

# Calibrating soybean parameters in CLM-Crop using an MCMC approach

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## **CLM-Crop Development**

- The CLM-Crop model is a agricultural module that relies on a suite of parametric inputs that govern plant growth under a given atmospheric forcing and available resources. (Drewniak et al., 2013)
- CLM-Crop development used measurements of GPP, NEE, and carbon from AmeriFlux data to choose parameter values that optimize crop productivity in the model.



Photo courtesy of David Cook

- Components of CLM-Crop integrated into the CLM4.5
  - Separate organs pool
  - Fertilization
  - Soybean nitrogen fixation
  - Nitrogen retranslocation
  - New carbon nitrogen ratios

### **Motivation**

- Differences between CLM3.5 and CLM4 crop models
- Major concern is crops are not as productive in CLM4
- Need to calibrate the parameters governing growth
- But over 100 parameters
  - Carbon nitrogen ratios :
    - Leaf (pre- and post-grain fill)
    - Stem (pre- and post-grain fill)
    - Organ
- Focus on soybean crop first
- Use data from Bondville, IL AmeriFlux site



# **Calibration Strategy**

Parallel Markov chain Monte Carlo (MCMC) strategy for the CLM-Crop model: multiple chains are run in parallel and exchange information as they evolve

CLM-Crop model calibration setup:



sample posterior distribution for model parameters

 $\pi(\theta|y) \propto \pi(y|\theta)\pi(\theta)$ 

parallel MCMC chains

evolution of live stem C/N value in 8 Markov chains



Performed soy carbon nitrogen ratio calibration; explored C, N & litter pools as calibration parameters

### **MCMC Implementation**

• Parallel adaptive MCMC strategy to determine the posterior distribution of parameters given observations

• Assume prior and likelihood:

$$\log(\pi(\theta)) = -\frac{1}{2}(\theta - \overline{\theta})^T \Sigma_{\theta}^{-1}(\theta - \overline{\theta})$$
$$\log(\pi(y|\theta)) = -\frac{1}{2}(F(\theta) - H(y))^T \Sigma_{\text{obs}}^{-1}(F(\theta) - H(y))$$

• compute posterior using Bayes rule:  $\pi(\theta|y) \propto \pi(y|\theta)\pi(\theta)$ 

• use a parallel MCMC implementation to accelerate and diagnose the convergence by running several Markov chains in parallel while adjusting a Gaussian proposal distribution according to their spread

Standard MCMC algorithm:

Given the target density  $\pi(x)$ , pick a symmetric **proposal density**  $Q(x'|x_t)$ , start with  $x_0$ , t = 0 then proceed as follows: 1. generate a proposed new sample value from  $Q(x'|x_t)$ 

2. calculate 
$$\alpha = \frac{\pi(x')}{\pi(x_t)}$$

3. pick u from uniform 0 and 1, and if  $u < \alpha$  set  $x_{t+1} = x'$ , else set  $x_{t+1} = x_t$ ; set t = t+1, go to 1.

# Validation of the Calibration Strategy

- Simple synthetic example:
  - proposal is auto-tuned
  - chains are run in parallel
  - [Craiu et al 2009]; [Zeng et al 2013]



- CLM-Crop: complex, relatively inexpensive: 2 min/simulated year
- Parallel chains:
  - Convergence is accelerated by adapting the proposal
  - Diagnostics are obtained from (inter-/intra- chain variance)
  - [Zeng et al 2013]



# CLM-Crop Model Calibration: Results using Bondville site observations

- Preliminary calibration results for soy carbon nitrogen ratio with validation
- Performance metric: match the filtered peak and slope during emergence



Obtained new calibrated parameters for soy that give significantly better fit with data

#### Calibration results: soybean (2004)





#### Validation results: soybean (2002)





# **Moving Forward**

- Need to expand
  - Number of parameters
  - Crop types
- Computationally expansive takes too long to converge
- Work on developing surrogate models for parameter calibration



Photo http://landcovertrends.usgs.gov/east/eco70Report.html

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