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

R. Quinn Thomas<sup>1\*</sup>, Evan Brooks<sup>1</sup>, Annika Jersild<sup>1</sup>, Eric Ward<sup>2</sup>, Randolph Wynne<sup>1</sup>, Timothy J. Albaugh<sup>1</sup>, Heather Dinon Aldridge<sup>3</sup>, Harold E. Burkhart<sup>1</sup>, Jean-Christophe Domec<sup>4,5</sup>, Thomas R. Fox<sup>1</sup>, Carlos A. Gonzalez-Benecke<sup>6</sup>, Timothy A. Martin<sup>7</sup>, Asko Noormets<sup>8</sup>, David A. Sampson<sup>9</sup>, 5

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Robert O. Teskey<sup>10</sup> 7

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<sup>1</sup>Department of Forest Resources and Environmental Conservation, Virginia Tech, USA,

<sup>2</sup>Climate Change Science Institute and Environmental Sciences Division, Oak Ridge National 10

Laboratory, USA, 11

<sup>3</sup>State Climate Office of North Carolina, North Carolina State University, USA, 12

<sup>4</sup>Bordeaux Sciences Agro, UMR 1391 INRA-ISPA, 33175 Gradignan Cedex, France, 13

<sup>5</sup>Nicholas School of the Environment, Box 90328, Duke University, Durham, NC 27708, USA, 14

<sup>6</sup>Department of Forest Engineering, Resources and Management, Oregon State University, USA,

<sup>7</sup>School of Forest Resources and Conservation, University of Florida, USA, 16

<sup>8</sup>Department of Forestry and Environmental Resources, North Carolina State University, USA, 17

<sup>9</sup>Decision Center for a Desert City, Arizona State University, USA, 18

<sup>10</sup>Warnell School of Forestry and Natural Resources, University of Georgia, USA, 19

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\*Corresponding author: R. Quinn Thomas (rqthomas@vt.edu)

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Abstract





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25 Predicting how forest carbon cycling will change in response to climate change and management 26 depends on the collective knowledge from measurements across environmental gradients, 27 ecosystem manipulations of global change factors, and mathematical models. Formally 28 integrating these sources of knowledge through data assimilation, or model-data fusion, allows 29 30 the use of past observations to constrain model parameters and estimate prediction uncertainty. However, the influence of different experimental treatments on those predictions depends on the 31 exact methods and techniques used for data assimilation. Here, we introduce a hierarchical 32 33 Bayesian DA approach (Data Assimilation of Pine Plantation Ecosystem Research, DAPPER) that uses observations of carbon stocks, carbon fluxes, water fluxes, and vegetation dynamics 34 from loblolly pine plantation ecosystems across the Southeastern U.S. to constrain parameters in 35 36 a modified version of the 3-PG forest growth model. The observations included major experiments that manipulated atmospheric carbon dioxide (CO<sub>2</sub>) concentration, water, and 37 nutrients, along with non-experimental studies that spanned environmental gradients across an 38 8.6 x 10<sup>5</sup> km<sup>2</sup> region. We optimized regionally representative posterior distributions for the most 39 sensitive model parameters, which dependably predicted data from plots withheld from the data 40 assimilation. The posterior distributions of parameters associated with ecosystem responses to 41 CO<sub>2</sub>, precipitation, and nutrient addition, along with the corresponding regional changes in 42 production associated with nutrient fertilization and drought, depended on how the experimental 43 data were assimilated. In particular, assimilating nutrient addition experiments reduced the 44 predicted sensitivity to nutrient fertilization while assimilated water manipulation experiments 45 increased the sensitivity to drought. Further, it was necessary to assimilate data from the CO2 46 experimental enrichment site before other studies to constrain the parameters associated with the 47

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influence of CO<sub>2</sub> on canopy photosynthesis. The ambient CO<sub>2</sub> plots were numerous and had a large contribution to the cost function compared to the low number of elevated CO<sub>2</sub> plots (289 ambient vs. 5 elevated plots). Overall, we demonstrated how three decades of research in southeastern U.S. planted pine forests can be used to develop data assimilation techniques that use multiple locations, multiple data streams, and multiple ecosystem experiment types to optimize parameters. This approach allows for future predictions to be consistent with a rich history of ecosystem research across a region.

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1 Introduction



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Forest ecosystems absorb and store a large fraction of anthropogenic carbon dioxide (CO<sub>2</sub>) 58 emissions (Le Quere et al., 2015; Pan et al., 2011) and supply wood products to a growing 59 human population (Shvidenko et al., 2005). Therefore, predicting future carbon sequestration and 60 timber supply is critical for adapting forest management practices to future environmental 61 conditions and for using forests to assist with reduction of atmospheric CO<sub>2</sub> concentrations. The 62 key sources of information for developing these predictions are results from global change 63 ecosystem manipulation experiments, observations of forest dynamics across environmental 64 65 gradients, and process-based ecosystem models. The challenge is integrating these three tools into a common framework for creating probabilistic predictions, or forecasts (based on (Luo et 66 al., 2011a)), that provide information on both the expected future state of the forest and the 67 68 probability distribution of those future states. 69 Data assimilation (DA), or data-model fusion, is an increasingly used framework for integrating 70 71 ecosystem observations into ecosystem models (Luo et al., 2011a; Niu et al., 2014; Williams et al., 2005). DA integrates observations with ecosystem models through statistical, often Bayesian, 72 methods that generate probability distributions for ecosystem model parameters and initial states. 73 DA allows for the explicit accounting of observational uncertainty (Keenan et al., 2011), the 74 incorporation of multiple types of observations with different time scales of collection 75

(Richardson et al., 2010), and the representation of prior knowledge through informed parameter

prior distributions or specific relationships among parameters (Bloom and Williams, 2015).

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

leverage environmental gradients and from ecosystem manipulation experiments will allow for

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80 forecasts to be consistent with the rich history of global change research in forest ecosystems. 81 Ecosystem manipulation experiments provide a controlled environment in which data collected 82 can be used to describe how forests acclimate and operate under altered environmental 83 conditions (Medlyn et al., 2015). These data may be used to constrain model parameters that are 84 85 associated with specific physiological functions associated with, for example, carbon allocation and turnover as related to the controlled manipulation. Furthermore, the assimilation of 86 experiments may increase parameter identifiability (reducing equifinality (Luo et al., 2009)), 87 88 where two parameters have compensating controls on the same processes, by isolating the response to a manipulated driver. For example, carbon assimilation and primary productivity can 89 be modeled as a light and temperature controlled process that is adjusted by nutrients, water, and 90 91 atmospheric CO<sub>2</sub> concentration. In this case, the productivity may mathematically be equal between a parameterization that has high potential conversion of light to photosynthesis (high 92 quantum yield) but low relative nutrient availability and a parameterization with low quantum 93 94 yield but high relative nutrient availability. Therefore, the challenge is that the same rate of production can emerge from different contributions of environmental controls. 95 96 For future predictions with changing environmental conditions, the relative contribution of each 97 environmental control should be separated in order to correctly parameterize the sensitivity to 98 changes in the environment. Key examples of existing and past ecosystem experiments that have 99 the potential to isolate specific parameters in DA include CO<sub>2</sub> enrichment, water manipulation, 100 nutrient addition, and elevated soil temperature experiments. Many of these experiments are 101 common, particularly when including nutrient addition experiments in managed forests. Other 102

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types of experiments are less common, but the few sites with the experiments, such as wholeecosystem CO<sub>2</sub> enrichment, include intensive measurements of numerous carbon pools and fluxes required for model optimization.

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Developing optimized parameters that apply to a region requires assimilating observations that span environmental gradients to support the application of model predictions to a range of climatic conditions, nutrient availabilities, and soil water dynamics. Therefore, the DA of multiple research sites across a region is an important extension of prior DA research focused on DA at a single site with multiple types of observations (Keenan et al., 2012; Richardson et al., 2010; Weng and Luo, 2011). Incorporating multiple locations that include global change experiments in DA is associated with numerous challenges. First, prior research has demonstrated that high frequency observations (i.e., daily, or more frequent, net ecosystem exchange observations) can overwhelm the contribution of low frequency observations (i.e., annual tree diameter measurements) to the cost-function used for optimization (Richardson et al., 2010), resulting in a parameter set that predominately represents the high-frequency dynamics. DA of ecosystem experiments and regional observations can present similar issues because key contrasts isolated in an ecosystem experiment with relatively few plots may be overwhelmed by the contribution of more numerous regional observations from non-manipulated plots. For example, whole ecosystem CO<sub>2</sub> enrichment experiments are uncommon but are the only observations representing ecosystem dynamics in an environment with over 550 ppm atmospheric CO<sub>2</sub> (McCarthy et al., 2010). Therefore, DA techniques may be required that assign additional weight to unique, but rare, experiments in the DA approach. As an example, a multi-

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stage Bayesian approach could be used where the observations from the unique experiment are assimilated first and the posteriors from that assimilation are used as priors for the assimilation of the remaining observations. Second, DA requires using highly simplified ecosystem models because many DA methods use millions of iterations to explore parameter distributions and these iterations have to be applied to both control and manipulated treatments. However, in tension with the need for simple models in DA, more complex models that simulate carbon, water, and nutrient dynamics are also needed to fully leverage the diversity of ecosystem manipulation experiments. Monthly time-scale models of ecosystem processes may be well suited to overcome these challenges for application to predicting changes in biomass over decades in response to global change. First, the contribution of monthly flux and annual biomass measurements to the optimized cost function is more similar in monthly than daily models (12:1 vs. 365:1). Second, they are computationally more efficient than daily models commonly used in DA, allowing data spanning hundreds of plots and multiple decades to be assimilated. Finally, DA is able to calibrate parameters associated with carbon, nitrogen, and water cycles so that they are appropriate for an aggregated monthly time step, helping prevent potential issues associated when applying daily parameterizations to coarser temporal time-steps. Southeastern U.S. planted pine forests are ideal ecosystems for exploring the application of DA to carbon cycle and forest production predictions. These ecosystems are dominated by loblolly pine (*Pinus taeda* L.), thus allowing for a single parameter set to be applicable to a large region containing many soil types and climatic gradients. Loblolly pine represents more than one half of the standing pine volume in the southern United States (11.7 million ha) and is by far the single most commercially important forest tree species for the region, with more than 1 billion

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addition (Albaugh et al., 2016; Carlson et al., 2014; Raymond et al., 2016), water exclusion (Bartkowiak et al., 2015; Tang et al., 2004; Ward et al., 2015; Will et al., 2015), and water addition experiments (Albaugh et al., 2004; Allen et al., 2005; Samuelson et al., 2008). The region also includes a long-term ecosystem CO<sub>2</sub> enrichment study (McCarthy et al., 2010). Furthermore, many of these experiments are multi-factor with water exclusion-by-nutrients (Will et al., 2015), water addition-by-nutrients (Albaugh et al., 2004; Allen et al., 2005; Samuelson et al., 2008), and CO<sub>2</sub>-by-nutrients treatments (McCarthy et al., 2010; Oren et al., 2001). Beyond experimental treatments, Southeastern U.S. loblolly pine ecosystems include at least two eddycovariance sites with high frequency measurements of carbon and water fluxes along with biometric observations over many years (Noormets et al., 2010; Novick et al., 2015), and sites with multi-year sap flow data (Ewers et al., 2001; Gonzalez-Benecke and Martin, 2010; Phillips and Oren, 2001). Finally, there are available studies that include plots that span the regional environmental gradients and extend back to the 1980s (Burkhart et al., 1985). Overall, the high availability of observations of biomass stocks, leaf area index (LAI), carbon fluxes, water fluxes, and vegetation dynamics that span the past 35 years in loblolly pine ecosystems, including plots with experimental manipulation and plots across environmental gradients, is well suited to potentially constrain model parameters and predictions of how carbon cycling responds to environmental change. Our objective was to develop a DA approach that integrated diverse data from multiple locations, including ecosystem experiments, for predicting how forest productivity may respond to global

seedlings planted annually (Fox et al., 2007; McKeand et al., 2003). There is also a rich history

of experimental research focused on global change factors including region-wide nutrient

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change. We applied DA techniques to optimize a monthly-time step, simple forest productivity model using southeastern U.S.-wide experimental (nutrient addition, CO<sub>2</sub> enrichment, and water manipulations) and non-experimental data from 35 years of loblolly pine plantation research in the region. Our DA approach, DAPPER (Data Assimilation of Pine Plantation Ecosystem Research), is unique in its focus on simultaneously assimilating observations from multiple locations, experimental types, and data streams into a simple ecosystem model that includes carbon, water, and (implicitly) nutrients using a hierarchal Bayesian technique to develop parameter distributions. We used the DAPPER system to evaluate the sensitivity of biomass predictions and parameter distributions to the inclusion of ecosystem experiments in DA and to predict the regional sensitivity of forest production to nutrient fertilization and drought.

## 2 Methods

# 185 2.1 Ecosystem Model

We used a modified version of the Physiological Principles Predicting Growth (3-PG) Model to simulate vegetation dynamics in loblolly pine stands (Bryars et al., 2013; Gonzalez-Benecke et al., 2016; Landsberg and Waring, 1997). 3-PG is a stand-level vegetation model that runs at the monthly time-step and includes vegetation carbon dynamics and a simple soil water bucket model (Figure 1). While a complete description of the 3-PG model and our modifications can be found in the Supplemental Material, the key concept for interpreting the results is that gross primary productivity (GPP) was simulated using a light-use efficiency approach where the absorbed photosynthetically active radiation (APAR) was converted to carbon based on a quantum yield. Quantum yield was simulated using a parameterized maximum quantum yield

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 $(\alpha)$  that was modified by environmental conditions including air temperature, atmospheric CO<sub>2</sub>, 195 available soil water and soil fertility. The available soil water and soil fertility modifiers were 196 values between 0 and 1, while the atmospheric CO<sub>2</sub> modifier had a value of 1 at 350 ppm and 197 values greater than 1 at higher CO<sub>2</sub> concentrations. 198 199 Elevated CO<sub>2</sub> modified tree physiology by increasing quantum yield, based on an increasing but 200 saturating relationship with atmospheric CO<sub>2</sub>. We also added a function where the allocation to 201 foliage relative to stem biomass decreased as atmospheric CO<sub>2</sub> increased. Available soil water 202 and quantum yield were positively related through a logistic relationship between relative 203 available soil water and the quantum yield modifier, where relative available soil water was the 204 ratio of simulated available soil water to a plot-level maximum available soil water. Soil fertility 205 and quantum yield were proportionally related, where quantum yield was scaled by an estimate 206 of relative stand-level fertility where a value of 1 was the maximum fertility. The fertility 207 modifier (FR) was constant throughout a simulation of a plot and was either based on site 208 characteristics or directly optimized as a stand-level parameter. Here we used site-index, a 209 measure of the height of a stand at a specified age (25 years), and the 35-year mean annual 210 temperature as site characteristics to predict FR. For a given climate, site index captures 211 differences in soil fertility, where a lower site index corresponded to a site with lower fertility. 212 However, regional variation in site index also included the influence of climate on growth rates 213 that were already accounted for in the other environmental modifiers in the 3-PG model. To 214 account for the climatic influence on site index, a long-term climate variable (35-year mean 215 annual temperature) was included in the empirical relationship that predicted FR as an 216 increasing, but saturating, function of site index. For plots with nutrient fertilization, FR was a 217

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218 directly optimized parameter. For our application of the 3-PG model using DA, we removed the previously simulated dependence of total root allocation on FR(Bryars et al., 2013; Gonzalez-219 Benecke et al., 2016). Therefore, plots with lower FR could be interpreted to have lower 220 221 quantum yield. Other environmental conditions influenced GPP, including temperature, frosts days, and vapor pressure deficit with a description of these modifiers found in the Supplemental 222 Material. 223 224 Each month, net primary production (a parameterized and constant proportion of GPP) was 225 226 allocated to foliage, stem (stemwood, stembark, and branches), coarse roots, and fine roots. Differing from previous applications of 3-PG to loblolly pine ecosystems, we modified the 227 model to simulate fine roots and coarse roots separately. 3-PG also simulated simple population 228 229 dynamics by including stem density as a state variable. Stem density and stem biomass pools were reduced by both density-dependent and density-independent mortality (a new 230 modification), with the former based on the concept of self-thinning. Finally, we added a simple 231 232 model of hardwood understory vegetation to enable the use of estimates of gross primary productivity and evapotranspiration from eddy-covariance tower studies with significant 233 understories. Details of the model can be found in the Supplemental Material. 234 235 The water cycle was a simple bucket model with transpiration predicted using a Penman-236 Monteith approach (Bryars et al., 2013; Gonzalez-Benecke et al., 2016; Landsberg and Waring, 237 1997). The canopy conductance used in the Penman-Monteith subroutine was modified by 238 environmental conditions. The modifiers include the same available soil water and vapor 239 pressure deficit modifier as used in the GPP calculation. Maximum canopy conductance 240

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241 occurred when simulated LAI exceeded a parameterized value of leaf area index (LAI). Evaporation was equal to the precipitation intercepted by the canopy. Runoff occurred when the 242 available soil water exceeded a plot-specific maximum available soil water. As in prior 243 applications of 3-PG, available soil water was not allowed take a value below a minimum 244 available soil water, resulting in an implicit irrigation in very dry conditions. 245 246 The 3-PG model used in this study simulated the monthly change in eleven state variables per 247 plot: four stocks for loblolly pines, five stocks for understory hardwoods, loblolly pine stem 248 density (stems ha<sup>-1</sup>), and available soil water. The key fluxes that were used for DA included 249 monthly GPP, monthly evapotranspiration (ET), annual root turnover, and annual foliage 250 turnover. In total, 46 parameters were required by 3-PG with 31 of the parameters optimized 251 252 using DA (Table 1, Supplemental Table 1, SI Table 2). The model required mean daily maximum temperature, mean daily minimum temperature, daily PAR, total frost days, total rain 253 at the monthly time scale, monthly atmospheric CO<sub>2</sub>, and latitude. Each plot also required 254 255 maximum available soil water, site index, mean annual temperature, and the initial condition of the eleven state variables as model inputs (Figure 2). 256 257 258 2.2 Observations We used thirteen different data streams from 294 plots at 187 unique locations spread across the 259 region to constrain model parameters (Table 2; Figure 3). The data streams covered the period 260 261 between 1981 to 2015. All data streams were not available in all plots (Table 2; Table 3). The most common set of data streams were annual or less frequent observations of stand stem 262 biomass (defined as the sum of stemwood, stembark and branches), winter foliage biomass, and 263 living tree counts. The stem and foliage biomass were optimized using regional allometric 264

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models based on measurements of tree diameter, height, and plot level-stem size distributions (Gonzalez-Benecke et al., 2014). The most comprehensive set of data streams was from Duke Forest where annual measurements and allometric-based estimates were made of stem biomass (loblolly pine and hardwood), coarse root biomass (loblolly pine and hardwood), fine root biomass (combined loblolly pine and hardwood), stem count (loblolly pine only), leaf turnover (combined loblolly pine and hardwood), and fine root production (combined loblolly pine and hardwood). The Duke Forest dataset (DK3 combined with the Duke FACE CO<sub>2</sub> fertilization study) also included monthly observations of LAI, gross ecosystem production (GEP; modeled gross primary productivity from net ecosystem exchange measured at an eddy-covariance tower), and ET. The set of data streams associated with a particular site and experimental design is shown in Table 3. The measurement uncertainty associated with each data stream is listed in Table 2. Since the model used a monthly time-step, and plots with only biomass and stem density observations were more common than plots with monthly flux estimates, the data used in the optimization cost function were not dominated by high frequency data streams (GEP and ET).

## 2.3 Data assimilation method

We used a hierarchal Bayesian framework to approximate the posterior probability distributions of model parameters in Table 1, the model process uncertainty parameters, and the latent model states and fluxes. The latent model states represented the 'true' stock or flux before measurement uncertainty was included in the observation. Our hierarchal approach was designed to partition uncertainty that is attributable to uncertainty in parameters, model process, and measurements (Hobbs and Hooten, 2015). Previous forest ecosystem DA efforts have either focused on parameter uncertainty, by using measurement uncertainty as the variance term in a Gaussian cost

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function, or on total uncertainty by directly estimating the Gaussian variance term. The latter 288 combines measurement uncertainty and process uncertainty into the same parameter and is 289 unable to be used for developing prediction intervals, as prediction intervals only include 290 291 parameter and process errors (Dietze et al., 2013; Hobbs and Hooten, 2015). Here, our focus was on estimating the probability distribution of forest biomass before uncertainty is added through 292 measurement. 293 294 First, we estimated the probability of a latent state or flux  $(z_{i,m,p})$  for each data point (i) from each 295 296 data stream (m) in a plot (p) using the 3-PG model with the plot FR. This included the optimized parameters  $(\theta_F)$ , fixed parameters  $(\theta_C)$ , soil characteristic inputs (S), climate inputs (C), site 297 index (SI), fertility (FR<sub>p</sub>), and initial conditions (I) required by the 3-PG to simulate each plot, 298  $f(\theta_F, \theta_c, C, S, I, FR_p)$ . The latent state  $(z_{i,m,p})$  was assumed to be normally distributed with the mean 299 from the 3-PG simulation and an optimized, data stream-specific, process variance  $\sigma_{m,(process)}^2$ 300 301 p(process process parameters)= 302  $P(z_{i,m,p}|f(\theta_F,\theta_C,C,S,I,FR_p),\sigma_{m (process)}^2)$ 303  $\sim$ Normal $(z_{i,m,p}|f(\theta_F,\theta_C,C,S,I,FR_p),\sigma_{m (process)}^2)$ Equation 1 304 305 306 The unobserved true state related to the observed state through a data observation model. In the sampling model, the measured state  $(y_{i,m,p})$  was a random sample from a normal distribution 307 with a mean of the true state and a data point-specific standard deviation ( $\sigma_{i,m,p}^2$ ). 308 309 p(data|process,data parameters)= 310

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311	$P\left(\boldsymbol{y}_{i,m,p} \middle  \boldsymbol{z}_{i,m,p}, \sigma_{i,m,p}^2\right) \sim Normal\left(\boldsymbol{y}_{i,m,p} \middle  \boldsymbol{z}_{i,m,p}, \sigma_{i,m,p}^2\right)$	Equation 2
312		
313	This standard deviation $(\sigma^2_{i,m,p})$ represented measurement uncertainty and was similar to the	
314	denominator in least-squares approach that is commonly used in DA (Bloom and Williams,	
315	2015; Keenan et al., 2011).	
316		
317	Each parameter $(\theta_F)$ that was optimized using the Bayesian method had a prior probability that is	
318	specified in Table 1. The prior distribution for the standard deviation $\sigma^2_{m,(process)}$ parameters	
319	were uniformly distributed:	
320		
321	p(process parameters priors)×p(priors)= $P(\sigma_m^2)$ × $P(\theta_F)$	Equation 3
322		
323	where	
324		
325	$P(\sigma_m^2) \sim unif(0.001,100)$	Equation 4
326		
327	and	
328		
329	$P(\theta_F)$ ~See Supplemental Table 1	Equation 5
330		
331	Finally, following the description of the plot specific FR <sub>p</sub> described above, the probability for	
332	fertilized treatments was based on a comparison to the control treatment F	R.
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$$P(FR_p|\theta_F,E) = \begin{cases} 1 \text{ if non-fertilized} \\ 1 \text{ if fertilized and } FR_p \ge FR \text{ of control plot} \\ 0 \text{ if fertilized and } FR_p < FR \text{ of control plot} \end{cases}$$
 Equation 6

Our complete Bayesian model for estimating the posterior distributions for the parameters ( $\theta_{\rm F}$ ),

process uncertainty ( $\sigma_{m,(process)}^2$ ), and unobserved true states ( $z_{i,m,p}$ ) was:

$$339 \qquad P\left(\theta_{F},\sigma_{m}^{2},z_{i,m,p}\left|y_{i,m,p},\sigma_{i,m,p}^{2},\theta_{C},S,C,SI,I\right.\right) \varpropto$$

340 
$$P(z_{i,m,p}|f(\theta_F,\theta_C,FR_p,E),\sigma_m^2)P(y_{i,m,p}|z_{i,m,p},\sigma_{i,m,p}^2)P(FR_p|\theta_F,E)P(\theta_F)P(\sigma_m^2)$$
 Equation 7

We numerically estimated the posterior distributions using the Monte-Carlo Markov Chain –

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

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

345 Trudinger et al., 2007; Williams et al., 2005; Zobitz et al., 2011). We adapted the size of the

jump for each parameter (i.e., how far a proposed new value can potentially be from the current

value) to ensure the acceptance rate of the parameter set is between 22% and 43% (Ziehn et al.,

348 2012). All MCMC-MH chains were run for 30 million iterations with the first 15 million

iterations discarded as the burn-in. Three chains were run and compared for convergence and we

sampled every 1000<sup>th</sup> parameter in the final 15 million iterations of the MCMC-MH chain. This

thinned chain was used in the analysis described below. The 3-PG model and MCMC-MH

algorithm were programed in FORTRAN 90 and used OpenMP to parallelize the simulation of

each plot within an iteration of the MCMC-MH algorithm.

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#### 2.4 Model simulations

Each plot simulated required initial conditions for each model state, climate inputs, soil characteristic inputs, and site index. We used the first observation at the plot as the initial conditions for the loblolly pine vegetation states (foliage biomass, stem biomass, coarse root biomass, fine root biomass, and stem number). When observations of coarse biomass and fine root biomass were not available, these stocks were initialized as a mean region-wide proportion of the observed stem biomass. However, the value of initial root biomass in plots without observations was not important because the plots without root observations did not contribute to the root cost function and root biomass does not influence any other functions in the model. In the two plots with flux observations (US-Dk3 and US-NC2), hardwood understory was also initialized using the first set of observations. Initial fine root and coarse biomass was distributed between loblolly pine and hardwoods based on their relative contribution of total initial foliage biomass. The initialized available soil water was assumed to be equal to the maximum available soil water because most plots were initialized in winter months when plant demand for water is minimal. The maximum available soil water in each plot was extracted from the SSURGO soils dataset (Staff, 2016). We assumed that the minimum available soil water was zero. The value we used corresponded to the maximum available soil water for the top 1.5 m of the soil. Because we focused on a region-wide optimization, we used region-wide 4-km estimates of observed monthly meteorology as inputs and to collect the 35-year mean annual temperature for each plot (Abatzoglou, 2013). Site index was based on height measurements at age 25 in each plot or calculated by combining observations of height at younger ages with an empirical model

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(Dieguez-Aranda et al., 2006).

We simulated the experiments by altering the environmental modifiers or by modifying the environmental inputs. Nutrient addition experiments were simulated by directly estimating FR, rather than calculating from Equation 2, and by requiring the optimized FR in the fertilized plot to be equal to or greater than the FR in the control plots. Throughfall exclusion experiments were simulated by decreasing rain inputs by 30% in the treatment plots. This assumed that the fractional reduction in precipitation and throughfall were equal. The SETRES Irrigation experiments were simulated by adding 650 mm to precipitation between April and October. CO<sub>2</sub> enrichment experiments were simulated by setting the atmospheric CO<sub>2</sub> input equal to the treatment mean from the elevated CO<sub>2</sub> rings (570 ppm). While not an experiment, one plot (US-NC2) included a thinning treatment during the period of observation. We simulated the thinning by specifying a decrease in the stem count that matched the proportion removed at the site, with the biomass of each tree equivalent to the average of trees in the plot.

## 2.4 Model experiments and analysis

Our analysis focused on comparing parameter distributions and predictions among simulations that used different experimental treatments to estimate the posterior distributions (Table 4). To examine the influence of the Duke FACE CO<sub>2</sub> fertilization, we compared a one stage vs. a two-stage data assimilation process. The one stage process assimilated all observations in all plots and experiments simultaneously. In this approach, the elevated CO<sub>2</sub> plots only represented 5 of the 294 plots across the region and thus a relatively minor contribution to the likelihood (cost-function) calculation. The two-stage process used the observations from Duke FACE, US-Dk3

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flux site, the other flux site in North Carolina (US-NC2) to estimate parameter posteriors using the priors in Table 1 and SI Table 1. These sites were grouped together because they were the most data rich, had the high frequency data streams (monthly GEP, ET, and LAI), and were relatively close in geography. FR was directly estimated for all plots in the first stage, with the FR of a fertilized plot required to be equal to or higher than its control plot. The FR of the CO<sub>2</sub> experiment was equal to the corresponding control plot estimated FR. The FR of the control plot was required to be greater than 0 and, if associated with a nutrient fertilization plot, less than the FR of the fertilized plot. For the second DA stage, the posterior distributions from the first stage were used as priors for the assimilation of the region-wide observations from the PINEMAP, FPC RW 18, FMRC Thinning, SETRES, and Waycross studies (Table 4). We compared the CO<sub>2</sub> quantum yield enhancement parameter (Calpha 700) between the one and two stage approaches to evaluate how the estimation of CO<sub>2</sub> fertilization of plant growth depended on how the Duke FACE data are used in data assimilation. We also estimated the distribution of the percentage increase in net primary productivity (NPP) associated with the elevated CO<sub>2</sub> treatment using the one and two stage data assimilation approaches. The distribution of the percentage increase in NPP was calculated by randomly selecting 1000 parameter sets, with replacement, from the 1-stage converged MCMC chains. This calculation was repeated using the 2-stage approach. Based on the results from comparing the one and two stage approaches (see results below), we proceeded using the two-stage approach to examine the influence of the water manipulation and nutrient fertilization experiments on posterior distributions and predictions. To evaluate the

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influence of water manipulation experiments, we repeated the second stage of the data assimilation without the plots where water was added or subtracted. To evaluate the influence of the nutrient manipulation experiments, we first repeated the first stage of data assimilation without the nutrient addition plots in the Duke FACE experiment and used those posteriors as priors to the second stage. This ensured that the priors to the second stage of data assimilation did not include information from nutrient addition experiments. The second stage then excluded the other nutrient manipulation experiments in the region. To examine how the exclusion of the water manipulation experiments influenced parameter inference and predictions, we first examined how the parameter distributions changed from initial priors through the two assimilation stages. With respect to the water manipulation experiments, we focused on the shape of the relationship between available soil water and the quantum yield and stomatal conductance modifier (governed by parameters SW1 and SW2) with and without assimilating the water manipulation experiments. To illustrate the capacity to estimate the probability distribution of predictions using the posterior uncertainty in parameters, we analyzed a focal site in Georgia, near the center of the loblolly pine range (circle in Figure 2). At the focal site, we predicted the sensitivity of stem biomass at age 25 (hereby referred to as STEM<sub>25</sub>) to a 30% increase and a 30% decrease in annual precipitation with and without assimilating the water experiments. A 30% percent decrease in precipitation mirrors the magnitude of reduction in the experimental throughfall reduction studies used in DA (Table 3 and Figure 3). Our prediction distributions were calculated by integrating across the parameter uncertainty by repeating simulations using 1000 random draws from the converged chain of the posteriors. Finally, we predicted the regional response to a reduction in precipitation from

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historical using the median posterior parameter values from the data assimilation with and without the water experiments included. Our regional corresponded to the native range of loblolly and used the HUC12 (USGS 12-digit Hydrological Unit Code) watershed as the scale of simulation. For each HUC12 in the region we used the mean site index, 30-year mean annual temperature, available soil water aggregated to the HUC12 level, and monthly meteorology as inputs (Figure 2). We simulated forest development from 1989 to 2014 using actual precipitation and again with a 30% reduction in precipitation. We focused our analysis on the percent change in STEM<sub>25</sub> between the two simulations. To examine how the exclusion of the nutrient addition experiment influenced parameter inference and prediction, we focused on the difference in maximum quantum yield parameter ( $\alpha$ ) and the relationship between site index and soil fertility modifier (FR) with and without assimilating the nutrient experiments. Additionally, we simulated how stem biomass at age 25  $(STEM_{25})$  responded to a complete removal of nutrient limitation (FR = 1) for the focal site in Georgia. As in the precipitation sensitivity described above, we represented the percentage change in STEM<sub>25</sub> between simulations with estimated FR and FR = 1 as a distribution by integrating across parameter uncertainty. We predicted the regional response to nutrient fertilization by setting the FR at all HUC12 units (see previous paragraph) equal to 1 using the median posterior parameter values from data assimilation where nutrient addition experiments were either included or not. We focused on the regional pattern in the percentage change in stem biomass with the predicted FR (current level fertility) and FR = 1 (nutrient limitation removed). Finally, we assessed overall model performance of the 2-stage approach for data assimilation

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with all experimental types included in DA, excluding the nutrient addition experiments, and excluding the nutrient addition experiments using an out-of-sample approach. The approach held 40 random FMRC thinning study plots (Table 3) out from the assimilation, predicted the 40 plots using the median parameter values, and compared the predicted stem biomass to the observed stem biomass. These were plots without any manipulations of nutrients or water, were located throughout the region, and had measurement ages up to 30 years old. For each plot, we only used the most recent observed values to increase the time length between initialization and validation. We repeated the validation for four unique sets of 40 FMRC thinning study plots.

#### 3 Results

Our multi-site, multi-experiment, multi-data stream DA approach was able to constrain most parameters in the 3-PG model (31 of 46 parameters were optimized; Table 6; Supplemental Table 3; Supplemental Figure 1-3). The 31 optimized parameters were the most sensitive parameters in the 3-PG model, defined by the change in total biomass at age 25 for the focal site in Georgia to a 10% change in the parameter (Table 1; Supplemental Table 1). One exception was the light extinction coefficient (k), which showed high sensitivity but was assumed to be fixed because it strongly co-varied with the quantum yield parameter ( $\alpha$ ). Parameters associated with biomass allocation had priors with large variance but DA was able to provide posteriors with relatively low variance (pFS2, pFS20, pR, and pCRS; Supplemental Figure 1; Supplemental Table 3). The DA process also produced posterior distributions that had less variability than the prior distribution for the important parameters associated with light-use efficiency ( $\alpha$ , y, FR1, and FR2; Table 5). DA did not change the parameter distributions, i.e., the posterior and prior distributions were similar, for the parameters that governed the temperature sensitivity of

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491 quantum yield, the VPD sensitivity of quantum yield, and the maximum canopy conductance (Supplemental Figure 1-2; Supplemental Table 3). These parameters had strong priors supported 492 by previous research on loblolly pine physiology. Finally, the DA approach was able to estimate 493 the distributions of the process uncertainty parameters (Supplemental Figure 3; Supplement 494 Table 4). 495 496 The addition of the second stage of assimilation that used region-wide observations and 497 posteriors from the DK+NC2 assimilation modified the distributions of the parameters that 498 499 related to allocation and mortality but did not provide additional constraint on the physiological parameters (Table 5). In particular, the parameters associated with the self-thinning curve and 500 allocation of coarse roots had non-overlapping 95% credible intervals between the DK+NC2 and 501 502 RW assimilation. The larger estimate for Wsx1000 and lower value for thinPower in the DK+NC2 indicated self-thinning was lower at the sites in the DK+NC2 assimilation than the 503 average of the other sites in the region. The lower value for the pCRS parameter indicated that 504 505 less NPP was allocated to coarse roots in the DK+NC2 assimilation than the RW assimilation. 506 507 The two-stage assimilation was critical for constraining the CO<sub>2</sub> quantum yield enhancement 508 parameter (Calpha700). Both the mean of the posterior distribution and the range of the 95% credible interval were smaller for fCalpha700 when all observations were assimilated 509 simultaneously (1-stage approach) than the distribution estimated using the 2-stage approach 510 (Duke and NC2 assimilated before the region-wide assimilation) (Figure 5a; Table 5). Despite 511 the same data used in both approaches, the differences in fCalpha700 led to a predicted lower 512 enhancement of NPP associated with elevated CO<sub>2</sub> in the experiment. The 1-stage assimilation 513

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514 approach had a median increase in NPP between the control and elevated CO<sub>2</sub> treatments of 15% compared to a 27% in the two-stage approach (Figure 5b). 515 516 The RW assimilation constrained the soil fertility parameters that were necessary to enable 517 regional simulations. Our regional model using the 2-stage approach performed well compared 518 519 to stem biomass data not used in the assimilation. The mean bias in stem biomass of the four outof-sample validation sets was -6.7 % and the RMSE was 21.2 Mg ha<sup>-1</sup> (Figure 4). 520 521 522 Excluding the nutrient addition experiments from the DA increased the simulated level of nutrient limitation but did not change the predictive capacity of the independent non-manipulated 523 validation set. DA without nutrient fertilization experiments had a greater and more uncertain 524 525 value for the maximum quantum yield parameter ( $\alpha$ ; Figure 6a; Table 5). This parameter was shared across all plots and modified by the environmental conditions at each plot. To compensate 526 for the higher  $\alpha$  parameter when nutrient fertilization experiments were excluded from DA, the 527 two soil fertility parameters (FR1 and FR2) combined to predict a 10% lower FR values for a 528 given site index and mean annual temperature (Figure 6b). Subsequently, the prediction for the 529 percentage change in STEM<sub>25</sub> associated with maximum fertilization (i.e., setting FR = 1) at the 530 focal site in Georgia was 7% higher and had greater uncertainty when nutrient fertilization 531 experiments were excluded from the DA (Figure 6c). The RMSE and mean bias of the non-532 manipulated validation set was 20.4 Mg ha<sup>-1</sup> and -4.8 %, respectively (SI Figure 1a) 533 534 Excluding the water manipulation experiments from the DA reduced the sensitivity to available 535 soil water but, similar to the inclusion of the nutrient addition experiments, did not change the 536

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predictive capacity of the independent non-manipulated validation set. The combined differences in the SW1 and SW2 parameters between the DA with and without the water manipulation experiments decreased the sensitivity of quantum yield and canopy conductance to a reduction in available soil water (Figure 7a). For example, at an available soil water to maximum available soil water ratio of 0.50, the quantum yield and canopy conductance modifier decreased from 0.95 without water experiments to 0.8 with water experiments (Figure 7a). At the focal site in Georgia, the sensitivity of STEM<sub>25</sub> to a reduction in annual precipitation (Figure 7b) was larger when the water experiments were included in the DA (-8.5% median change in STEM<sub>25</sub> for a 30% reduction in precipitation) than when the experiments were excluded (-4.1% median change in STEM<sub>25</sub> for a 30% reduction in precipitation). Similarly, the predictions of STEM<sub>25</sub> change associated with a 30% increase in precipitation (median: 3.8%) were higher when water experiments were included than when not included (median: 1.1%). The magnitude of uncertainty in the predictions did not differ substantially between forecasts with and without water experiments (Figure 7b). The RMSE and mean bias of the non-manipulated validation set was 19.3 Mg ha<sup>-1</sup> and -5.8 %, respectively (SI Figure 1b) Regionally (i.e., the native range of loblolly pines), using the two-stage approach (RW), the most productive areas were the coastal plains and the interior of Mississippi and Alabama (Figure 8). These patterns were largely driven by patterns in the soil fertility factor (FR; Figure 9), reflecting the sensitivity of the 3-PG model to the FR parameters (Table 1). The area weighted mean STEM<sub>25</sub> response to fertilization (represented by setting FR = 1) across the region was 28% with the highest response occurring in the far west of the region, the Piedmont of Georgia, the interior of the gulf coast, and the northern reach of the region (Figure 10a). These were all areas with the

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560 lowest soil fertility parameter. The least responsive region to nutrient addition was in Florida (Figure 10a). Excluding the nutrient addition experiments from the DA increased the sensitivity 561 to nutrient addition (Figure 10b), as shown for the focal Georgia site (Figure 6b), but did not 562 563 change the spatial patterns of the response. 564 The sensitivity of forest production to a 30% reduction in precipitation varied across the region. 565 The most sensitive areas, the Piedmont of Georgia and the western edge of the region, predicted 566 up to a 13.1% decline in STEM<sub>25</sub> (Figure 11a). These were warm areas with relatively low 567 568 precipitation before the 30% reduction (Figure 2c). The least sensitive area was the interior of the gulf coast (<1% decline; Figure 11a), the area with the highest precipitation in the region (Figure 569 2c). The regional mean reduction in STEM<sub>25</sub> associated with a 30% decrease in precipitation was 570 571 5.7% (Figure 11a). Excluding the water manipulation experiments from DA reduced the regional mean sensitivity to 1.7% (Figure 11b). 572 573 574 **Discussion** 575 576 Using DA to parameterize models applied to forecasting ecosystem change requires detangling the vegetation responses to temperature, precipitation, nutrients, and elevated CO<sub>2</sub>. To address 577 this challenge, we introduced a regional-scale hierarchical Bayesian approach (DAPPER) that 578 assimilated data across environmental gradients and ecosystem manipulation experiments into a 579 modified version of the 3-PG model to estimate parameters and generate uncertainty estimates 580 on predictions of carbon and water cycling across the whole native range of loblolly pine. 581 Furthermore, we organized observations of carbon stocks, carbon fluxes, water fluxes, vegetation 582

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structure, and vegetation dynamics that spanned 35 years of forest research (Figure 3; Table 3) in a region with large and dynamic carbon fluxes (Lu et al., 2015). By combining the DAPPER system with the regional set of observations, we were able to estimate parameters in a model with high predictive capacity (Figure 4) and with quantified uncertainty on parameters (Table 5). We also found that the predictions of forest productivity response to rising CO<sub>2</sub>, altered precipitation, and altered nutrient availability were highly sensitive to the types of experiments used in DA as well as the methodological approach applied. We found that including nutrient and water manipulation experiments aided in distinguishing the mechanisms driving patterns in biomass across the region. Including these experiments in the data-assimilation did not improve the predictive capacity of the independent validation set of non-manipulation plots. However, including nutrient and water manipulation did change the underlying mechanisms explaining the patterns in stem biomass. Without the nutrient and water manipulation experiments, the same biomass predictions were attributable to a higher level of nutrient limitation and a lower level of water limitation. This resulted in differing sensitivities to changes in nutrient or water availability. Overall, this finding highlights a key challenge when parameterizing ecosystem models that will be used for global change predictions, that different combinations of environmental drivers can produce similar predictions of current observations. Ecosystem manipulation experiments are an important tool for addressing this challenge. Parameter and process identifiability, or equifinality, presents a challenge when parameterizing ecosystem models using DA (Luo et al., 2009). One important source of equifinality is the tradeoff between parameters governing the potential productivity of the vegetation and the

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downregulation of productivity due to nutrient limitation. When using observational data at a single site, a single parameter is often optimized to set a photosynthetic rate per absorbed light, i.e., a quantum yield. This single parameter combines the potential photosynthesis set by climate and the influence of nutrient limitation on photosynthesis into a single parameter. However, separating these two processes into two or more parameters is challenging because a high potential quantum yield parameter ( $\alpha$ ) and high nutrient limitation (FR) can mathematically yield the same photosynthetic rate as low potential quantum yield and low nutrient limitation. The former implies a larger potential response to nutrient addition than the latter. We found that including nutrient addition experiments in DA helped overcome this challenge. In the case of the 3-PG model used in this study, the maximum quantum parameter ( $\alpha$ ) and soil fertility parameters (FR1 and FR2) were more constrained and inferred lower levels of nutrient limitation across the region when nutrient fertilization experiments were included in the DA. This finding likely extends to other models that include the concept of potential productivity and productivity downregulated by nutrient limitation. For example, the applications of the Data Assimilation Link Ecosystem Carbon (DALEC) model (Williams et al., 2005) to DA often assumed nine of the ten parameters associated with photosynthesis were fixed, thus using a single parameter to represent both the quantum yield (defined as nitrogen use efficiency in DALEC) and the magnitude of nitrogen limitation of a site (Fox et al., 2009). The use of a single parameter, rather than using nutrient addition experiments to separate into multiple parameters, is appropriate when assuming nutrient availability is static. Applications of DA to predictions of ecosystems with changing nutrient availability, either through management, elevated CO<sub>2</sub>, or nitrogen addition, would benefit from using nutrient addition studies to quantify the magnitude of nutrient limitation. Studies of known nutrient gradients could be used in lieu of nutrient addition studies,

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629 but effort must be made to account for confounding abiotic factors, such as available soil water or climatic conditions, that may co-vary with nutrient availability. 630 631 632 Another challenge in DA is deciding how to weigh different types of data used in model fitting (Gao et al., 2011; Wutzler and Carvalhais, 2014). Here we demonstrate that DA efforts should 633 also consider how to weigh different types of ecosystem experiments. In our analysis, we 634 included three types of experiments: nutrient addition, water manipulation, and CO<sub>2</sub> fertilization. 635 The nutrient addition and water manipulation experiments were represented by multiple sites 636 637 across the region while the CO<sub>2</sub> fertilization only occurred at a single location (Figure 3). We found that the parameter that represents the increase in maximum quantum yield under elevated 638 CO<sub>2</sub> was substantially lower when all observations, sites, and experiments were assimilated 639 640 simultaneously than when the CO<sub>2</sub> fertilization experiment was given greater weight. The greater weight was applied by first assimilating the CO<sub>2</sub> fertilization experiment and using the posteriors 641 as priors for assimilating the remaining observations. Providing additional weight on the single 642 643 site with unique environmental conditions (i.e., atmospheric CO<sub>2</sub> at 570 ppm) using a two-stage data-assimilation, we were able to more accurately represent the observed differences in NPP 644 between the ambient and elevated CO<sub>2</sub> treatments at the Duke site (McCarthy et al., 2010). 645 Given than only a few of the parameters were significantly different between the Duke site and 646 the other studies across the region, it may be possible to optimize one parameter for the Duke site 647 and another parameter for the other studies in a 1-stage approach that combines all the plots into 648 a single assimilation. However, the 2-stage approach was required to identify which parameters 649 were different between the Duke site and the other studies. Overall, we suggest that DA efforts 650 using multiple studies and multiple experiment types identify whether particular experiments at 651

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652 limited number of sites have the potential to uniquely constrain specific parameters. In this case, additional weight may be needed to avoid having the signal of the unique experiment 653 overwhelmed by the large amount of data from the other sites and experiments. 654 655 Our analysis highlights that nutrient limitation of productivity was widespread across the region. 656 The largest potential gains in productivity from nutrient addition were predicted in central 657 Georgia, an area with warm annual temperatures but poor soils, as expressed in the low site 658 index. The baseline fertility used in our regional analysis was derived from an empirical model 659 660 of site index that was developed using field plots with minimal management (Sabatia and Burkhart, 2014). Subsequently our estimate of baseline fertility is likely on the low end of forest 661 stands currently in production. Further, we recognize that the site index model had uncertainty 662 663 that could be formally incorporated into the hierarchal Bayesian approach in future applications. 664 The soil fertility modifier has commonly been used to calibrate the 3-PG for applications to a 665 666 single site, with recent work focused on developing an approach to predicting the soil fertility modifier from environmental conditions (Gonzalez-Benecke et al., 2016; Subedi et al., 2015). 667 We have extended prior efforts to develop a simple predictive model of FR in two ways. First, 668 we simultaneously calibrated the parameters in the empirical FR model alongside the other 669 parameters in the 3-PG model. Prior studies have assumed fixed values for the 3-PG model 670 parameters, fitted FR for plots with observations, and developed a relationship between FR and 671 site index. Our Bayesian approach to simultaneously calibrating the 3-PG parameters and the FR 672 model allowed for the estimation of uncertainty and covariation among parameters in the 3-PG 673 and FR models. Second, we included a climate term (mean annual temperature) in the 674

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relationship between site index and FR. This resulted in a lower FR for a given site index in warmer locations. By including the climate term, FR can be interpreted as relative to the climate at a given location and the potential productivity of a plot can be optimized by setting FR equal to 1. When a climate term is not used in the empirical FR model, FR is relative to the greatest site index in the region, which does not occur in the northern extent of the region even in fertilized plots due to climatic constraints. Our simulations show that loblolly pine productivity was not strongly sensitive to changes in precipitation at present day temperatures and atmospheric CO<sub>2</sub>. We simulated a 30% reduction in annual precipitation and found a maximum of a 13.1% reduction in productivity. A 30% reduction in precipitation is plausible but is more extreme than most Multivariate Adaptive Constructed Analogs (MACA) downscaled climate model projections for the Representative Concentration Pathway (RCP) 8.5 scenario from the CMIP5 Project (comparing the 1971-2000 period to the 2070-2099) (Abatzoglou and Brown, 2012; Taylor et al., 2012). Central Georgia was the most responsive to precipitation reduction, paralleling the spatial patterns in the response to nutrient addition, suggesting that the region is able to support high productivity but is sensitive to nutrient and precipitation levels. The simulated sensitivity was likely due to poor soils (low site index) and low baseline precipitation relative to the warm climate. Our predictions of low sensitivity to precipitation reduction or addition were derived from assimilating observations from throughfall exclusion and irrigation experiments across the region. Prior publications from the studies used in DA also reported low sensitivities to water manipulations, indicating that our predictions are likely not biased (Albaugh et al., 2004; Samuelson et al., 2014; Ward et al., 2015; Wightman et al., 2016). For example, the throughfall exclusion experiment at the focal site in

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698 Georgia, reported a 13% reduction in stem production during a dry year but a 0% reduced during a wet year, resulting in a 7% reduction of productivity over a 2-year period in response to a 30% 699 reduction in throughfall (Samuelson et al., 2014). Our predicted 8.5% reduction to a 30% 700 reduction in precipitation compares well to the observed change, noting that our sensitivity 701 integrated over a 25-year rotation and included a mix of relatively wet and dry years. 702 703 The 3-PG model included a highly simplified representation of interactions between the water 704 and carbon cycles that resulted in parameterizations that, while consistent with observations, may 705 706 contain assumptions that require additional investigation. For example, transpiration is modeled as a potential canopy transpiration that occurred if leaf area was not limiting transpiration. The 707 LAI at which leaf area was no longer limiting was a parameter that was optimized (LAIgex in SI 708 709 Table 3), resulting in a value of 2.3. Interestingly, this optimized value is consistent with the scant literature on this topic. In their analysis of multi-year measurements of transpiration in 710 711 loblolly pine, Phillips and Oren (2001) observed that transpiration per unit leaf area was 712 relatively insensitive to increases in leaf area above LAI of approximately 2.5. Iritz and Lindroth (1996) reviewed transpiration data from a range of crop species and found only small increases 713 in transpiration above LAI of 3-4. These authors suggest that the threshold-type responses 714 observed were related to the range of LAI at which self-shading increases most rapidly, therefore 715 limiting increases in transpiration. The resulting model behavior of "flat" transpiration above 2.3 716 LAI, with gradually decreasing photosynthesis above that value, results in increasing water use 717 efficiency at higher LAI values. The parameterization of the relationships between transpiration 718 and photosynthesis in 3-PG would likely benefit from additional data beyond the two eddy-719 covariance studies with ET observations used here. For example, canopy conductance estimates, 720

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and their associated uncertainty have been derived from assimilating observations from sap-flow measurements into a model that scales from the sensor measurements to canopy transpiration using LAI observations (Bell et al., 2015). This sap-flow to canopy conductance scaling approach (the State Space Canopy Conductance (StaCC) model (Bell et al., 2015)) produces a probability distribution of monthly canopy conductance that could be integrated into the DAPPER system by treating the posterior estimates of StaCC as the distribution of the data in equation 2. Second, the optimized parameters that described the relationship between relative available soil water and the modifier of photosynthesis and transpiration predicted a modifier value greater than zero when the relative available soil water was zero. This resulted in positive values from photosynthesis and transpiration when the average available soil water during the month was zero. In practice, the monthly available soil water was rarely zero during simulations, which presents a challenge constraining the shape of the available soil water modifier. The priors for the two available soil moisture modifiers (SW1 and SW2) had ranges that permitted the modifier to be zero. Therefore, additional data is likely needed during very dry conditions to develop a more physically based parameterization. Alternatively, the parameterization of a nonzero soil moisture modifier at zero available soil water may be due to trees having access to water at soil depths deeper than the top 1.5 m of soil represented by the bucket in 3-PG. Overall, it is important to view the parameterization presented here as a phenomenological relationship that is consistent with observations from throughfall exclusion and irrigation experiments as well as observations across regional gradients in precipitation. Beyond the specifics of the 3-PG modeling efforts, the DA of regional observations into a monthly, computationally tractable ecosystem model can potentially inform Earth system

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modeling efforts. While the details of physiology differ between 3-PG and global land-surface models, the concepts governing NPP allocation are similar. Therefore, DA using the 3-PG model can be used to parameterize the allocation patterns of similar plant types in a global model. One land-surface model, the Community Land Model (CLM), includes parameters that govern the ratio of stem to leaf allocation, ratio of coarse root to stem allocation, and the ratio of leaf to fine root allocation, parameters that are also optimized in DAPPER. As an example, the ratio of fine root to leaf allocation in CLM 4.0 and 4.5 for temperate pine plant function type is set to 1, resulting in equal annual allocation of carbon to foliage and fine roots (Oleson et al., 2013). In contrast, we found that the median ratio of fine root to foliage allocation was substantially lower at 0.13 (Table 6). Therefore, simulations in the CLM with the lower value of root allocation would have higher allocation to aboveground tissues if the loblolly pine parameters from our analysis were used. This would increase carbon accumulation in woody tissues and could alter predictions of nutrient limitation because stems have higher C:N ratios. Other parameters, including the stem to coarse root ratio, are closer to the values used in the CLM.

## **5 Conclusions**

DA is increasingly used for ecological forecasting due to its ability represent prior knowledge, integrate observations into the parameterization, and estimate multiple components of uncertainty, including observation, parameter, and process representation uncertainty (Dietze et al., 2013; Luo et al., 2011b; Niu et al., 2014). Our application of DA to loblolly pine plantations of the southeastern U.S demonstrated that these ecosystems are well suited as a test-bed for the development of DA techniques, particularly techniques for assimilating ecosystem experiments. Further, we found that assimilating ecosystem manipulative experiments into a simple ecosystem

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767 model changed predictions quantifying how forest productivity responds to environmental change, highlighting the importance of networks of ecosystem manipulation experiments for 768 helping to parameterize and evaluate ecosystems models (Medlyn et al., 2015). 769 770 6 Data availability 771 772 Observations used in the DA can be found in the following: Duke FACE study can be found in McCarthy et al. (McCarthy et al., 2010), the PINEMAP studies are available through the TerraC 773 database (http://terrac.ifas.ufl.edu), the DK3 eddy-flux tower data are available through the 774 Ameriflux database (http://ameriflux-data.lbl.gov), the Waycross data can be found in Bryars et 775 al. (2003), the NC2 data are available upon request with Asko Noormets, the FMRC and FPC are 776 available through membership with the cooperatives. The parameter chains and 3-PG are 777 available upon request from R. Quinn Thomas. 778 779 Acknowledgments 780 Funding support came from USDA-NIFA Project 2015-67003-23485 and the Pine Integrated 781 Network: Education, Mitigation, and Adaptation project (PINEMAP), a Coordinated 782 783 Agricultural Project funded by the USDA National Institute of Food and Agriculture, Award 784 #2011-68002-30185. Additional funding support came from USDA-NIFA McIntire-Stennis Program. The Virginia Space Grant Consortium Graduate STEM Research Fellowship Program 785 provided partial support for A. Jersild. Computational support was provided by Virginia Tech 786 Advanced Research Computing. This research was also supported by grants from the French 787 Research Agency (MACACC ANR-13-AGRO-0005 and MARIS ANR-14-CE03-0007). We 788 thank Luke Smallman and Mat Williams for helpful discussions about data assimilation. We

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Table 1. A subset of parameters optimized using data assimilation, prior distributions, and the sensitivity of total biomass at age 25 to the parameter. These are the parameters referred to in the results and discussion, other optimized model parameter can be found in the supplemental material.

Parameter	Parameter description	Units	Sensitivity*	Prior distribution	Prior parameters	Reference for prior
α	Canopy quantum efficiency (pines)	mol C mol PAR <sup>-1</sup>	0.84	Uniform	Min = 0.02 $Max = 0.1$	Vague
у	Ratio NPP/GPP	-	0.84	Uniform	Max = 0.66 Min = 0.30	1
fCalpha700	Proportional increase in canopy quantum efficiency between 350 and 700 ppm CO2	-	0.08	Uniform	Min = 1.05 $Max = 2.0$	Vague
fCpFS700	Proportional decrease in allocation to foliage between 350 and 700 ppm CO2	-	0.00#	Uniform	Min = 0.50 Max = 1.00	Vague
SWconst	Moisture ratio deficit when downregulation is 0.5	-	0.06	Uniform	Min = 0.6 $Max = 1.8$	2, Vague
SWpower	Power of moisture ratio deficit	-	0.06	Uniform	Min = 1 $Max = 13$	2, Vague
FR1	Fertility rating parameter 1 (mean annual temperature coefficient)	-	0.23	Uniform	Min = 0.0 $Max = 1.0$	Vague
FR2	Fertility rating parameter 2 (site index age 25 coefficient)	-	0.39	Uniform	Min = 0.0 $Max = 1.0$	Vague
wSx1000	Maximum stem mass per tree at 1000 trees/ha	kg tree <sup>-1</sup>	0.43	Normal	Mean = 235 $Sd = 25$	3,4
thinPower	Power in self thinning law	-	0.25	Uniform	$ \begin{aligned} Min &= 1.1 \\ Max &= 1.80 \end{aligned} $	3,4
pCRS	Ratio of coarse roots to stem allocation	-	0.08	Uniform	Min = 0.15 Max = 0.35	5

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1(DeLucia et al., 2007); (Landsberg and Waring, 1997), (Bryars et al., 2013), (Gonzalez-Benecke et al., 2016), 5(Albaugh et al., 2005) \* Sensitivity is 1 when a 10% increase in the parameter results in a 10% change in total biomass. (Sensitivity is 0 when a 10% increase in the parameters does not change total biomass by a value greater than 0.01%.

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Table 2. Regional observational data streams used in data assimilation

Data stream	Measurement frequency	Measurement or estimation	Uncertainty	Stream ID for Table	
	requeriey	technique		4	
Foliage biomass	Annual or less	Allometric	Based on propagating the	1	
(Pine)		relationship	allometric model uncertainty		
			in Gonzalez-Benecke et al.		
E 1: 1:	. 1 1	A 11	2014. Varied by observation.	2	
Foliage biomass (hardwood)	Annual or less	Allometric relationship	Assumed zero	2	
Stem biomass (pine)	Annual or less	Allometric	Based on propagating the	3	
Stem biomass (pine)	Ailliuai oi iess	relationship	allometric model uncertainty	3	
		relationship	in Gonzalez-Benecke et al.		
			2014. Varied by observation.		
Stem biomass	Annual or less	Allometric	Assumed zero	4	
(hardwood)		relationship			
Coarse root biomass	Annual or less	Allometric	Standard deviation (SD) =	5	
(combined)		relationship	10% of observation		
Fine root biomass	Annual or less	Allometric	SD = 10% of observation	6	
(combined)		relationship			
Foliage biomass	Annual	Litterfall traps	SD = 2.5% of observation	7	
turnover (combined)		3.61	GD 100/ C 1	0	
Fine root biomass	Annual	Mini-	SD = 10% of observation	8	
turnover (combined)	. 1 1	rhizotrons	10// 1 11)	0	
Pine stem count	Annual or less	Counting individuals	1% (assumed small)	9	
Leaf area index	Monthly to	Litter traps or	If litter trap method: SD =	10	
(pine)	annual	LI 2000	2.5% of observation	10	
(pine)	umuu	E1 2000	If LI-2000 method: SD =		
			10% of observation		
Leaf area index	Monthly to	Litter traps or	If litter trap method: SD =	11	
(hardwood)	annual	LI 2000	2.5% of observation		
			If LI-2000 method: $SD =$		
			10% of observation		
Leaf area index	Only used if not	Litter traps or	If litter trap method: SD =	12	
(combined)	separated into	LI 2000	2.5% of observation		
	pine and		If LI-2000 method: SD =		
G . T	hardwood		10% of observation		
Gross Ecosystem Production	Monthly	Modeled from	SD = 10% of observation	13	
riouuction		flux eddy- covariance net			
		ecosystem			
		exchange			
Evapotranspiration	Monthly	Eddy-	SD = 10% of observation	14	
L vapou anspiration	1410Hully	covariance	5D 10/0 01 003C1 validii	17	

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Number

Table 3. Descriptions of the studies used in data assimilation.

16

12

1

Number

Experimental

Nutrient

addition,

irrigation, nutrient x irrigation

CO<sub>2</sub>, nutrient

addition, CO2

x nutrient

addition

None

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Study name

SETRES4

Duke

flux

1013

1014 1015

1016 1017 FACE<sup>5</sup> and

NC2 Flux

1





Reference

(Albaugh et al.,

(McCarthy et al.,

(Noormets et al.,

2010; Novick et al.,

2004)

2015)

2010)

•	of locations	of plots per site	treatments (plots)	streams (Table 2)	Years	Stand Ages (years)	
FMRC <sup>1</sup> Thinning Study	163	1	None	1, 3,9	1981 - 2003	8 - 30	(Burkhart et al., 1985)
FPC <sup>2</sup> Region- wide 18	18	2	Nutrient addition	1, 3,9	2011-2014	12-21	(Albaugh et al., 2015)
PINEMAP <sup>3</sup>	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)

1,3,5,6,9,

2,3,4,5,6,

7,8,9,10,

11,13,14

2,3,4,5,6,

7,9,10,11 ,12,13,14 1991-2006

1996-2004

2005-2014

8 - 23

13-22

12-22

Measurement

Measurement

Total 187 294 1981 - 2014 4 - 30

<sup>1</sup>Forest Modeling Research Cooperative; <sup>2</sup> Forest Productivity Cooperative; <sup>3</sup> Pine Integrated Network: Education, Mitigation, and Adaptation project (PINEMAP); <sup>4</sup> Southeast Tree Research and Education Site; <sup>5</sup> Free Air Carbon Enrichment

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

Simulation Name	Treatments included in assimilation	Number of plots
All	1-stage data assimilation. All plots and experiments in the	294
	region were used simultaneously.	
DK+NC2	1 <sup>st</sup> stage of 2-stage assimilation. All plots at the Duke eddy flux	13
	(DK3), Duke Free Air CO <sub>2</sub> Enrichment Study, and NC2 eddy	
	flux site; includes CO <sub>2</sub> enrichment and nutrient addition	
	experiments at the Duke site	
DK+NC2-fert	1 <sup>st</sup> stage of 2-stage assimilation. Same as DK+NC2 but without	10
	nutrient fertilization plots	
RW	2 <sup>nd</sup> stage of 2-stage assimilation. Region-wide assimilation of	281
	FRMC, FPC, PINEMAP, Waycross, and SETRES sites. Uses	
	the posteriors of the DK+NC2 simulation as priors. Includes	
	nutrient addition and water manipulation experiments. This	
	simulation is repeated four times for four different out-of-	
	sample validation plots.	
RW-fert	2 <sup>nd</sup> stage of 2-stage assimilation. Same as RW but without	222
	nutrient addition experiments; uses the posteriors of the	
	DK+NC2-fert simulation as priors	
RW-water	2 <sup>nd</sup> stage of 2-stage assimilation. Same as RW but without water	241
	manipulation experiments	

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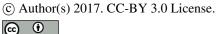




Table 5. Posterior means and 95% credible intervals for parameters listed in Table 1 using the data assimilation approaches listed in Table 4.

Parameter	RW	RW-fert	RW-water	DK+NC2	All
α	0.037	0.040	0.037	0.035	0.032
	(0.034 - 0.040)	(0.036 - 0.045)	(0.035 - 0.040)	(0.030 - 0.042)	(0.030 - 0.035)
y	0.48	0.48	0.48	0.48	0.52
	(0.46 - 0.51)	(0.45 - 0.51)	(0.46 - 0.51)	(0.45 - 0.51)	(0.50 - 0.54)
fCalpha700	1.31	1.31	1.31	1.32	1.11
	(1.22 - 1.40)	(1.22 - 1.40)	(1.22 - 1.40)	(1.23 - 1.41)	(1.08 - 1.15)
fCpFS700	0.84	0.83	0.84	0.84	0.99
	(0.75 - 0.93)	(0.75 - 0.93)	(0.75 - 0.93)	(0.76 - 0.93)	(0.95 - 1.0)
SWconst	1.48	1.31	1.8	1.30	1.57
	(1.09 - 1.85)	(0.95 - 1.70)	(1.47 - 2.15)	(0.89 - 1.76)	(1.08 - 1.79)
SWpower	1.61	1.29	2.93	2.20	1.47
	(0.90 - 2.46)	(0.78 - 1.98)	(1.48 - 3.82)	(1.47 - 3.44)	(1.09 - 2.26)
FR1	0.094	0.096	0.118	not fit	0.094
	(0.086 - 0.104)	(0.088 - 0.103)	(0.110 - 0.128)		(0.087 - 0.102)
FR2	0.144	0.124	0.179	not fit	0.153
	(0.133 - 0.154)	(0.108 - 0.142)	(0.156 - 0.182)		(0.140 - 0.168)
wSx1000	176	180	180	258	181
	(171 - 181)	(174 - 186)	(176 - 186)	(228 - 295)	(174 0 187)
thinPower	1.67	1.70	1.71	1.28	1.61
	(1.60 - 1.74)	(1.63 - 1.78)	(1.65 - 1.78)	(1.12 - 1.60)	(1.51 - 1.69)
pCRS	0.26	0.24	0.25	0.17	0.28
	(0.25 - 0.27)	(0.23 - 0.25)	(0.24 - 0.26)	(0.16 - 0.19)	(0.27 - 0.29)





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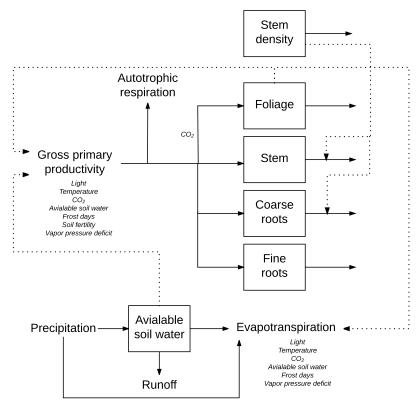


Figure 1. A diagram of the monthly time-step 3-PG model used in this study. The stocks are represented by the boxes and the fluxes by the arrows. An influence of a stock on a flux that is not directly related to that stock is represented by the dotted lines. The environmental influences on a flux is described using italics. A description of the model can be found in the supplemental information.





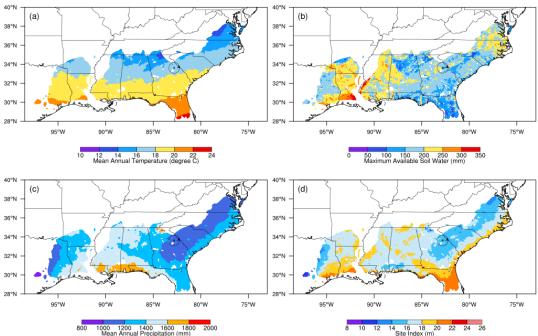


Figure 2. Key climatic and stand characteristic inputs to the regional 3-PG simulations: (a) Mean annual temperature (1979-2011) as a summary of the gradient in monthly temperature inputs used in simulations, (b) maximum available soil water for the top 1.5 meters of soil from SSURGO, (c) mean annual precipitation (1979-2011) as a summary of the gradient in monthly precipitation inputs used in simulations, and (d) site index. The focal site in Georgia highlighted in Figures 5c and 6b is represented by the circle containing the dot. The area shown is the natural range of loblolly pine (*Pinus taeda L*.).

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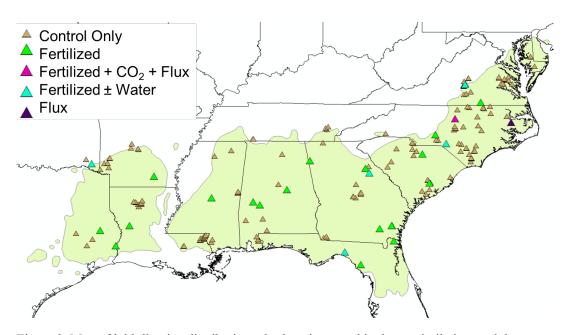


Figure 3. Map of loblolly pine distribution, plot locations used in data assimilation, and the experiment type associated with each plot. The control-only treatments were plots without any associated experimental treatment or flux measurements. Fertilized were plots with nutrient additions.  $CO_2$  were plots with free-air concentration enrichment treatments. The flux treatments were plots with eddy-covariance measurements of ecosystem-scale carbon and water exchange. The water treatments included throughfall exclusion and irrigation experiments.





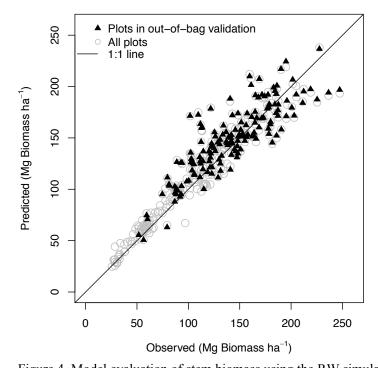
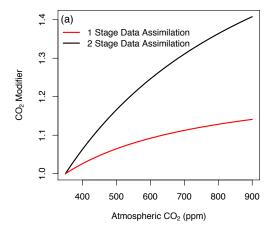


Figure 4. Model evaluation of stem biomass using the RW simulation described in Table 5. The gray circles correspond to predictions where all plots were used in data assimilation. The black triangles correspond to predictions where 120 plots were not included in data assimilation and represent an independent evaluation of model predictions (out-of-bag validation). For each plot, we used the measurement with the longest interval between initialization and measurement for evaluation.







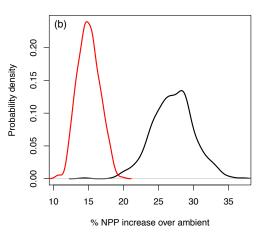


Figure 5. The influence of the data assimilation approach on predictions of how productivity responds to atmospheric CO<sub>2</sub>. (a) The relationship between atmospheric CO<sub>2</sub> concentration and the modifier of light-use efficiency when all plots and experiments are assimilated simultaneously (1 stage) and when the Duke and NC2 plots are assimilated before assimilating the remaining observations across the region (2 stage). (b) The probability distribution of predicted response of NPP to the elevated CO<sub>2</sub> at the Duke FACE experiment for the two assimilation approaches. Uncertainty was estimated by integrating the parameter uncertainty estimated through data assimilation (see Methods).

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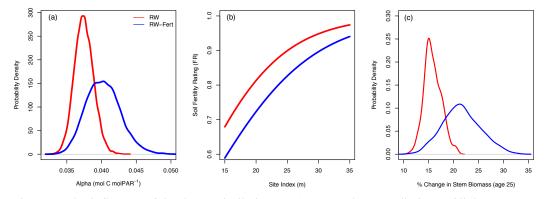


Figure 6. The influence of the data assimilation (DA) approach on predictions of light-use efficiency, nutrient limitation, and how productivity responds to nutrient addition. (a) The posterior distribution for the potential light-use efficiency parameter (alpha). (b) The relationship between site index and the nutrient limitation modifier in 3-PG (FR). (c) The predicted distribution of the response of stem biomass to nutrient fertilization (setting FR = 1) at the focal site in Georgia. The red line corresponds to DA that included nutrient addition experiments. The blue line corresponds to DA that did not include nutrient addition experiments.





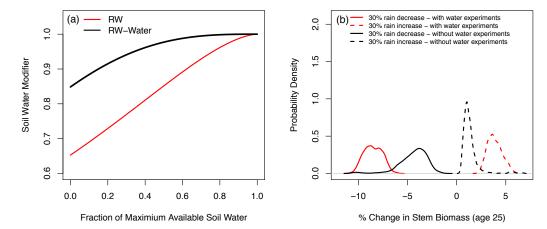


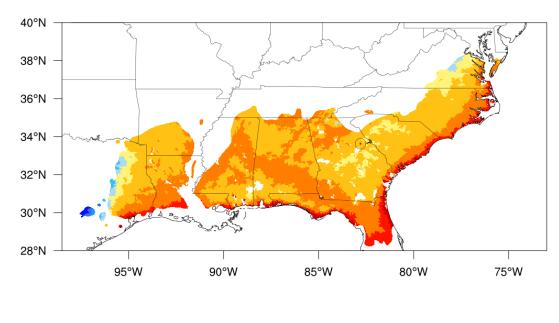
Figure 7. The influence of the data assimilation (DA) approach on predictions of water limitation and how productivity responds to a change in precipitation. (a) The relationship between fraction of maximum available soil water, as predicted by 3-PG, and the modifier of light-use efficiency and canopy conductance. (b) The predicted distribution of the response of stem biomass to a 30% increase (dashed lines) and a 30% decrease (solid lines) in precipitation at the focal site in Georgia. The red line corresponds to DA that included water manipulation experiments. The black line corresponds to DA that did not include water manipulation experiments.

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25 50 75 100 125 150 175 200 225 250 275 300 Age 25 stem biomass (Mg/ha)

1089 1090 1091

Figure 8. Regional predictions of stem biomass stocks for a 25-year-old stand planted in 1979. Parameters used the predictions were from the RW data assimilation approach described in Table 5. The focal site in Georgia highlighted in Figures 5c and 6b is represented by the circle containing the dot.



1095 1096 1097

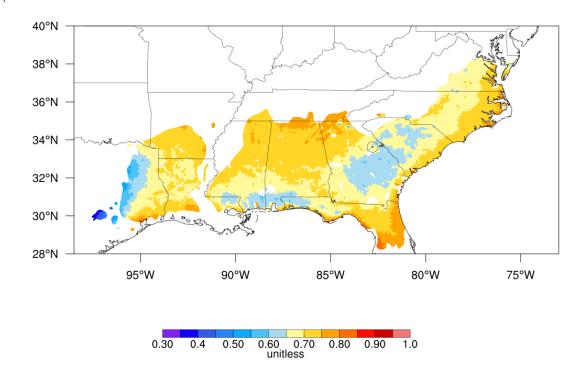


Figure 9: Regional predictions of the soil fertility factor (FR) used in 3-PG to define the nutrient status of the simulated stand. Parameters used the predictions were from the RW data assimilation approach described in Table 5. The focal site in Georgia highlighted in Figures 5c and 6b is represented by the circle containing the dot.





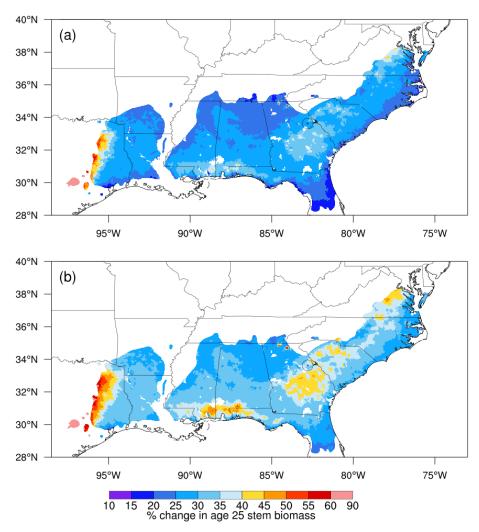


Figure 10: Regional predictions of the change in stem biomass of a 25-year stand when nutrient limitation is completely removed through nutrient addition (simulated by setting FR = 1). Predictions from data assimilation that included nutrient addition experiments are shown in (a) and prediction data assimilation that did not include nutrient addition experiments are shown in (b). The focal site in Georgia highlighted in Figures 5c and 6b is represented by the circle containing the dot.





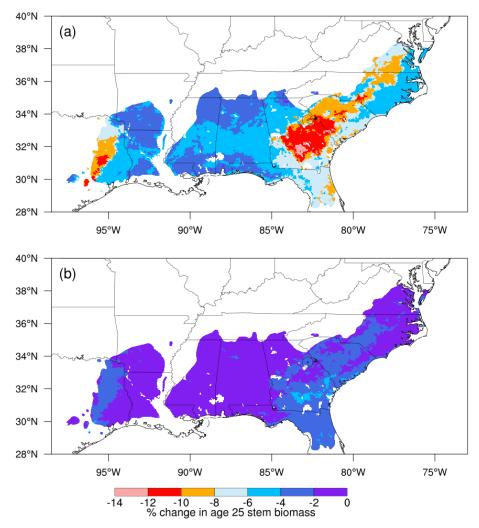


Figure 11. Regional predictions of the change in stem biomass of a 25-year stand when annual precipitation is reduced by 30%. Predictions from data assimilation that included water manipulation experiments are shown in (a) and prediction data assimilation that did not include water manipulation experiments are shown in (b). The focal site in Georgia highlighted in Figures 6c and 7b is represented by the circle containing the dot.