil nitrogen in the usually low
544 mineral nitrogen soils.

545 Root overproliferation is still controversial in experiments. For example, Gersani et al.
546 (2001) and O’Brien (2005) found competing plants generate more roots than those planted
547 isolated for pea and soybeans, respectively; whereas, McNickle and Brown (2014) found root
548 growth follows the availability of soil nutrients and individuals growth with competitors have the
549 same root growth as that predicted by the changed nutrient availability. Roots are far more
550 adaptive and complex than those simulated in models at modifying their growth patterns in
551 response to soil nutrient and water dynamics (Hodge, 2009). The root growth strategies in

552 response to competition also vary with species (Belter and Cahill, 2015). The mechanisms of
553 self-recognition of inter- and intra- roots also can lead to varied behavior of root growth (Chen et
554 al., 2012). However, all of the aforementioned studies considered only *plastic* root
555 overproliferation, where individuals produce more roots in the presence of other individuals than
556 they do in isolation, analogous to stem elongation of crowded seedlings (Dudley and Schmitt,
557 1996). A portion of root overproliferation may also be *fixed*, analogous to trees that still grow tall
558 even when grown in isolation. Dybzinski et al. (2019) showed that plant community nitrogen
559 uptake rate was independent of fine root mass in seedlings of numerous species, suggesting a
560 high degree of fixed fine root overproliferation. To improve root competition models, more
561 detailed experiments that control root growth should be conducted to quantify the marginal
562 benefits of roots in isolated, monoculture, and polyculture environments.

563 At high soil nitrogen, height-structured competition for light (also a game-theoretic
564 tragedy of the commons, Falster and Westoby, 2003; Givnish, 1982) prevails, and trees with
565 greater *relative* allocation to trunks prevail. The balance between these two competitive
566 priorities (fine roots vs. stems) can be observed in our model predictions as a shift from fine root
567 allocation to wood allocation as soil nitrogen increases. The increases in the critical height,
568 which is the height of the shortest tree in canopy layer, from low nitrogen to high nitrogen
569 indicates a shift from the importance of competition for soil nitrogen to the importance of
570 competition for light as ecosystem nitrogen increases (Fig. S6). Because the most competitive
571 type shifts from high fine root allocation to low fine root allocation as ecosystem total nitrogen
572 increases, increases in NPP and plant biomass across the nitrogen gradient are greater than the
573 increases in NPP and plant biomass under the fixed strategy (Fig. 3). This greatly reduces the
574 carbon cost of belowground competition. The slight decrease in the fraction of NPP allocated to

575 leaves at elevated [CO₂] occurs because of increases in total NPP and constant absolute NPP
576 allocation to foliage. It is consistent with FACE experiments that show leaf area index (LAI) in
577 closed-canopy forests is not responsive to elevated [CO₂] (Norby et al., 2003).

578 Our model predicts that the ratio of plant biomass under elevated [CO₂] relative to plant
579 biomass under ambient [CO₂] should increase with increasing nitrogen due to the shift of carbon
580 allocation from fine roots to woody tissues. In contrast, the analytic model of Dybzinski *et al.*
581 (2015) predicts that the ratio of plant biomass under elevated [CO₂] relative to plant biomass
582 under ambient [CO₂] should be largely independent of total nitrogen because of an increasing
583 shift in carbon allocation from long-lived, low-nitrogen wood to short-lived, high-nitrogen fine
584 roots under elevated [CO₂] and with increasing nitrogen. This significant difference between
585 these two predictions traces back to differences in how fine root stoichiometry is handled in the
586 two models. In the model of Dybzinski *et al.* (2015), the fine root C:N ratio is flexible and the
587 marginal nitrogen uptake capacity per unit of carbon allocated to fine roots depends on its
588 nitrogen concentration. Like the model presented here, the model of Dybzinski *et al.* (2015)
589 predicts decreasing fine root mass with increasing nitrogen availability. *Unlike* the model
590 presented here (which has constant fine root nitrogen concentration), the model of Dybzinski et
591 al. (2015) predicts increasing fine root nitrogen concentration with increasing nitrogen
592 availability. As a result, there is less nitrogen to allocate to wood as nitrogen increases in the
593 model of Dybzinski *et al.* (2015) than there is in the model presented here. These countervailing
594 factors even out the ratio of plant biomass under elevated [CO₂] relative to plant biomass under
595 ambient [CO₂] across the nitrogen gradient in Dybzinski *et al.* (2015), whereas their absence
596 amplifies this ratio with increasing nitrogen in the model presented here. Our ability to diagnose

597 and understand this discrepancy highlights the utility of deploying closely-related analytical and
598 simulation models (Weng et al., 2017).

599 **4.2 Model complexity and uncertainty**

600 Compared with the conventional pool-based vegetation models that use pools and fluxes
601 to represent plant demographic processes at a land simulation unit (e.g., grid or patch), VDMs
602 add two more layers of complexity. The first is the inclusion of stochastic birth and mortality
603 processes of individuals (i.e., demographic processes). These processes allow the models to
604 predict population dynamics and transient vegetation structure, such as size-structured
605 distribution and crown organization (e.g., Moorcroft et al., 2001; Strigul et al., 2008). With
606 changes in vegetation structure, allocation and mortality rates can change, generating a different
607 carbon storage accumulation curve compared with those predicted by pool-based models where
608 vegetation structure is not explicitly represented (e.g., Weng et al., 2015). The second is the
609 simulated shift in dominant plant traits during succession due to the shifting of competitive
610 outcomes among different PFTs, which changes the allocation between fast- and slow-turnover
611 pools and thus the parameters of allocation and the residence time of carbon in the ecosystem.

612 Together, these mechanisms may alter long-term predictions of terrestrial carbon cycling
613 due to changes in PFT-based parameters (Dybzinski et al., 2011; Farrior et al., 2013; Weng et al.,
614 2015). As described in the Introduction, current pool-based models can be described by a linear
615 system of equations characterized by the key parameters of allocation, residence time, and
616 transfer coefficients (Eq. 1) with the rigid assumption of unchangeable plant types (Luo et al.,
617 2012; Xia et al., 2013). In VDMs however, allocation, residence time, leaf traits, phenology,
618 mortality, plant forms, and their responses to climate change are all strategies of competition
619 whose success varies with the environmental conditions and the traits of the individuals they are

620 competing against. To make predictions of carbon cycle responses to the novel conditions of
621 climate change, we must understand what determines the most competitive strategy, how the
622 most competitive strategy changes with conditions, and how the most competitive strategy
623 impacts the carbon cycle.

624 Many trade-offs between plant traits can shift in response to environmental and biotic
625 changes, limiting the applicability of varying a single trait, as we have in this study. For example,
626 allocation, leaf traits, mycorrhizal types, and nitrogen fixation can all change with ecosystem
627 nitrogen availability (Menge et al., 2017; Ordoñez et al., 2009; Phillips et al., 2013; Vitousek et
628 al., 2013). The unrealistic effects of model simplification can be corrected by adding important
629 tradeoffs that are missing. For example, the positive feedback between root allocation and SOM
630 decomposition plays a role in mitigating the effects of tragedies of the commons of root over-
631 proliferation (e.g., Gersani et al., 2001; Zea-Cabrera et al., 2006) due to a negative feedback
632 induced by root turnover. High root allocation increases the decomposition rate of SOM and the
633 supply of mineral nitrogen because of the high turnover rate of root litter, which favors a strategy
634 of high wood allocation and reduces the competitive optimal fine root allocation. This negative
635 feedback indicates that the model structure is flexible and that we can incorporate correct
636 mechanisms step by step to improve model prediction skills. Testing single strategies is still a
637 necessary step to improving our understanding of the system and prediction skills of the models,
638 though it could lead to unrealistic responses sometimes.

639

640 **4.3 Implications for Earth system modeling**

641 As shown in model inter-comparison studies, the mechanisms of modeling allocation
642 differ very much, leading to high variation in their predictions (e.g., De Kauwe et al. 2014).

643 Calibrating model parameters to fit data may not increase model predictive skill because data are
644 often also highly variable. Franklin et al. (2012) suggest that in order to build realistic and
645 predictive allocation models, we should correctly identify and implement fundamental principles.
646 Our model predicts similar patterns to those of Valentine and Mäkelä (2012), which are very
647 different in their details but share fundamental principles, including 1) evolutionary- or
648 competitive-optimization, 2) capped leaves and fine roots, 3) structurally unlimited stem
649 allocation (i.e., for optimizing carbon use), and 4) height-structure competition for light and root-
650 mass-based competition for soil resources. The principles 2 and 3 are commonly used in models
651 (De Kauwe et al., 2014; Jiang et al., 2019). However, the different rules of implementing them
652 (e.g., allometric equation, functional relationships, etc.) lead to highly varied predictions (as
653 shown in De Kauwe et al., 2014), though the formulations may be very similar. In competitively-
654 optimal models, such as this study and also Valentine and Mäkelä (2012), the competition
655 processes generate similar emergent patterns by selecting those that can survive in competition,
656 regardless the details of those differences.

657 In this study, similar as in Valentine and Mäkelä (2012), there is a hypothesis for the
658 tradeoffs between light capture and nitrogen uptake via allocation based on insights gained from
659 simpler models (e.g., Dybzinski et al., 2015; Mäkelä et al., 2008) for predicting allocation as an
660 emergent property of competition. One advantage of building a model in this way is that the
661 vegetation dynamics are predicted from first principles, rather than based on the correlations
662 between vegetation properties and environmental conditions. With these first principles, the
663 models can produce reasonable predictions, though the details of physiological and demographic
664 processes vary among models. For vegetation models designed to predict the effects of climate
665 change, the important operational distinction is that the fundamental rules cannot or will not

666 change as climate changes. Nor, presumably, will the underlying ecological and evolutionary
667 processes change as climate changes. The emergent properties can change as climate changes
668 however, and the models built on the “scale-appropriate” unbreakable constraints and ecological
669 and evolutionary processes will be able to accurately predict changes in emergent ecosystem
670 properties.

671 This modeling approach also demands improvement in model validation and benchmarking
672 systems (Collier et al., 2018; Hoffman et al., 2017). As shown in this study, allocation responses
673 to elevated CO₂ at different nitrogen levels in monoculture runs are opposite to those in
674 competitive-allocation runs. For example, in monoculture runs, elevated [CO₂] increases wood
675 allocation and decreases fine root allocation at low nitrogen; whereas in competitive-allocation
676 runs elevated [CO₂] leads to low wood allocation and high fine root allocation. Simply
677 calibrating against short-term observational data may improve the agreements with observations
678 but would not change model predictions because these results emerge from the fundamental
679 assumptions of the models. An updated model benchmarking system should have the metrics of
680 competitive plant traits during the development of ecosystems and their responses to changes in
681 climate.

682

683 **5 Conclusions**

684 Our study illustrates that including the competition processes for light and soil resources in
685 a game-theoretic vegetation demographic model can substantially change the prediction of the
686 contribution of ecosystems to the global carbon cycle. Allowing the model to track the
687 competitive allocation strategies can generate significantly different ecosystem-level predictions
688 (e.g., biomass and ecosystem carbon storage) than those of fixed strategies. Building such a

689 model requires differentiating between the unbreakable tradeoffs of plant traits and ecological
690 processes from the emergent properties of ecosystems. Drawing on insights from closely-related
691 analytical models to develop and understand more complicated simulation models seems, to us,
692 indispensable. Evaluating these models also requires an updated model benchmarking system
693 that includes the metrics of competitive plant traits during the development of ecosystems and
694 their responses to climate changes.

695

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701

702 **Codes and data availability**

703 The codes of the BiomeE model are available at GitHub:

704 <https://github.com/wengensheng/BiomeESS>

705 The simulated data from simulation experiments and Python scripts used in this study will be
706 made publicly available at the publish of this paper.

707

708 **Reference**

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