

1 **Leveraging 35 years of *Pinus taeda* research in the southeastern U.S. to constrain**
2 **forest carbon cycle predictions: regional data assimilation using ecosystem**
3 **experiments**

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35 **Abstract**

36 Predicting how forest carbon cycling will change in response to climate change and management
37 depends on the collective knowledge from measurements across environmental gradients,
38 ecosystem manipulations of global change factors, and mathematical models. Formally
39 integrating these sources of knowledge through data assimilation, or model-data fusion, allows
40 the use of past observations to constrain model parameters and estimate prediction uncertainty.
41 Data assimilation (DA) focused on the regional scale has the opportunity to integrate data from
42 both environmental gradients and experimental studies to constrain model parameters. Here, we
43 introduce a hierarchical Bayesian DA approach (Data Assimilation to Predict Productivity for
44 Ecosystems and Regions, DAPPER) that uses observations of carbon stocks, carbon fluxes,
45 water fluxes, and vegetation dynamics from loblolly pine plantation ecosystems across the
46 Southeastern U.S. to constrain parameters in a modified version of the 3-PG forest growth
47 model. The observations included major experiments that manipulated atmospheric carbon
48 dioxide (CO₂) concentration, water, and nutrients, along with non-experimental surveys that
49 spanned environmental gradients across an 8.6 x 10⁵ km² region. We optimized regionally
50 representative posterior distributions for model parameters, which dependably predicted data
51 from plots withheld from the data assimilation. While the mean bias in predictions of N
52 fertilization experiments, irrigation experiments, and CO₂ enrichment experiments was low,
53 future work needs to focus modifications to model structure that decrease the bias in predictions
54 of drought experiments. Predictions of how growth responded to elevated CO₂ strongly
55 depended on whether ecosystem experiments were assimilated and whether the assimilated field
56 plots in the CO₂ study were allowed to have different mortality parameters than the other field
57 plots in the region. We present predictions of stem biomass productivity under elevated CO₂,

58 decreased precipitation, and increased nutrient availability that include estimates of uncertainty
59 for the Southeastern U.S. Overall, we: 1) demonstrated how three decades of research in
60 southeastern U.S. planted pine forests can be used to develop DA techniques that use multiple
61 locations, multiple data streams, and multiple ecosystem experiment types to optimize
62 parameters, and 2) developed a tool for the development of future predictions of forest
63 productivity for natural resource managers that leverage a rich dataset of integrated ecosystem
64 observations across a region.
65

66 **1 Introduction**

67 Forest ecosystems absorb and store a large fraction of anthropogenic carbon dioxide (CO₂)
68 emissions (Le Quere et al., 2015; Pan et al., 2011) and supply wood products to a growing
69 human population (Shvidenko et al., 2005). Therefore, predicting future carbon sequestration and
70 timber supply is critical for adapting forest management practices to future environmental
71 conditions and for using forests to assist with the reduction of atmospheric CO₂ concentrations.
72 The key sources of information for developing these predictions are results from global change
73 ecosystem manipulation experiments, observations of forest dynamics across environmental
74 gradients, and process-based ecosystem models. The challenge is integrating these three sources
75 into a common framework for creating probabilistic predictions that provide information on both
76 the expected future state of the forest and the probability distribution of those future states.

77
78 Data assimilation (DA), or data-model fusion, is an increasingly used framework for integrating
79 ecosystem observations into ecosystem models (Luo et al., 2011; Niu et al., 2014; Williams et
80 al., 2005). DA integrates observations with ecosystem models through statistical, often Bayesian,
81 methods that can generate probability distributions for ecosystem model parameters and initial
82 states. DA allows for the explicit accounting of observational uncertainty (Keenan et al., 2011),
83 the incorporation of multiple types of observations with different time scales of collection
84 (MacBean et al., 2016; Richardson et al., 2010), and the representation of prior knowledge
85 through informed parameter prior distributions or specific relationships among parameters
86 (Bloom and Williams, 2015).

87
88 Using DA to parameterize ecosystem models with observations from multiple locations that

89 leverage ecosystem manipulation experiments and environmental gradients will allow for
90 predictions to be consistent with the rich history of global change research in forest ecosystems.
91 Ecosystem manipulation experiments provide a controlled environment in which data collected
92 can be used to describe how forests acclimate and operate under altered environmental
93 conditions (Medlyn et al., 2015) and can potentially allow for the optimization of model
94 parameters associated with the altered environmental factor in the experiment. Furthermore, the
95 assimilation of data from ecosystem manipulation experiments may increase parameter
96 identifiability (reducing equifinality; Luo et al., 2009), where two parameters have compensating
97 controls on the same processes, by isolating the response to a manipulated driver. Observations
98 that span environmental gradients include measures of forests ecosystem stocks and fluxes across
99 a range of climatic conditions, nutrient availabilities, and soil water dynamics. These studies
100 leverage time and space to quantify the sensitivity of forest dynamics to environmental variation.
101 However, covariation of environmental variation can pose challenges separating the responses to
102 individual environmental factors. Overall, assimilating observations from a region that includes
103 environmental gradients and manipulation experiments is a useful extension of prior DA research
104 focused on DA at a single site with multiple types of observations (Keenan et al., 2012;
105 Richardson et al., 2010; Weng and Luo, 2011).

106
107 Southeastern U.S. planted pine forests are ideal ecosystems for exploring the application of DA
108 to carbon cycle and forest production predictions. These ecosystems are dominated by loblolly
109 pine (*Pinus taeda* L.), thus allowing for a single parameter set to be applicable to a large region
110 containing many soil types and climatic gradients. Loblolly pine represents more than one half of
111 the standing pine volume in the southern United States (11.7 million ha) and is by far the single

112 most commercially important forest tree species for the region, with more than 1 billion
113 seedlings planted annually (Fox et al., 2007; McKeand et al., 2003). There is also a rich history
114 of experimental research located across the region focused on global change factors that have
115 included nutrient addition (Albaugh et al., 2016; Carlson et al., 2014; Raymond et al., 2016),
116 water exclusion (Bartkowiak et al., 2015; Tang et al., 2004; Ward et al., 2015; Will et al., 2015),
117 and water addition experiments (Albaugh et al., 2004; Allen et al., 2005; Samuelson et al., 2008).
118 The region also includes a multi-year ecosystem CO₂ enrichment study (McCarthy et al., 2010).
119 Furthermore, many of these experiments are multi-factor with water exclusion by nutrient
120 addition (Will et al., 2015), water addition by nutrient addition (Albaugh et al., 2004; Allen et al.,
121 2005; Samuelson et al., 2008), and CO₂ by nutrients addition treatments (McCarthy et al., 2010;
122 Oren et al., 2001). Beyond experimental treatments, Southeastern U.S. loblolly pine ecosystems
123 include at least two eddy-covariance sites with high frequency measurements of C and water
124 fluxes along with biometric observations over many years (Noormets et al., 2010; Novick et al.,
125 2015), and sites with multi-year sap flow data (Ewers et al., 2001; Gonzalez-Benecke and
126 Martin, 2010; Phillips and Oren, 2001). Finally, there are studies that include plots that span the
127 regional environmental gradients and extend back to the 1980s (Burkhart et al., 1985). Overall,
128 the multi-decadal availability of observations of C stocks (or biomass), leaf area index (LAI), C
129 fluxes, water fluxes, and vegetation dynamics in plots with experimental manipulation and plots
130 across environmental gradients, is well suited to potentially constrain model parameters and
131 predictions of how carbon cycling responds to environmental change.

132

133 Using loblolly pine plantations across the southeastern U.S as a focal application, our objectives
134 were to 1) develop and evaluate a new DA approach that integrates diverse data from multiple

135 locations and experimental treatments with an ecosystem model to estimate the probability
136 distribution of model parameters, 2) examine how the predictive capacity and optimized
137 parameters differ between an assimilation approach that only uses environmental gradients and
138 an assimilation approach that uses both environmental gradients and ecosystem manipulations,
139 and 3) demonstrate the capacity of the DA approach to predict, with uncertainty, regional forest
140 dynamics by simulating how forest productivity responds to drought, nutrient fertilization, and
141 elevated atmospheric CO₂ across the Southeastern U.S.

142

143 **2 Methods**

144

145 **2.1 Observations**

146 We used thirteen different data streams from 294 plots at 187 unique locations spread across the
147 native range of loblolly pine trees to constrain model parameters (Table 1; Figure 1). The data
148 streams covered the period between 1981 to 2015. The Forest Modeling Research Cooperative
149 (FMRC) Thinning Study provides the largest number of plots that span the region (Burkhart et
150 al., 1985). In this study, we only used the control plots that were not thinned. The Forest
151 Productivity Cooperative (FPC) Region-wide 18 (RW18) study included control and nutrient
152 fertilization addition plots that span the region (134.4 kg ha⁻¹ N + 13.44 kg ha⁻¹ P biannually)
153 (Albaugh et al., 2015). The PINEMAP study included four locations dispersed across the region
154 that included a replicated factorial experiment with control, nutrient fertilization (224 kg ha⁻¹ N +
155 27 kg ha⁻¹ P + micronutrients once at project initiation), throughfall reduction (30% reduction),
156 and fertilization by throughfall treatments (Will et al., 2015). The SETRES study was located at
157 a single location and included replicated control, irrigation (~650 mm of added water per year),

158 nutrient fertilization ($\sim 100 \text{ kg N ha}^{-1} + 17 \text{ kg P ha}^{-1}$ with micronutrients applied annually with
159 absolute amount depending on foliar nutrient ratios), and fertilization by irrigation treatments
160 (Albaugh et al., 2004). The Waycross study was a single site with a non-replicated fertilization
161 treatment. The annual application of nutrient fertilization was focused on satisfying the nutrient
162 demand by the trees and resulted in one most productive stands in the region (Bryars et al.,
163 2013). These five studies included data streams of stand stem biomass (defined as the sum of
164 stemwood, stembark and branches) and live stem density. Waycross and SETRES included LAI
165 measurements from litterfall traps (Waycross) or estimates from LICOR LAI-2200 (SETRES).
166 SETRES also included fine root and coarse root measurements. In the PINEMAP, SETRES, and
167 RW18 studies we only used foliage biomass estimates from the control plots. We excluded the
168 foliage biomass estimates from the treatment plots because they were derived from allometric
169 models that may not have captured changes in allometry due to the experimental treatment. We
170 did use LAI measurements from both control and treatment plots where available (SETRES).
171
172 We also included observations the Duke FACE study where the atmospheric CO_2 was increased
173 by 200 ppm above ambient concentrations. Based on the data presented in McCarthy et al.
174 (2010) the study included six control plots, four CO_2 fumigated rings (including the unfertilized
175 half of the prototype), two nitrogen fertilization treatments ($115 \text{ kg N ha}^{-1} \text{ yr}^{-1}$ applied annually) ,
176 and one CO_2 by nitrogen addition treatment (fertilized half of prototype). The Duke FACE study
177 included observations of stem biomass (loblolly pine and hardwood), coarse root biomass
178 (loblolly pine and hardwood), fine root biomass (combined loblolly pine and hardwood), stem
179 density (loblolly pine only), leaf turnover (combined loblolly pine and hardwood), fine root
180 production (combined loblolly pine and hardwood), and monthly LAI (loblolly pine and

181 hardwood).

182

183 Finally, we included two Ameriflux sites with eddy-covariance towers in loblolly pine stands.

184 The US-DK3 site was located in the same forest as the Duke FACE site described above (Novick

185 et al., 2015). The US-NC2 site was located in coastal North Carolina (Noormets et al., 2010).

186 We used monthly gross ecosystem production (GEP; modeled gross primary productivity from

187 net ecosystem exchange measured at an eddy-covariance tower) and evapotranspiration (ET)

188 estimates from the sites. The monthly GEP and ET were gap-filled by the site PI. The GEP was

189 a flux partitioned product created by the site PI. The biometric data from the US-DK3 site was

190 assumed to be the same as the first control ring. The biometric data from the US-NC2 site

191 included observations of stem biomass (loblolly pine and hardwood), coarse root biomass

192 (loblolly pine and hardwood), fine root biomass (combined loblolly pine and hardwood), stem

193 density (loblolly pine only), leaf turnover (combined loblolly pine and hardwood), and fine root

194 production (combined loblolly pine and hardwood).

195

196 **2.2 Ecosystem Model**

197 We used a modified version of the Physiological Principles Predicting Growth (3-PG) model to

198 simulate vegetation dynamics in loblolly pine stands (Bryars et al., 2013; Gonzalez-Benecke et

199 al., 2016; Landsberg and Waring, 1997). 3-PG is a stand-level vegetation model that runs at the

200 monthly time-step and includes vegetation carbon dynamics and a simple soil water bucket

201 model (Figure 2). While a complete description of the 3-PG model and our modifications can be

202 found in the Supplemental Material Section 1, the key concept for interpreting the results is that

203 gross primary productivity (GPP) was simulated using a light-use efficiency approach where the

204 absorbed photosynthetically active radiation (APAR) was converted to carbon based on a
205 quantum yield (Supplemental Material Section 1.1). Quantum yield was simulated using a
206 parameterized maximum quantum yield (α) that was modified by environmental conditions
207 including atmospheric CO₂, available soil water (ASW) and soil fertility (Supplemental Material
208 Section 1.2-1.3). The ASW and soil fertility modifiers were values between 0 and 1, while the
209 atmospheric CO₂ modifier had a value of 1 at 350 ppm (thus values greater than 1 at higher CO₂
210 concentrations).

211
212 Elevated CO₂ modified tree physiology by increasing quantum yield, based on an increasing but
213 saturating relationship with atmospheric CO₂ (Supplemental Material Section 1.2). Based on
214 initial results from the data assimilation, we also added a function where the allocation to foliage
215 relative to stem biomass decreased as atmospheric CO₂ increased (Supplemental Material Section
216 1.2). ASW and quantum yield were positively related through a logistic relationship between
217 relative ASW and the quantum yield modifier, where relative ASW was the ratio of simulated
218 ASW to a plot-level maximum ASW. Soil fertility and quantum yield were proportionally
219 related, where quantum yield was scaled by an estimate of relative stand-level fertility (a value of
220 1 was the maximum fertility). The fertility modifier (FR) was constant throughout a simulation
221 of a plot and was either based on site characteristics or directly optimized as a stand-level
222 parameter (Supplemental Material Section 1.3). For plots with nutrient fertilization, FR was a
223 directly optimized parameter or set to 1, depending on the level of fertilization (see below). For
224 unfertilized plots, we used site index (SI), a measure of the height of a stand at a specified age
225 (25 years), to estimate FR. This approach is in keeping with previous efforts (Gonzalez-Benecke
226 et al., 2016; Subedi et al., 2015); however, SI does not solely represent nutrient availability of an

227 ecosystem. For a given climate SI captures differences in soil fertility, where a lower SI
228 corresponded to a site with lower fertility, but regional variation in SI also included the influence
229 of climate on growth rates that were already accounted for in the other environmental modifiers
230 in the 3-PG model. When a climate term is not used in the empirical FR model, FR is relative to
231 the highest SI in the region, which does not occur in the northern extent of the region even in
232 fertilized plots due to climatic constraints. Thus, we also included the historical (1970-2011) 35-
233 year mean annual temperature (MAT) as an additional predictor, resulting in an empirical
234 relationship that predicted FR as an increasing, but saturating, function of SI within areas of
235 similar long-term temperature. For our application of the 3-PG model using DA, we removed
236 the previously simulated dependence of total root allocation on FR (Bryars et al., 2013;
237 Gonzalez-Benecke et al., 2016) because we separated coarse and fine roots. Other environmental
238 conditions influenced GPP, including temperature, frost days, and vapor pressure deficit (VPD).
239 A description of these modifiers can be found in Supplemental Material Section 1.2.

240

241 Each month, net primary production (a parameterized and constant proportion of GPP) was
242 allocated to foliage, stem (stemwood, stembark, and branches), coarse roots, and fine roots
243 (Supplemental Material Section 1.4). Differing from previous applications of 3-PG to loblolly
244 pine ecosystems, we modified the model to simulate fine roots and coarse roots separately. 3-PG
245 also simulated simple population dynamics by including stem density as a state variable. Stem
246 density and stem biomass pools were reduced by both density-dependent mortality, based on the
247 concept of self-thinning (Landsberg and Waring, 1997), and density-independent mortality, a
248 new modification where a constant proportion of individuals die each month (Supplemental
249 Material Section 1.5). Finally, we added a simple model of hardwood understory vegetation to

250 enable the assimilation GEP and ET observations from eddy-covariance tower studies with
251 significant understories (Supplemental Material Section 1.7).

252

253 The water cycle was a simple bucket model with transpiration predicted using a Penman-
254 Monteith approach (Bryars et al., 2013; Gonzalez-Benecke et al., 2016; Landsberg and Waring,
255 1997)(Supplemental Material Section 1.6). The canopy conductance used in the Penman-
256 Monteith subroutine was modified by environmental conditions. The modifiers included the
257 same ASW and VPD modifier as used in the GPP calculation. Maximum canopy conductance
258 occurred when simulated LAI exceeded a parameterized value of LAI (LAI_{gcx}). Evaporation
259 was equal to the precipitation intercepted by the canopy. Runoff occurred when the ASW
260 exceeded a plot-specific maximum ASW. As in prior applications of 3-PG, ASW was not
261 allowed take a value below a minimum ASW, resulting in an implicit irrigation in very dry
262 conditions. This assumption may cause the model to be less sensitive to low ASW but the
263 optimized parameterization may compensate.

264

265 The 3-PG model used in this study simulated the monthly change in eleven state variables per
266 plot: four stocks for loblolly pines, five stocks for understory hardwoods, loblolly pine stem
267 density (stems ha⁻¹), and ASW. The key fluxes that were used for DA included monthly GEP,
268 monthly ET, annual root turnover, and annual foliage turnover. In total, 46 parameters were
269 required by 3-PG. The model required mean daily maximum temperature, mean daily minimum
270 temperature, mean daily PAR, total frost days per month, total rain per month, annual
271 atmospheric CO₂, and latitude. Each plot also required maximum ASW, SI, MAT, and the initial
272 condition of the eleven state variables as model inputs (Figure 3).

273
274 We used the first observation at the plot as the initial conditions for the loblolly pine vegetation
275 states (foliage biomass, stem biomass, coarse root biomass, fine root biomass, and stem number).
276 When observations of coarse biomass and fine root biomass were not available, these stocks
277 were initialized as a mean region-wide proportion of the observed stem biomass. However, the
278 value of initial root biomass in plots without observations was not important because root
279 biomass did not influence any other functions in the model. The hardwood understory stocks at
280 US-DK3 and US-NC2 were also initialized using the first set of observations. Initial fine root
281 and coarse biomass were distributed between loblolly pine and hardwoods based on their relative
282 contribution of total initial foliage biomass. The initialized ASW was assumed to be equal to the
283 maximum ASW because most plots were initialized in winter months when plant demand for
284 water was minimal. The maximum ASW in each plot was extracted from the SSURGO soils
285 dataset (Soil Survey Staff, 2013). The value we used corresponded to the maximum ASW for the
286 top 1.5 m of the soil. We assumed that the minimum ASW was zero. Because we focused on a
287 region-wide optimization, we used region-wide 4-km estimates of observed monthly
288 meteorology as inputs and to calculate the 35-year MAT for each plot (Abatzoglou, 2013). SI
289 was based on height measurements at age 25 in each plot or calculated by combining
290 observations of height at younger ages with an empirical model (Diegues-Aranda et al., 2006).

291
292 We simulated ecosystem manipulation experiments in the 3-PG model by altering the
293 environmental modifiers or by modifying the environmental inputs. Nutrient addition
294 experiments were simulated by setting FR equal to 1 for the studies that applied nutrients at
295 regular interval to remove nutrient deficiencies (RW18, SETRES, Waycross). FR was directly

296 estimated for fertilized plots in two of the studies either because nutrients were only added once
297 at the beginning of the study (PINEMAP), thus potentially not removing nutrient limitation, or
298 nitrogen was the only element added (Duke FACE), thus allowing the potential for nutrient
299 limitation by other elements. For these plots, we also assumed that the FR of the fertilized plot
300 was equal to or larger than the control plot. Throughfall exclusion experiments were simulated
301 by decreasing the throughfall by 30% in the treatment plots. The SETRES irrigation experiments
302 were simulated by adding 650 mm to ASW between April and October. CO₂ enrichment
303 experiments were simulated by setting the atmospheric CO₂ input equal to the treatment mean
304 from the elevated CO₂ rings (570 ppm). One plot (US-NC2) included a thinning treatment during
305 the period of observation. We simulated the thinning by specifying a decrease in the stem count
306 that matched the proportion removed at the site, with the biomass of each tree equivalent to the
307 average of trees in the plot.

308

309 **2.3 Data assimilation method**

310 We used a hierarchical Bayesian framework to estimate the posterior distributions of parameters,
311 latent states of stocks and fluxes, and process uncertainty parameters. The latent states
312 represented a value of the stock or flux before uncertainty was added through measurement. The
313 approach was as follows.

314

315 Consider a stock or flux (m) for a single plot (p) at time t ($q_{p,m,t}$). $q_{p,m,t}$ is influenced by the
316 processes represented in the 3-PG model and a normally distributed model process error term,

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$$318 \quad q_{p,m,t} \sim N(f(\boldsymbol{\theta}, FR_p), \sigma_m) \quad \text{Equation 1}$$

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where θ is a vector of parameters that are optimized, FR_p is the site fertility, and σ_m is the model process error. Not shown are the vector of parameters that were not optimized (Supplemental Material Table 1), the plot ASW, an array climate inputs, and the initial conditions because these were assumed known and not estimated in the hierarchical model. The process error assumed that the error linearly scales with the magnitude of the prediction:

$$\sigma_m^2 = \gamma_m + \rho_m f(\theta, FR_p) \quad \text{Equation 2}$$

While the structure of the Bayesian model allowed for all data streams to have process uncertainty that scales with the prediction, in this application we only allowed stem biomass, GEP, and ET process uncertainty to scale because they had large variation across space (stem biomass) and through time (i.e., there should be lower process uncertainty in the winter when GEP is lower). For the other data streams, the linear scaling term was removed by fixing ρ_m at 0.

FR_p did not have an explicit probability distribution. Rather the probability density evaluated to 1 if the plot was not fertilized, thus causing FR_p to be estimated from SI and MAT (Supplemental Material Equation 15), or if it was a fertilized plot and has an FR_p equal or higher than that of its non-fertilized control plot. The probability density evaluated to 0 if the estimated FR_p in a fertilized plot was less than the FR_p in the control plot or FR_p was not contained in the interval between 0 and 1.

$$\text{FR}_p \sim \begin{cases} 1 & \text{if non-fertilized, } \text{FR}_p \geq 0, \text{ and } \text{FR}_p \leq 1 \\ 1 & \text{if } \text{FR}_p = 1 \text{ and fertilization levels are assumed to remove nutrient deficiencies} \\ 0 & \text{if } \text{FR}_p < 1 \text{ and fertilization levels are assumed to remove nutrient deficiencies} \\ 1 & \text{if fertilized but levels are not assumed to remove deficiencies and } \text{FR}_p \geq \text{FR of control plot} \\ 0 & \text{if fertilized but levels are not assumed to remove deficiencies and } \text{FR}_p < \text{FR of control plot} \\ 0 & \text{if } \text{FR}_p < 0 \text{ or } \text{FR}_p > 1 \end{cases}$$

342 Equation 3

343

344 Our model included the effect of observational errors for measurements of stocks and fluxes.

345 For a single stocks or flux for a plot at time t there was an observation ($y_{p,m,t}$). The normally

346 distributed observation error model was:

347

$$y_{p,m,t} \sim N(q_{p,m,t}, \tau_{p,m,t}^2) \quad \text{Equation 4}$$

349

350 where $\tau_{p,m,t}^2$ represented the measurement error of the observed state or flux. By including the

351 observational error model, $q_{p,m,t}$ represented the latent, or unobserved, stock or flux. The variance

352 was unique to each observation because it was represented as a proportion of the observed value.

353 The $\tau_{p,m,t}^2$ was assumed known (Table 1) and not estimated in the hierarchical model.

354

355 The hierarchical model required prior distributions for all optimized parameters, including the

356 parameters for the 3-PG model (θ), FR_p , and the process error parameters. The prior

357 distributions for θ are specified in Table 3. Some parameters were informed by previous

358 research in loblolly pine ecosystems while other parameters were ‘uninformative’ with flat

359 distributions that with broad, but physically reasonable, bounds. The prior distributions for the

360 process error parameters were non-informative and had a uniform distribution with upper and

361 lower bounds that spanned the range of reasonable error terms.

362 $\gamma_m \sim U(0.001, 100)$ Equation 5

363 $\rho_m \sim U(0, 10)$ Equation 6

364

365 By combining the data, process, and prior models, our joint posterior that includes all thirteen

366 data streams, plots, months with observations, and fitted parameters was

367

368 $p(\boldsymbol{\theta}, \boldsymbol{\gamma}, \boldsymbol{\rho}, \mathbf{q} | \mathbf{y}, \boldsymbol{\tau}, \text{priors}) \propto$

369
$$\prod_{p=1}^P \prod_{m=1}^M \prod_{t=1}^T N(q_{p,m,t} | f(\boldsymbol{\theta}, FR_p), \gamma_m + \rho_m f(\boldsymbol{\theta}, FR_p))$$

370
$$\prod_{p=1}^P \prod_{m=1}^M \prod_{t=1}^T N(y_{p,m,t} | q_{p,m,t}, \tau_{p,m,t}^2)$$

371
$$\prod_{p=1}^P p(FR_p) \prod_{f=1}^F p(\boldsymbol{\theta}_f) \prod_{m=1}^M p(\gamma_m) \prod_{m=1}^M p(\rho_m)$$

372 Equation 7

373 where bolded components represent vectors, P is the total number of plots, M is the total number

374 of data streams, T is the total months with observations, and F is the total number of 3-PG

375 parameters that are optimized.

376

377 We numerically estimated the joint posterior distribution using the Monte-Carlo Markov Chain –

378 Metropolis Hasting (MCMC-MH) algorithm (Zobitz et al., 2011). This approach has been widely

379 used to approximate parameter distributions in ecosystem DA research (Fox et al., 2009;

380 Trudinger et al., 2007; Williams et al., 2005; Zobitz et al., 2011). Briefly, the algorithm proposed

381 new values for the model parameters, uncertainty parameters, latent states, and FR. The proposed
382 values were generated using a random draw from a normal distribution with a mean equal to the
383 previously accepted value for that parameter and standard deviation equal to the parameter-
384 specific jumping size. The ratio of the proposed calculation of Equation 7 to the previously
385 accepted calculation of Equation 7 was used to determine if the proposed parameter was
386 accepted. If the ratio was greater than or equal to 1 the proposed value was always accepted. If
387 the ratio was less than 1, a random number between 0 and 1 was drawn and the proposed value
388 was accepted if the ratio was greater than the random number. This allowed less probable
389 parameter sets to be accepted, thus sampling the posterior distribution. We adapted the size of
390 the jump size for each parameter to ensure the acceptance rate of the parameter set was between
391 22% and 43% (Ziehn et al., 2012) by adjusting the jump size if the acceptance rate for a
392 parameter was outside the 22 – 43% range. All MCMC-MH chains were run for 30 million
393 iterations with the first 15 million iterations discarded as the burn-in. Four chains were run and
394 tested for convergence using the Gelman–Rubin convergence criterion, where a value for the
395 criterion less than 1.1 indicated an acceptable level of convergence. We sampled every 1000th
396 parameter in the final 15 million iterations of the MCMC-MH chain and used this thinned chain
397 in the analysis described below. The 3-PG model and MCMC-MH algorithm were programed in
398 FORTRAN 90 and used OpenMP to parallelize the simulation of each plot within an iteration of
399 the MCMC-MH algorithm.

400

401 **2.4 Data assimilation evaluation**

402 Using the observations, model, and hierarchical Bayesian method described above, we
403 assimilated both the non-manipulated and manipulated plots (Base assimilation; Table 4). We

404 assessed model performance first by calculating the RMSE and bias of stem biomass predictions
405 (the most common data stream). In the evaluation, we only used the most recent observed values
406 to increase the time length between initialization and validation. Second, we assessed the
407 predictive capacity by comparing model predictions to data not used in the parameter
408 optimization in a cross-validation study. In this evaluation, we repeated the Base assimilation
409 without 160 FMRC thinning study plots (Table 2), predicted the 160 plots using the median
410 parameter values, and calculated the RMSE and bias stem biomass of the independent set of
411 plots. Rather than holding out all 160 plots from a single assimilation and not generating a
412 converged chain, we divided the 160 plots into four unique sets of 40 plot and repeated the
413 assimilation for each set. Finally, we compared the predicted responses to experimental
414 manipulation to the observed responses. We focused the comparison on the percentage
415 difference in stem biomass between the control and treatment plots. We used a paired t-test to
416 test for differences between the predicted and observed responses within an experimental type
417 (irrigated, drought, nutrient addition, and elevated CO₂). We combined the single and multi-
418 factor treatments for analysis. For the analysis of the nutrient addition studies we only used plots
419 where FR was assumed to be 1 so that we were able to simulate the treatments without requiring
420 the optimization of a site-specific FR parameter.

421

422 During preliminary analysis, we found that the Base assimilation predicted lower stem biomass
423 than observed in the elevated CO₂ plots in the Duke FACE study. Further analysis investigating
424 the cause of the bias in the CO₂ plots showed that three parameters (wSx1000, ThinPower, and
425 pCRS) were required to be unique to the Duke FACE study in order to reduce the bias.
426 Therefore, the Base assimilation included unique parameters for wSx1000, ThinPower, and

427 pCRS parameters in all plots in the Duke FACE and US-DK3 studies. To highlight the need for
428 the site-specific parameters, we repeated the Base assimilation approach without the three
429 additional parameters for the Duke studies (NoDkPars assimilation).

430

431 **2.5 Sensitivity to inclusion of ecosystem experiments**

432 We also evaluated how parameter distributions and the associated environmental sensitivity of
433 model predictions depended on the inclusion of ecosystem experiments in data assimilation.
434 First, we repeated the Base assimilation, this time excluding the plots that included the
435 manipulated treatments (NoExp). We removed all manipulation types at once, rather than
436 individual experimental types, because all experimental types involved multi-factor studies. The
437 NoExp assimilation had the same number of data streams as the Base assimilation because it
438 included the control treatments from the experimental studies. The NoExp assimilation
439 represented the situation where only observations across environmental gradients were available.
440 Second, we compared the parameterization of the ASW, soil fertility, and atmospheric CO₂
441 environmental modifiers from the Base to the NoExp assimilation. The modifiers equations are
442 described in Supplemental Material Section 1.2 and 1.3. Third, we repeated the same
443 independent validation exercise for the 160 FMRC plots as described above for the Base
444 assimilation. Fourth, we predicted the treatment plots in the irrigated, drought, nutrient addition
445 (only plots where FR was assumed to be 1), and elevated CO₂ plots. As for the Base
446 assimilation, we used a t-test to compare the experimental response between the NoExp
447 assimilation and observed and between the NoExp and Base assimilations. Since the
448 experimental treatments were not used in the optimization, this was an independent evaluation of
449 predictive capacity.

450

451 **2.6 Regional predictions with uncertainty**

452 To demonstrate the capacity of the data assimilation system to create regional predictions with
453 uncertainty, we simulated the regional response to a decrease in precipitation, an increase in
454 nutrient availability, and an increase in atmospheric CO₂ concentration, each as a single factor
455 change from a 1985-2011 baseline. Each prediction included uncertainty by integrating across
456 the parameter posterior distributions using a Monte-Carlo sample of the parameter chains. Our
457 region corresponded to the native range of loblolly pine and used the HUC12 (USGS 12-digit
458 Hydrological Unit Code) watershed as the scale of simulation. For each HUC12 in the region we
459 used the mean SI, 30-year mean annual temperature, ASW aggregated to the HUC12 level, and
460 monthly meteorology from Abatzoglou (2013) as inputs (Figure 3). The SI of each HUC12 was
461 estimated from biophysical variables in the HUC12 using the method described in Sabatia and
462 Burkhart (2014). This SI corresponded to an estimated SI for stands without intensive
463 silvicultural treatments or advanced genetics of planted stock.

464

465 To sample parameter uncertainty, we randomly drew 500 samples from the Base assimilation
466 MCMC chain and simulated forest development from a 1985 planting to age 25 in 2011 in each
467 HUC. We chose age 25 as the final age because it is a typical age of harvest in the region. For
468 each sample, we repeated the regional simulation with 1) a 30% reduction in precipitation, 2) FR
469 set to 1, and 3) atmospheric CO₂ increased by 200 ppm. Within a parameter sample, we
470 calculated the percent change in stem biomass at age 25 between control simulation and the three
471 simulations with the environmental changes. We focused our regional analysis on the
472 distribution of the percent change in stem biomass.

473

474 **3 Results**

475 **3.1 Data assimilation evaluation**

476 Our multi-site, multi-experiment, multi-data stream DA approach (Base assimilation) increased
477 confidence in the model parameters (Table 5). Averaged across parameters, the posterior 99%
478 quantile range from the Base assimilation was 60% less than the prior range. The largest
479 reduction in parameter uncertainty was for the parameters associated with light-use efficiency
480 (α) and the conversion of GPP to NPP (γ), which on average had ranges that were 85% lower
481 in the posterior than the prior. Parameters associated with allocation and allometry had a 63%
482 reduction in the range while parameters associated with mortality processes had 70% reduction
483 in the range. Parameters associated with environmental modifiers had the least reduction in the
484 range with a 40% decrease. In addition to the parameters associated with the 3-PG model, the
485 model process error parameters for each data stream were well constrained with large reductions
486 in the range (> 99% decrease; Supplemental Material Table 2)

487

488 The Base assimilation reliably predicted data from the regionally distributed non-manipulated
489 plots that were not used in the optimization. The mean bias in stem biomass of the cross-
490 validation was -3.7 % and the RMSE was 21.8 Mg ha⁻¹ (Figure 4a). Furthermore, the response of
491 stem biomass to irrigation (df = 7, p = 0.18), nutrient addition (df = 26, p = 0.29), and elevated
492 CO₂ (df = 4, p = 0.43) was not significantly different between the observed and the Base
493 assimilation (Figure 5). The Base assimilation was significantly more sensitive to drought than
494 observed (n = 31, p < 0.001; Figure 5).

495

496 The plots at the Duke Forest study had a higher carrying capacity of stem biomass before self-
497 thinning (WSx1000), smaller self-thinning parameter (ThinPower), and lower allocation to
498 coarse root (pCRS) than values optimized from the other plots across the region (Table 6). The
499 DA approach without these three study specific parameters (NoDkPars) predicted significantly
500 lower accumulation of stem biomass in response to elevated CO₂ than observed (df = 4, p =
501 0.002; Figure 5). The NoDKPars assimilation optimized the CO₂ fertilization parameter
502 (fCalpha700) to a value that predicted 45% less light-use efficiency at 700 ppm (1.13 in
503 NoDKPar vs. 1.33 in Base; Table 6) than the Base assimilation.

504

505 3.2 Sensitivity to inclusion of ecosystem experiments

506

507 Excluding the experimental treatments from the data assimilation did not strongly influence the
508 predictive capacity of the model. The RMSE validation plots in NoExp assimilation decreased
509 slightly compared to Base assimilation (21.8 to 18.0 Mg ha⁻¹) while the bias slightly increased (-
510 3.7 to -4.1%)(Figure 4b). Excluding the experimental treatments resulted in a significantly lower
511 response of stem biomass to elevated CO₂ than observed (df = 4, p < 0.001; Figure 5).

512 Furthermore, there was a slight negative response of stem biomass to CO₂ in the NoExp
513 assimilation because the parameter governing the change in foliage allocation at elevated CO₂
514 (fCpFS700) was unconstrained by observations (Table 6). This led to convergence on the lower
515 bound of the prior distribution (0.5) where foliage allocation decreased with increased
516 atmospheric CO₂. The predictions of irrigation, drought, and nutrient addition experiments were
517 not significantly different between the Base and NoExp assimilations (Figure 5).

518

519 The parameters and associated response functions in the 3-PG for nutrients, ASW, and
520 atmospheric CO₂ differed between the Base and NoExp assimilations (Figure 6). First, the
521 parameterization of the soil fertility rating (FR) showed a stronger dependence on SI in the
522 NoExp assimilation than in the Base assimilation (Figure 6a). For a given SI there was a lower
523 FR, thus stronger nutrient limitation, when experimental treatments were excluded from
524 assimilation. Second, the parameterization of the function relating photosynthesis and canopy
525 conductance to ASW resulted in lower photosynthesis and maximum conductance when soil
526 available water was less than 50% in the NoExp than Base assimilations (Figure 6b). Finally, the
527 response of photosynthesis to atmospheric CO₂ was functionally zero in the NoExp assimilation,
528 thus highlighting the importance of the elevated CO₂ treatments in the Duke FACE study for
529 constraining the parameterization of the CO₂ response function (Figure 6c).

530

531 **3.3 Regional predictions with uncertainty**

532 Regionally (i.e., the native range of loblolly pines), stem biomass at age 25 ranged from 52 Mg
533 ha⁻¹ to 292 Mg ha⁻¹ with the most productive areas located in the coastal plains and the interior of
534 Mississippi and Alabama (Figure 7a). The least productive locations were the western and
535 northern extents of the native range. The width of the 95% quantile interval for each HUC12 unit
536 ranged from 6.2 to 29.8 Mg ha⁻¹ with largest uncertainty located in most the productive HUC12
537 units and in the far western extent of the region (Figure 7b).

538

539 The predicted change in stem biomass at age 25 from an additional 200 ppm of atmospheric CO₂
540 (over the 1985-2011 concentrations) was similar to the change associated with a removal of
541 nutrient limitation (by setting FR = 1) (Figure 8a,c). The median change associated with

542 elevated CO₂ for a given HUC12 unit ranged from 19.2 to 55.7% with a regional median of
543 21.7% (Figure 8a). The change associated the removal of nutrient limitation ranged from 6.9 to
544 303.7% for a given HUC12 unit, with regional median of 24.1% (Figure 8b). The response to
545 elevated CO₂ was more consistent across space than the response to nutrient addition. The
546 largest potential gains in productivity from nutrient addition were predicted in central Georgia,
547 the northern extent of the region, and the western extents, areas with the lowest SI (Figure 3).

548

549 Stem biomass was considerably less responsive to a 30% decrease in precipitation, than to
550 nutrient addition and an increase in atmospheric CO₂. The median change in stem biomass when
551 precipitation was reduced from the 1985-2011 levels ranged from -11.6 to -0.1% for a given
552 HUC12 unit with a regional median of -5.1% (Figure 8c). Central Georgia was the most
553 responsive to precipitation reduction reflecting the relatively low annual precipitation and warm
554 temperatures (Figure 3).

555

556 For a given location, the predicted response to elevated CO₂ had larger uncertainty than the
557 predicted response to precipitation reduction and nutrient limitation removal (Figure 8c,d,f). The
558 uncertainty, defined as the width of the 95% quantile interval, was consistent across the region
559 for the response to elevated CO₂ (Figure 8b). The uncertainty in the response to precipitation
560 reduction and nutrient limitation removal was largest in the regions with the largest predicted
561 change (Figure 8df).

562

563 **4 Discussion**

564 Using DA to parameterize models for predicting ecosystem change requires disentangling the

565 vegetation responses to temperature, precipitation, nutrients, and elevated CO₂. To address this
566 challenge, we introduced a regional-scale hierarchical Bayesian approach (DAPPER) that
567 assimilated data across environmental gradients and ecosystem manipulation experiments into a
568 modified version of the 3-PG model. Furthermore, we synthesized observations of carbon stocks,
569 carbon fluxes, water fluxes, vegetation structure, and vegetation dynamics that spanned 35 years
570 of forest research in a region (Table 1, Figure 1) with large and dynamic carbon fluxes (Lu et al.,
571 2015). By combining the DAPPER system with the regional set of observations, we were able to
572 estimate parameters in a model with high predictive capacity (Figure 4) and with quantified
573 uncertainty on parameters (Table 5) and regional simulations (Figures 7 and 8).

574

575 Our hierarchical approach (Equation 7) was designed to partition uncertainty among parameters,
576 model process, and measurements (Hobbs and Hooten, 2015). Separating the parameter and
577 process uncertainty is required to estimate prediction intervals, as prediction intervals only
578 include parameter and process errors (Dietze et al., 2013; Hobbs and Hooten, 2015). Previous
579 forest ecosystem DA efforts have either focused on parameter uncertainty, by using
580 measurement uncertainty as the variance term in a Gaussian cost function (Bloom and Williams,
581 2015; Keenan et al., 2012; Richardson et al., 2010) or on total uncertainty by directly estimating
582 the Gaussian variance term (Ricciuto et al., 2008). Our approach allowed the estimation of the
583 probability distribution of forest biomass before uncertainty is added through measurement.

584 Considering that the method of DA can potentially have a large influence on posterior parameter
585 distributions (Trudinger et al., 2007), future research should focus on comparing the hierarchical
586 approach presented here to other approaches by using the same data constraints with alternative
587 cost functions.

588

589 **4.1 Sensitivity to inclusion of ecosystem experiments**

590 The most important experimental manipulation for constraining model parameters was the Duke
591 FACE CO₂ fertilization study because the CO₂ fertilization parameters (fC_{alpha}700 and
592 fC_{pFS}700) converged on the lower bounds of their prior distributions when the experiments
593 were excluded from the assimilation. In contrast, excluding the nutrient fertilization, drought,
594 and irrigation studies did not substantially alter the predictive capacity of the model. This
595 finding suggests that data assimilation using plots across environmental gradients alone can
596 constrain parameters associated with water and nutrient sensitivity. However, regardless of
597 whether the experiments were included in the assimilation, the optimized model predicted higher
598 sensitivity to drought than observed, highlighting that future studies should focus on improving
599 the sensitivity to drought.

600

601 The 3-PG model included a highly-simplified representation of interactions between the water
602 and carbon cycles that resulted in parameterizations that may contain assumptions that require
603 additional investigation. First, transpiration was modeled as a function of a potential canopy
604 transpiration that occurred if leaf area was not limiting transpiration. The LAI at which leaf area
605 was no longer limiting was a parameter that was optimized (LAI_{g_{cx}} in Table 5), resulting in a
606 value of 2.2. Interestingly, this optimized value is consistent with the scant literature on this
607 topic. In their analysis of multi-year measurements of transpiration in loblolly pine, Phillips and
608 Oren (2001) observed that transpiration per unit leaf area was relatively insensitive to increases
609 in leaf area above LAI of approximately 2.5. Iritz and Lindroth (1996) reviewed transpiration
610 data from a range of crop species and found only small increases in transpiration above LAI of 3-

611 4. These authors suggest that the threshold-type responses observed were related to the range of
612 LAI at which self-shading increases most rapidly, therefore limiting increases in transpiration.
613 The resulting model behavior of "flat" transpiration above 2.2 LAI, with gradually decreasing
614 photosynthesis above that value, results in increasing water use efficiency at higher LAI values.
615 Second, the relationship between relative ASW and the modifier of photosynthesis and
616 transpiration predicted a modifier value greater than zero when the relative ASW was zero. This
617 resulted in positive values from photosynthesis and transpiration when the average ASW during
618 the month was zero. In practice, the monthly ASW was rarely zero during simulations, which
619 presents a challenge constraining the shape of the ASW modifier. The priors for the two ASW
620 modifiers (SWconst and SWpower) had ranges that permitted the modifier to be zero. Therefore,
621 additional data are likely needed during very dry conditions to develop a more physically based
622 parameterization. Alternatively, the parameterization of a non-zero soil moisture modifier at zero
623 ASW may be due to trees having access to water at soil depths deeper than the top 1.5 m of soil
624 represented by the bucket in 3-PG. Overall, it is important to view the parameterization presented
625 here as a phenomenological relationship that is consistent with observations from drought and
626 irrigation experiments as well as observations across regional gradients in precipitation.

627

628 Constraining the sensitivity to atmospheric CO₂ differs from constraining the sensitivity to ASW
629 because, unlike the multiple constraints on water sensitivity (drought, irrigation, and gradient
630 studies), environmental conditions created by the few elevated CO₂ plots provided unique
631 constraint on parameters. Our finding demonstrated that DA efforts should test for bias in
632 unique ecosystem experiments before finalizing a set of model parameters used in optimization.
633 In particular, we found that the parameter governing the photosynthetic response to elevated CO₂

634 (fCalpha700) was substantially lower when all parameters were assumed to be shared across all
635 plots than when the CO₂ fertilization experiment was allowed to have unique parameters. The
636 need for the three unique parameters at the Duke FACE study parameters can be explained by
637 the constraint provided by multiple data streams and multiple plots. An assumption of the model
638 was that an increase in stem biomass caused a decrease stem density through self-thinning,
639 unless the average tree stem biomass was below a parameterized threshold (WSx1000).
640 Therefore, an increase in photosynthesis and stem biomass through CO₂ fertilization could cause
641 a decrease in stem density. For a single study, it is straightforward to simultaneously fit the CO₂
642 fertilization and self-thinning parameters to fit stem biomass and stem density observations for
643 the site. However, regional DA presents a challenge because the self-thinning parameters are
644 well constrained by the stem biomass and stem density observations across the region but the
645 CO₂ fertilization parameters are not. As a result of the regional DA, the self-thinning parameters
646 caused a stronger decrease in stem density than observed in the Duke FACE study. Therefore,
647 the optimization favored a solution where there was a lower response to CO₂, thus a smaller
648 decrease in stem density. Allowing the Duke FACE study to have unique self-thinning
649 parameters that resulted in lower rates of self-thinning and allowed for simulated stem biomass
650 to respond to CO₂ in a way that matched the observations without penalizing the optimization by
651 degrading the fit to the stem density.

652

653 Our finding that the Duke FACE study required unique self-thinning parameters to reduce bias in
654 the simulated stem biomass suggests that when using DA to optimize parameters that are shared
655 across plots, careful examination of prediction bias in key sites that provide unique constraint on
656 certain parameters (like the Duke FACE) is critical. Based on this example, we suggest that DA

657 efforts using multiple studies and multiple experiment types identify whether particular
658 experiments at limited number of sites have the potential to uniquely constrain specific
659 parameters. In this case, additional weight or site-specific parameters may be needed to avoid
660 having the signal of the unique experiment overwhelmed by the large amount of data from the
661 other sites and experiments. Additionally, the finding suggests that multi-site DA should
662 consider using hierarchical approaches to predicting mortality, particularly because mortality is
663 often not simulated as mechanistically as growth. A hierarchical approach, where each plot has a
664 set of mortality parameters that are drawn from a regional distribution, could avoid having
665 unexplained variation in mortality rates lead to bias in the parameterization of growth related
666 processes (i.e., growth responses to CO₂, drought, nutrient fertilization, etc.). The hierarchical
667 approach to mortality could also highlight patterns in mortality rates across a region and allow
668 for additional investigations in the mechanisms driving the patterns.

669

670 **4.2 Regional predictions with uncertainty**

671 Our predictions of how stem biomass responses to elevated CO₂, nutrient addition, and drought
672 were designed to illustrate the capacity of the DAPPER approach to simulate the uncertainty in
673 future predictions. By using DA, our regional predictions and the uncertainty are consistent with
674 observations but are associated with key caveats. First, only parameter uncertainty was
675 presented in the regional simulations. There is additional uncertainty associated with model
676 process error. We showed the parameter uncertainty because it isolated the capacity to
677 parameterize the individual environmental response functions in the model. Second, the
678 response to drought may be too strong because of the bias in the model predictions of the
679 drought studies. However, there is potential that the drought studies underestimated the

680 sensitivity to ASW since they are relatively short term (< 5 years) and manipulate local ASW
681 without manipulating large scale ASW (i.e., regional water tables). Third, the large responses to
682 N fertilization at the western and northern extents of the study region may be too high. The large
683 responses are attributed to the low SI and the low predicted site fertility index (FR_p). The low SI
684 may be attributable to water limitation and temperature limitation that is not fully accounted for
685 in the parameterization. Additional nutrient addition experiments in the northern and western
686 extent along with further development of the representation of nutrient availability in the 3-PG
687 model may allow for a more robust representation of soil fertility. Finally, the baseline fertility
688 used in our regional analysis was derived from an empirical model of SI that was developed
689 using field plots with minimal management (Sabatia and Burkhart, 2014). Subsequently our
690 estimate of baseline fertility is likely on the low end of forest stands currently in production and
691 the response to nutrient addition may be higher than a typical stand under active management.

692

693 **5 Conclusions**

694 DA is increasingly used for developing predictions from ecosystem models that include
695 uncertainty estimation, due to its ability represent prior knowledge, integrate observations into
696 the parameterization, and estimate multiple components of uncertainty, including observation,
697 parameter, and process representation uncertainty (Dietze et al., 2013; Luo et al., 2011; Niu et
698 al., 2014). Our application of DA to loblolly pine plantations of the southeastern U.S
699 demonstrated that these ecosystems are well suited as a test-bed for the development of DA
700 techniques, particularly techniques for assimilating ecosystem experiments. We found that
701 assimilating observations across environmental gradients can provide substantial constraint on
702 many model parameters but that ecosystem manipulative experiments, particularly elevated CO_2

703 studies, were critical for constraining parameters associated forest productivity in a more CO₂
704 enriched atmosphere. This highlights the importance of whole-ecosystem manipulation CO₂
705 experiments for helping to parameterize and evaluate ecosystem models. Finally, we present an
706 approach for the development of future predictions of forest productivity for natural resource
707 managers that leverage a rich dataset of integrated ecosystem observations across a region.
708

709 **6 Data availability**

710 Observations used in the DA can be found in the following: Duke FACE study can be found in
711 McCarthy et al. (2010), the PINEMAP studies are available through the TerraC database
712 (<http://terrac.ifas.ufl.edu>), the US-DK3 eddy-flux tower data are available through the Ameriflux
713 database (<http://ameriflux-data.lbl.gov>), the Waycross data can be found in Bryars et al. (2013),
714 the US-NC2 data are available upon request from Asko Noormets, the FMRC and FPC are
715 available through membership with the cooperatives. The parameter chains and 3-PG model
716 code are available upon request from R. Quinn Thomas.

717

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734

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Table 1. Regional observational data streams used in data assimilation.

Data stream	Measurement frequency	Measurement or estimation technique	Uncertainty	Stream ID for Table 3
Foliage biomass (Pine)	Annual or less	Allometric relationship	Based on propagating the allometric model uncertainty in Gonzalez-Benecke et al. 2014. Varied by observation.	1
Foliage biomass (hardwood)	Annual or less	Allometric relationship	Assumed zero	2
Stem biomass (pine)	Annual or less	Allometric relationship	Based on propagating the allometric model uncertainty in Gonzalez-Benecke et al. 2014. Varied by observation.	3
Stem biomass (hardwood)	Annual or less	Allometric relationship	Assumed zero	4
Coarse root biomass (combined)	Annual or less	Allometric relationship	Assumed zero*	5
Fine root biomass (combined)	Annual or less	Allometric relationship	SD = 10% of observation	6
Foliage biomass production (combined)	Annual	Litterfall traps	SD = 10% of observation	7
Fine root biomass production (combined)	Annual	Mini-rhizotrons	SD = 10% of observation	8
Pine stem density	Annual or less	Counting individuals	1% (assumed small)	9
Leaf area index (pine)	Monthly to annual	Litter traps or LI 2000	SD = 10% of observation	10
Leaf area index (hardwood)	Monthly to annual	Litter traps or LI 2000	SD = 10% of observation	11
Leaf area index (combined)	Only used if not separated into pine and hardwood	Litter traps or LI 2000	SD = 10% of observation	12
Gross Ecosystem Production	Monthly	Modeled from flux eddy-covariance net	SD = 10% of observation	13

Evapotranspiration	Monthly	ecosystem exchange Eddy- covariance	SD = 10% of observation	14
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931 *the relatively low number of observations prevented convergence when using the observational
932 uncertainty model so observational uncertainty was assumed to be zero to allow convergence.
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Table 2. Descriptions of the studies used in data assimilation.

Study name	Number of locations	Number of plots per site	Experimental treatments (plots)	Data streams (Table 2)	Measurement Years	Measurement Stand Ages (years)	Reference
FMRC ¹ Thinning Study	163	1	None	1, 3,9	1981 - 2003	8 - 30	Burkhart et al. (1985)
FPC ² Region-wide 18	18	2	Nutrient addition	1, 3,9	2011-2014	12-21	Albaugh et al. (2015)
PINEMA P ³	4	16	Nutrient addition, 30% throughfall, Nutrient x throughfall	1, 3,9	2011-2015	3 – 13	Will et al. (2015)
Waycross	1	2	Nutrient addition	3,9,10	1991-2010	4-23	Bryars et al. (2013)
SETRES ⁴	1	16	Nutrient addition, irrigation, nutrient x irrigation	1,3,5,6, 9,10	1991-2006	8 - 23	Albaugh et al. (2004)
Duke FACE ⁵ and US-DK3 Flux	1	12	CO ₂ , nutrient addition, CO ₂ x nutrient addition	2,3,4,5, 6,7,8,9, 10,11,1 3,14	1996-2004	13-22	McCarthy et al. (2010); Novick et al. (2015)
NC2 Flux	1	1	None	2,3,4,5, 6,7,9,1 0,11,12 ,13,14	2005-2014	12-22	Noormets et al. (2010)
Total	187	294			1981 - 2014	4 - 30	

934 ¹Forest Modeling Research Cooperative; ² Forest Productivity Cooperative; ³ Pine Integrated
 935 Network: Education, Mitigation, and Adaptation project (PINEMAP); ⁴ Southeast Tree Research

936 and Education Site; ⁵ Free Air Carbon Enrichment
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Table 3. The prior distributions of all 3-PG model parameters optimized using data assimilation.

Parameter	Parameter description	Units	Prior distribution	Prior parameters	Reference for prior (see footnote)
Allocation and structure					
pFS2	Ratio of foliage to stem allocation at stem diameter = 2 cm	-	uniform	min = 0.08 max = 1.00	uninformed
pFS20	Ratio of foliage to stem allocation at stem diameter = 20 cm	-	uniform	min = 0.10 max = 1.00	uninformed
pRF	Ratio of fine roots to foliage allocation	-	uniform	min = 0.05 max = 2.00	uninformed
pCRS	Ratio of coarse roots to stem allocation	-	uniform	min = 0.15 max = 0.35	1
SLA0	Specific leaf area at stand age 0	m ² kg ⁻¹		mean = 5.53 sd = 0.44	2
SLA1	Specific leaf area for mature aged stands	m ² kg ⁻¹	normal	mean = 3.58 sd = 0.11	2
tSLA	Age at which specific leaf area = 0.5(SLA0 + SLA1)	Years	normal	mean = 5.97 sd = 2.15	2
fCpFS700	Proportional decrease in allocation to foliage between 350 and 700 ppm CO ₂	-	uniform	min = 0.50 max = 1.00	uninformed
StemConstant	Constant in stem mass vs. diameter relationship	-	normal	mean = 0.022 sd = 0.005	3
StemPower	Power in stem mass vs. diameter relationship	-	normal	mean = 2.77 sd = 0.2	3
Canopy photosynthesis, autotrophic respiration, and transpiration					
alpha	Canopy quantum efficiency (pines)	mol C mol PAR ⁻¹	uniform	min = 0.02 max = 0.06	uninformed
y	Ratio NPP/GPP	-	uniform	min = 0.30 max = 0.65	4
MaxConduct	Maximum canopy conductance	m s ⁻¹	uniform	min = 0.005 max = 0.03	2
LAIgcx	Canopy LAI for maximum canopy conductance	-	uniform	min = 2 max = 5	2,5,6
Environmental modifiers of photosynthesis and transpiration					

kF	Reduction rate of production per degree Celsius below zero	-	normal	mean = 0.18 sd = 0.016	2
Tmin	Minimum monthly mean temperature for growth	°C	normal	mean = 4.0 sd = 2.0	2,5,6
Topt	Optimum monthly mean temperature for growth	°C	normal	mean = 25.0 sd = 2.0	2,5,6
Tmax	Maximum monthly mean temperature for growth	°C	normal	mean = 38.0 sd = 2.0	2,5,6
SWconst	Moisture ratio deficit when downregulation is 0.5	-	uniform	min = 0.01 max = 1.8	uninforme d
SWpower	Power of moisture ratio deficit	-	uniform	min = 1 max = 13	uninforme d
CoeffCon d	Defines stomatal response to VPD	mbar ⁻¹	normal	mean = 0.041 sd = 0.003	2
fAlpha7 00	Proportional increase in canopy quantum efficiency between 350 and 700 ppm CO ₂	-	uniform	min = 1.00 max = 1.8	uninforme d
MaxAge	Maximum stand age used to compute relative age	Years	uniform	min = 16 max = 200	uninforme d
nAge	Power of relative age in fage	-	uniform	min = 0.2 max = 4.0	uninforme d
rAge	Relative age to where fage = 0.5	-	uniform	min = 0.01 max = 3.00	uninforme d
FR1	Fertility rating parameter 1 (mean annual temperature coefficient)	-	uniform	min = 0.0 max = 1.0	uninforme d
FR2	Fertility rating parameter 2 (site index age 25 coefficient)	-	uniform	min = 0.0 max = 1.0	uninforme d
Mortality					
wSx1000	Maximum stem mass per tree at 1000 trees/ha	kg tree ⁻¹	normal	mean = 235 sd = 25	2,5,6
ThinPow er	Power in self thinning law	-	uniform	min = 1.0 max = 2.5	2,5,6
ms	Fraction of mean stem biomass per tree on dying trees	-	uniform	min = 0.1 max = 1.0	uninforme d
Rttover	Average monthly root turnover rate	month ⁻¹	uniform	min = 0.017 max = 0.042	7

MortRate	Density independent mortality rate (pines)	month ⁻¹	uniform	min = 0.0002 max = 0.004	uninformed
<hr/>					
Understory hardwoods					
alpha_h	Canopy quantum efficiency (understory hardwoods)	mol C mol PAR ⁻¹	uniform	min = 0.005 max = 0.07	uninformed
pFS_h	Ratio of foliage to stem partitioning (understory hardwoods)	-	uniform	min = 0.2 max = 3.0	uninformed
pR_h	Ratio of foliage to fine roots (understory hardwoods)	-	uniform	min = 0.05 max = 2	uninformed
SLA_h	Specific leaf area (understory hardwoods)	m ² kg ⁻¹	normal	mean = 16 sd = 3.8	8
fAlpha700_h	Proportional increase in canopy quantum efficiency between 350 and 700 ppm CO ₂ (understory hardwood)	-	uniform	min = 1.00 max = 2.5	uninformed
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¹Albaugh et al., 2005; ²Gonzalez-Benecke et al., 2016; ³Gonzalez-Benecke et al., 2014 ⁴DeLucia et al., 2007; ⁵Bryars et al., 2013; ⁶Subedi et al., 2011; ⁷Matamala et al., 2003; ⁸LeBauer et al., 2010; uninformed priors had large, ecologically reasonable bounds.

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Table 4. Description of the different data assimilation approaches used.

Simulation Name	Treatments included in assimilation	Number of plots
Base	All plots and experiments in the region were used simultaneously. Includes unique pCRS, wSx1000, and ThinPower parameters for plots in the Duke FACE study	294
NoExp	Same as Base assimilation but excluding all plots with experimental manipulations. Includes control plots that are part of experimental studies.	208
NoDkPars	Same as Base assimilation but without pCRS, wSx1000, and ThinPower parameter for plots in the Duke FACE and US-DK3 studies	294

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Table 5. The optimized medians, range of the 99% quantile intervals of the posterior distributions and the 99% quantile range for priors with normally distributed priors or the range of the upper and lower bounds for priors with uniform distributions.

Parameter	Posterior median	Posterior 99% C.I. range	Prior range	Posterior/Prior Range
Allocation and structure				Parameter group mean = 0.38
pFS2	0.58	0.55 - 0.61	0.08 – 1.00	0.06
pFS20	0.57	0.55 - 0.59	0.10 – 1.00	0.05
pR	0.11	0.07 - 0.15	0.05 – 2.00	0.04
pCRS	0.26	0.25 - 0.27	0.15 - 0.35	0.11
pCRS (Duke)	0.21	0.18 - 0.23	0.15 - 0.35	0.20
SLA0	8.44	7.67 - 9.25	4.4 - 6.66	0.70
SLA1	2.84	2.72 - 2.96	3.59 - 4.16	0.43
tSLA	4.13	3.88 - 4.41	0.43 - 11.51	0.05
fCpFS700	0.74	0.60 - 0.90	0.50 – 1.00	0.60
StemConst	0.022	0.009 - 0.035	0.009 - 0.035	1.00
StemPower	2.78	2.29 - 3.27	2.25 - 3.29	0.95
Canopy photosynthesis, autotrophic respiration, and transpiration				Parameter group mean = 0.14
alpha	0.029	0.026 - 0.031	0.02 - 0.06	0.14
y	0.50	0.47 - 0.53	0.30 - 0.65	0.15
MaxCond	0.011	0.01 - 0.012	0.005 - 0.03	0.09
LAIg _{cx}	2.2	2.0 - 2.48	2.0 - 5.0	0.16
Environmental modifiers of photosynthesis and transpiration				Parameter group mean = 0.61
kF	0.16	0.12 - 0.2	0.14 - 0.22	1.04
T _{min}	-5.56	-8.88 - -2.69	-1.15 - 9.15	0.60
T _{opt}	23.42	21.1 - 26.31	19.85 - 30.15	0.51
T _{max}	39.56	34.71 - 44.39	32.85 - 43.15	0.94
SWconst	1.09	0.91 - 1.56	0.01 - 1.8	0.36
SWpower	8.86	3.39 - 12.98	1.00 – 13.00	0.80
CoeffCond	0.036	0.029 - 0.043	0.034 - 0.048	0.91
fC _{alpha} 700	1.33	1.18 - 1.52	1.0 - 1.80	0.43
MaxAge	151.5	54.4 - 199.6	16.0 - 200.0	0.79
nAge	3.35	1.77 - 3.99	1.00 – 4.00	0.74
rAge	2.25	0.81 - 2.99	0.01 – 3.00	0.73
FR1	0.073	0.061 - 0.086	0.00 – 1.00	0.03
FR2	0.17	0.15 - 0.19	0.0 – 1.0	0.04
Mortality				Parameter group mean = 0.37
wS _x 1000	176.9	169.6 - 184.4	165.6 - 294.4	0.15
wS _x 1000 (Duke)	243.3	196.89 - 305.02	165.6 - 294.4	0.76

ThinPower	1.68	1.60 - 1.78	1.00 - 2.5	0.12
ThinPowerv(Du ke)	1.26	1.00 - 1.85	1.00 - 2.5	0.56
mS	0.52	0.37 - 0.71	0.10 - 1.00	0.38
Rttover	0.023	0.017 - 0.031	0.017 - 0.042	0.55
MortRate	0.001	9e-04 - 0.0011	2e-04 - 0.004	0.06
Understory hardwoods				Parameter group mean = 0.28
alpha_h	0.02	0.02 - 0.02	0.005 - 0.07	0.01
pFS_h	1.78	1.54 - 2.06	0.2 - 3.0	0.19
pR_h	0.21	0.06 - 0.43	0.05 - 2.00	0.19
SLA_h	16.3	14.1 - 19.0	6.2 - 25.8	0.25
fCalpha700_h	1.84	1.58 - 2.17	1.0 - 2.50	0.74

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Table 6. Median and range of the 99% quantile intervals of the posterior distributions for the parameters in the NoExp and NoDkPars assimilations

Parameter	NoExp median	NoExp 99% range	NoDkPars median	NoDkPar 99%
Allocation and structure				
pFS2	0.63	0.61 - 0.68	0.57	0.55 - 0.60
pFS20	0.63	0.60 - 0.65	0.57	0.55 - 0.59
pR	0.11	0.06 - 0.16	0.11	0.08 - 0.15
pCRS	0.29	0.27 - 0.30	0.26	0.25 - 0.27
pCRS (Duke)	0.25	0.23 - 0.28	N/A	N/A
SLA0	7.47	6.57 - 8.41	8.56	7.73 - 9.32
SLA1	3.00	2.88 - 3.12	2.89	2.79 - 2.99
tSLA	4.75	4.30 - 5.26	4.12	3.90 - 4.38
fCpFS700	0.50	0.50 - 0.53	0.94	0.83 - 1.00
StemConst	0.022	0.01 - 0.04	0.02	0.01 - 0.04
StemPower	2.79	2.27 - 3.26	2.77	2.28 - 3.30
Canopy photosynthesis, autotrophic respiration, and transpiration				
alpha	0.030	0.028 - 0.033	0.029	0.026 - 0.031
y	0.48	0.45 - 0.51	0.49	0.46 - 0.52
MaxCond	0.017	0.015 - 0.021	0.011	0.011 - 0.012
LAIgcx	4.4	3.9 - 5.0	2.1	2.0 - 2.5
Environmental modifiers of photosynthesis and transpiration				
kF	0.15	0.11 - 0.20	0.16	0.11 - 0.20
Tmin	-7.8	-10.97 - -4.95	-6.04	-9.06 - -3.03
Topt	21.55	19.15 - 24.39	22.71	20.54 - 25.42
Tmax	40.56	36.51 - 45.62	39.82	35.62 - 44.56
SWconst	0.93	0.8 - 1.1	1.14	0.91 - 1.62
SWpower	6.27	2.98 - 11.49	7.99	3.29 - 12.95
CoeffCond	0.041	0.034 - 0.047	0.036	0.030 - 0.042
fCalpha700	1.01	1.00 - 1.06	1.15	1.10 - 1.25
MaxAge	152.84	54.18 - 199.5	152.0	49.2 - 199.3
nAge	3.36	1.93 - 3.99	3.36	1.89 - 3.99
rAge	2.26	0.80 - 2.99	2.24	0.83 - 2.99
FR1	0.12	0.09 - 0.14	0.08	0.07 - 0.09
FR2	0.20	0.16 - 0.24	0.17	0.15 - 0.19
Mortality				
wSx1000	191.6	180.2 - 210.2	181.32	173.26 - 196.32
wSx1000 (Duke)	235.1	175.0 - 297.5	N/A	N/A
ThinPower	1.76	1.61 - 1.92	1.59	1.46 - 1.72
ThinPower (Duke)	1.42	1.01 - 2.02	N/A	N/A

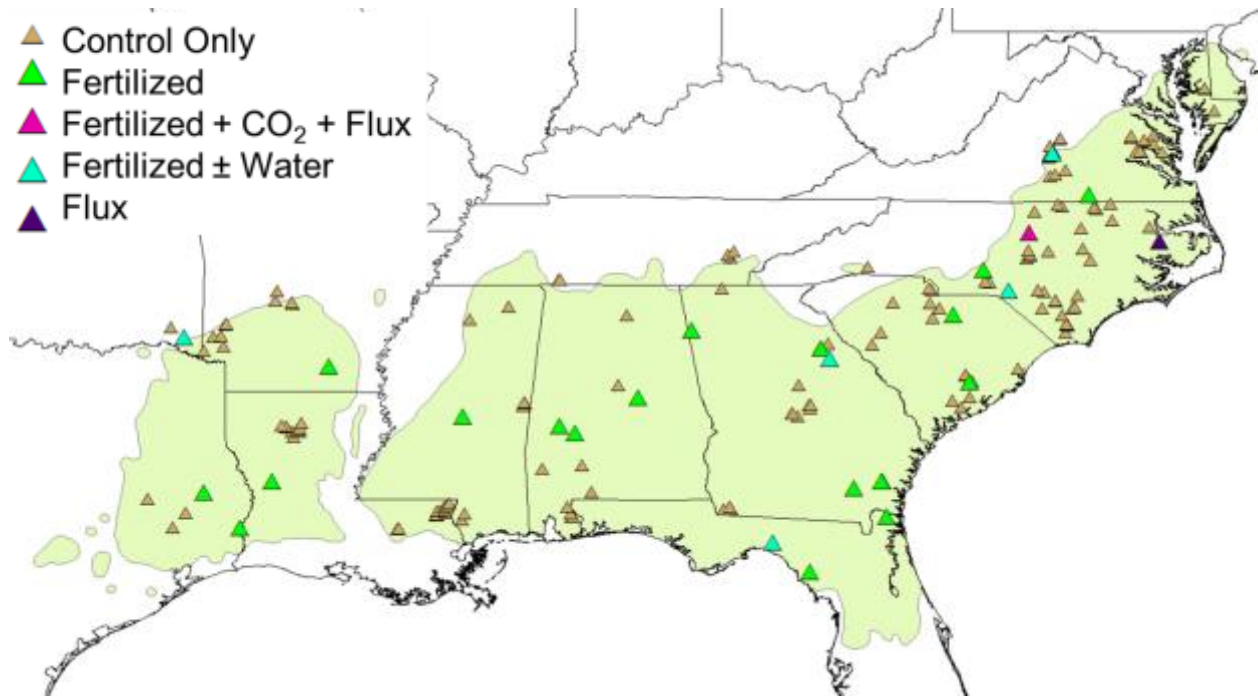
mS	0.54	0.33 - 0.80	0.5	0.25 - 0.71
Rttover	0.019	0.02 - 0.03	0.022	0.017 - 0.030
MortRate	0.0013	0.0011 - 0.0014	0.0011	9e-04 - 0.0013

Understory hardwoods

alpha_h	0.031	0.025 - 0.040	0.02	0.017 - 0.023
pFS_h	2.39	1.86 - 2.96	1.79	1.59 - 2.09
pR_h	0.25	0.05 - 0.67	0.21	0.06 - 0.41
SLA_h	12.37	9.96 - 15.07	16.42	14.37 - 18.55
fCalpha700_h	1.08	1.00 - 1.83	1.83	1.56 - 2.15

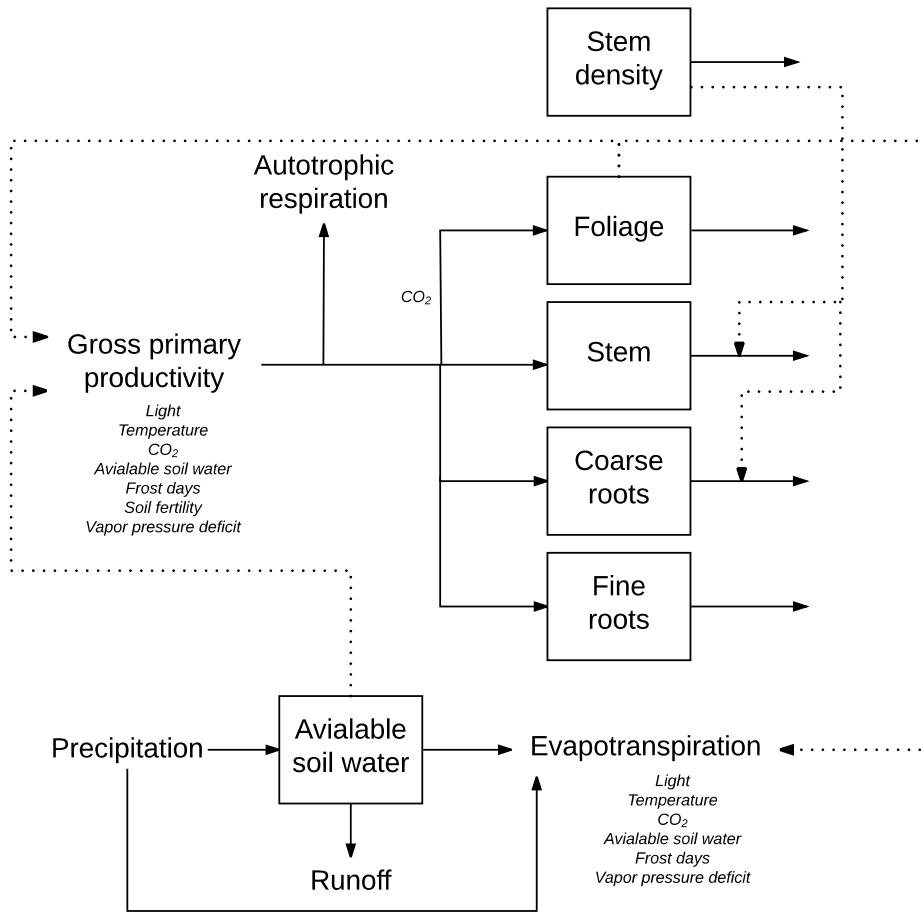
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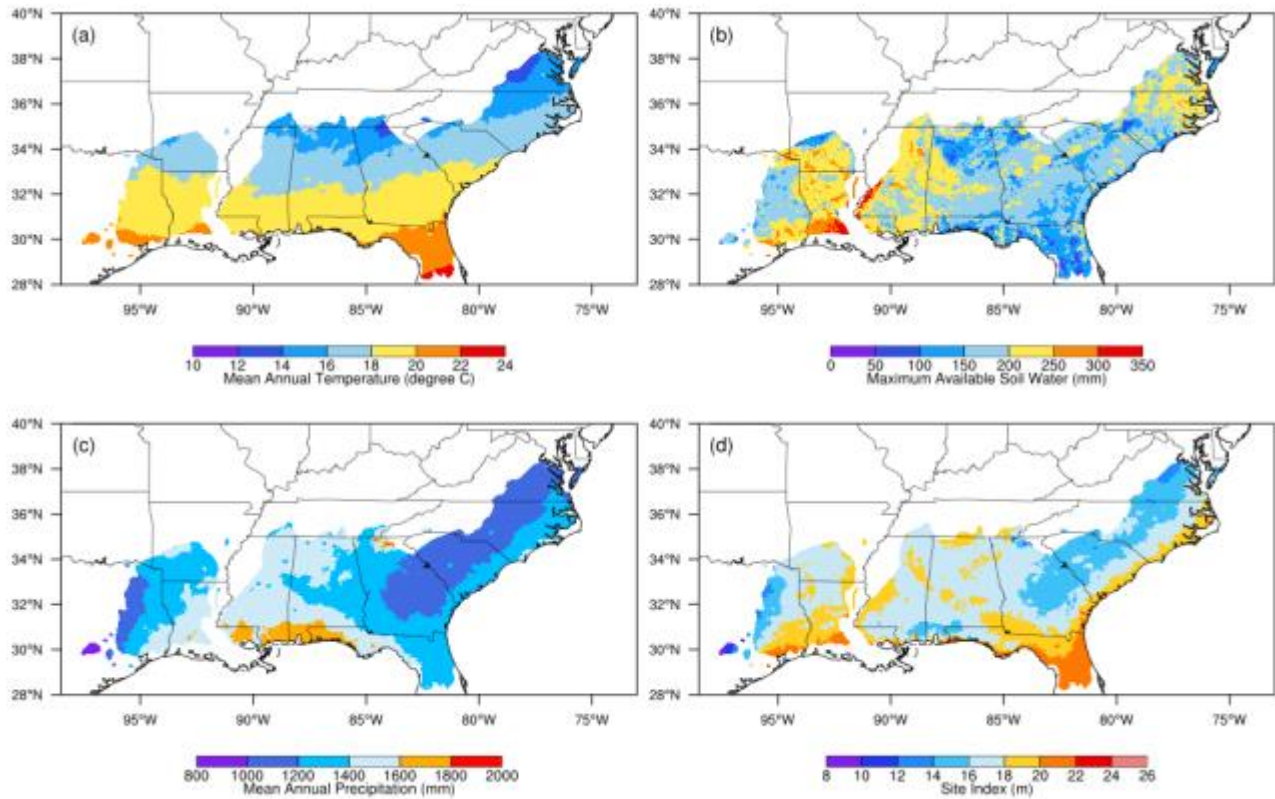
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 952 Figure 1. Map of loblolly pine distribution, plot locations used in data assimilation, and the
 953 experiment type associated with each plot. The control-only treatments were plots without any
 954 associated experimental treatment or flux measurements. Fertilized were plots with nutrient
 955 additions. CO₂ were plots with free-air concentration enrichment treatments. The flux treatments
 956 were plots with eddy-covariance measurements of ecosystem-scale carbon and water exchange.
 957 The water treatments included throughfall exclusion and irrigation experiments.
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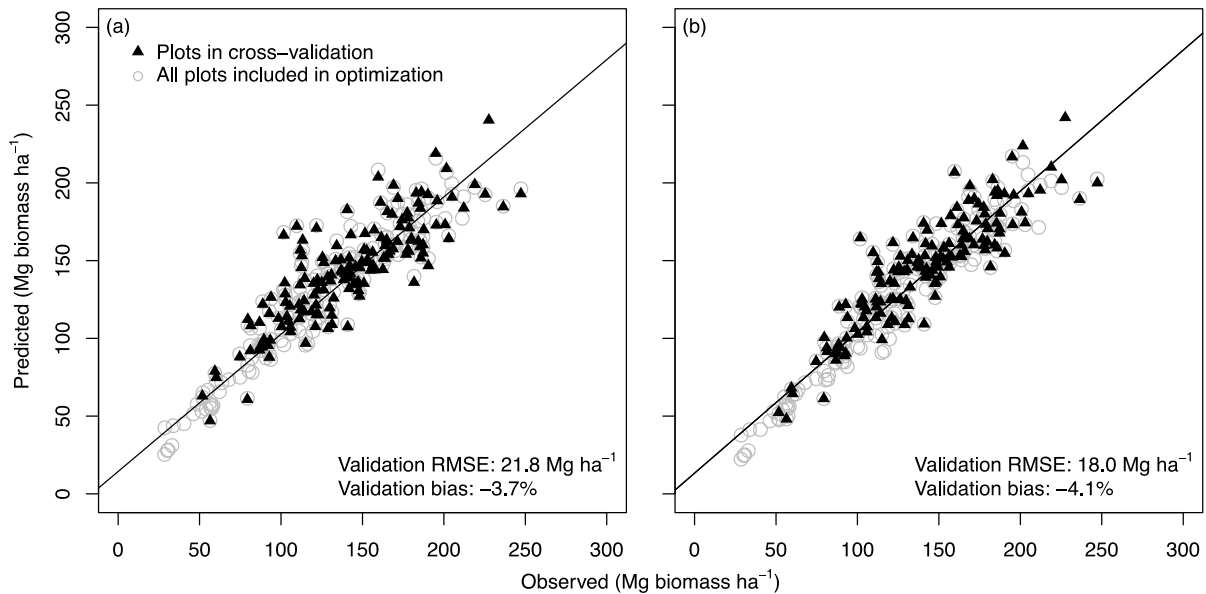
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Figure 2. A diagram of the monthly time-step 3-PG model used in this study. The stocks are represented by the boxes and the fluxes by the arrows. An influence of a stock on a flux that is not directly related to that stock is represented by the dotted lines. The environmental influences on a flux is described using italics. A description of the model can be found in the supplemental information.



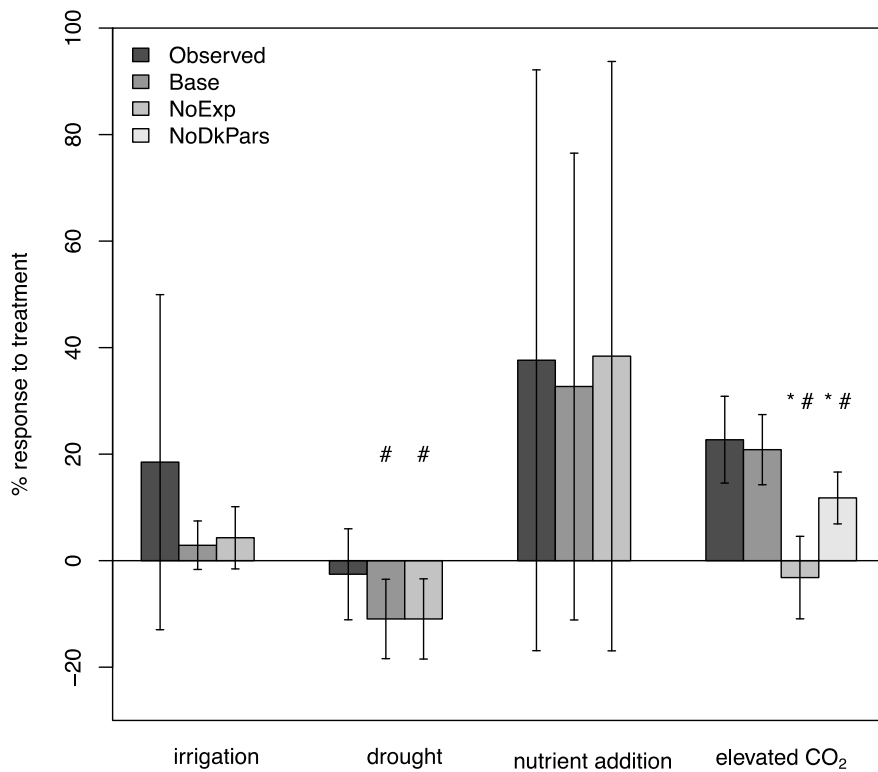
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 969 Figure 3. Key climatic and stand characteristic inputs to the regional 3-PG simulations: (a) Mean
 970 annual temperature (1979-2011) as a summary of the gradient in monthly temperature inputs
 971 used in simulations, (b) maximum available soil water for the top 1.5 meters of soil from
 972 SSURGO, (c) mean annual precipitation (1979-2011) as a summary of the gradient in monthly
 973 precipitation inputs used in simulations, and (d) site index. The area shown is the natural range of
 974 loblolly pine (*Pinus taeda* L.).

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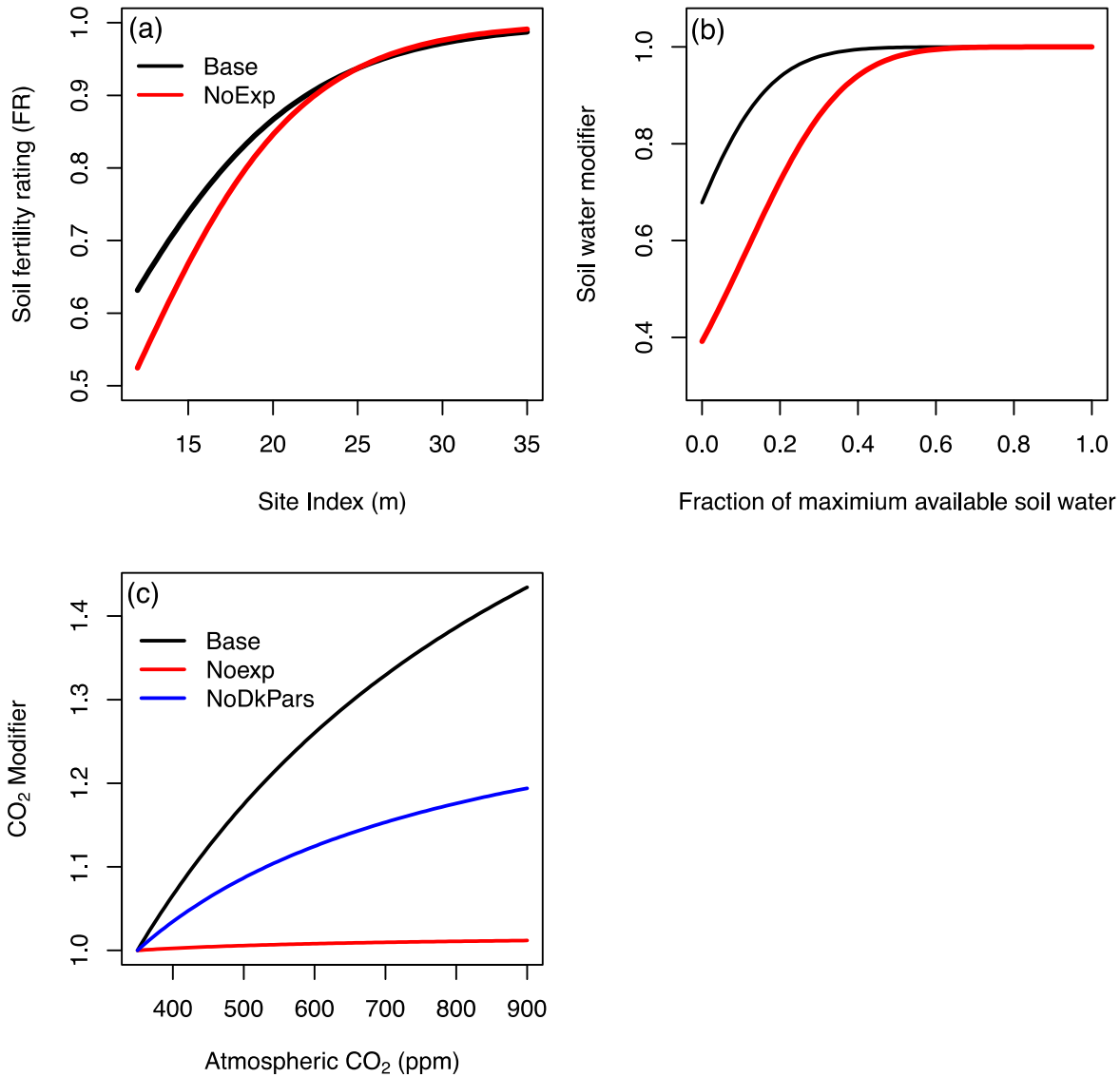
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 982 Figure 4. Model evaluation of stem biomass when assimilating (a) observations across
 983 environmental gradients and ecosystem manipulation experiments (Base; Table 4), and (b)
 984 assimilation only observations across environmental gradients (NoExp; Table 4). The gray
 985 circles correspond to predictions where all plots were used in data assimilation. The black
 986 triangles correspond to predictions where 160 plots were not included in data assimilation and
 987 represent an independent evaluation of model predictions (out-of-bag validation). For each plot,
 988 we used the measurement with the longest interval between initialization and measurement for
 989 evaluation.

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 992 Figure 5. The mean response, expressed as a percentage change in stem biomass from the
 993 control treatment, for irrigation, drought (as a reduction in throughfall), nutrient addition, and
 994 elevated CO₂ experiments. The observed response and the response simulated by the Base,
 995 NoExp, and NoDkPars assimilation approaches are shown. # signifies that value below marker
 996 was significantly different from the observed response ($p < 0.05$). * signifies that value below
 997 marker was significantly different from the response in the Base assimilation ($p < 0.05$). Error
 998 bars are ± 1 standard deviation.
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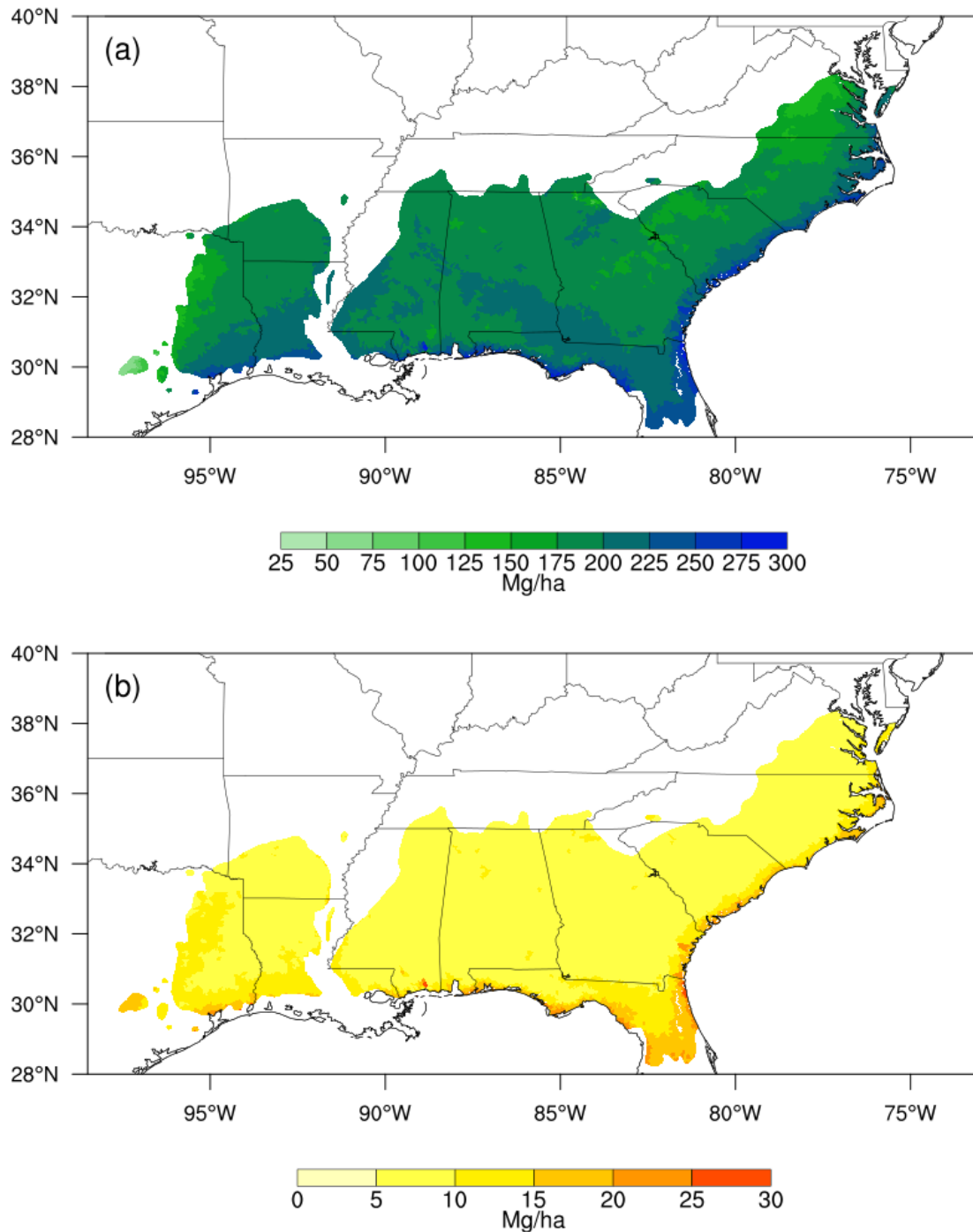
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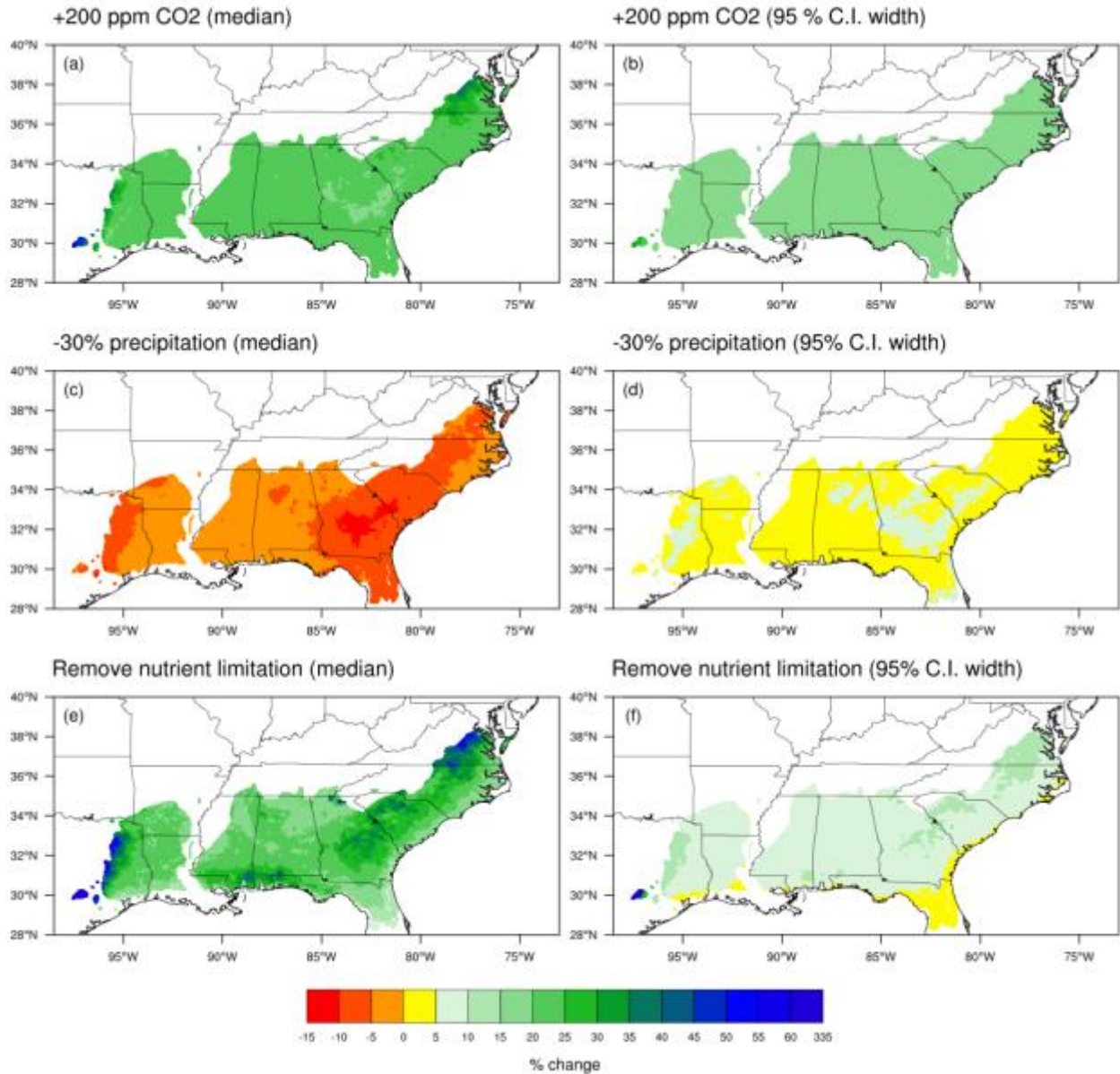
Figure 6. Optimized environmental response functions in the 3-PG model for the (a) soil fertility influence on photosynthesis), (b) available soil water influence on photosynthesis and conductance, and (c) atmospheric CO₂ influence on photosynthesis. The function shapes were derived from the parameters in the Base, NoExp, and NoDkPars assimilations (Table 4).

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Figure 7. (a) Regional predictions of stem biomass stocks for a 25-year-old stand planted in 1985. Parameters used in the predictions were from the Base assimilation approach described in Table 5. (b) The width of the 95% quantile interval associated with uncertainty in model parameters.



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Figure 8. Predictions of the percentage change in stem biomass at age 25 in response to (a,b) a 200 ppm increase in atmospheric CO₂ over 1985-2011 concentrations, (c,d) a 30% reduction in precipitation from 1985-2011 levels, and (e,f) a removal of nutrient limitation by setting the soil fertility rating in the model equal to 1. The left column is the median prediction and the right column is the width of the 95% quantile interval associated with parameter uncertainty. The predictions used the Base assimilation.