Towards a representation of priming on soil carbon decomposition in the global land biosphere model ORCHIDEE (version 1.9.5.2).

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Abstract

Priming of soil carbon decomposition encompasses different processes through which the decomposition of native (already present) soil organic matter is amplified through the addition of new organic matter, with new inputs typically being more labile than the native soil organic matter. Evidence for priming comes from laboratory and field experiments, but to date there is no estimate of its impact at global scale and under the current anthropogenic perturbation of the carbon cycle. Current soil carbon decomposition models do not include priming mechanisms, thereby introducing uncertainty when extrapolating short-term local observations to ecosystem and regional to global scale. In this study we present a simple conceptual model of decomposition priming, called PRIM, able to reproduce laboratory (incubation) and field (litter manipulation) priming experiments. Parameters for this model were first optimized against data from 20 soil incubation experiments using a Bayesian framework. The optimized parameter values were evaluated against another set of soil incubation data independent from the ones used for calibration and the PRIM model reproduced the soil incubations data better than the original, CENTURY-type soil decomposition model, whose decomposition equations are based only on first order kinetics. We then compared the PRIM model and the standard first order decay model incorporated into the global land biosphere model ORCHIDEE. A test of both models was performed at ecosystem scale using litter manipulation experiments from 5 sites. Although both versions were equally able to reproduce observed decay rates of litter, only ORCHIDEE-PRIM could simulate the observed priming ($R^2=0.54$) in cases where litter was added or removed. This result suggests that a conceptually simple and numerically tractable representation of priming adapted to global models is able to capture the sign and magnitude of the priming of litter and soil organic matter.
Keywords: soil carbon decomposition, global land biosphere model, priming effect, climate change.
1. Introduction

Soils are the largest reservoir of organic carbon (C) on land, holding three times as much as plant biomass globally (MEA, 2005). The dynamics of long-term soil organic matter formation (Schmidt et al., 2011) and its decomposition on time scales of future climate change (Jones et al. 2003) both remain poorly understood. The lack of a mechanistic understanding of soil carbon dynamics on time scales going from years to centuries induces important differences in the future projections of the global land carbon storage among global land biosphere models (Todd-Brown et al., 2013).

Different conceptual models have been proposed to explain empirical data on soil carbon decomposition, mainly incubation experiments (Wutzler and Reichstein, 2008; Manzoni and Porporato, 2009). Those conceptual models are usually calibrated to fit data (i.e. measurements of stock evolution or fluxes) from experiments on soil incubation, and on time scales going from hours to days (Panikov and Sizova, 1996; Blagodatsky and Richter 1998).

It was shown by Wutzler and Reichstein (2008) that conceptual decomposition models accounting for interactions between labile and more recalcitrant microbial-related carbon, often called “priming effects”, could better fit data from incubation experiments acquired over periods of about 100 days.

The conceptual models of soil carbon decomposition encapsulated in global land biosphere models usually ignore interactions between labile and recalcitrant carbon. All global land biosphere models part of the Earth System Models used for IPCC climate projections are based on donor-pool dominant transfer and first order decay (Luo et al. 2015).

Many of those global land biosphere models have soil carbon modules derived from the CENTURY (Parton et al., 1988) and RothC (Coleman and Jenkinson, 1999) models, in which the first order decay rates of different pools are modulated by soil temperature and moisture, as well as by soil texture (Friedlingstein et al., 2006).
Although the conceptual models with priming showed a more realistic behavior than first order decay models when applied to short term incubation data, one may still wonder if priming significantly influences the dynamics of soil carbon on time scales ranging from years to decades, and at large spatial scales. On the one hand, incorporating priming in a global land biosphere model has the disadvantage of introducing new parameters that are difficult to constrain and of generating a more complex - but unproven - dynamical behavior than the first order decay models. On the other hand, if the performances of first order decay models are not satisfactory at the large scale, structural changes of soil carbon models are needed and must be carefully tested.

The current situation with first-order decay dynamics in global land biosphere is that out of the 11 Earth System models used for the IPCC-AR5 CMIP5 simulations and benchmarked by Todd-Brown et al., (2013) against a global soil organic carbon (SOC) map, only six succeeded in representing the total mean C stocks at the global scale, but all failed to reproduce the spatial heterogeneity of SOC stocks as well as the SOC distribution under different vegetation cover (Todd-Brown et al., 2013). Possible causes of model failure include both errors in model structure but also errors in the different parameters controlling soil carbon dynamics. The optimization of the parameters of a first order decay model against a global SOC map could only partly reduce regional discrepancies with observations, with the optimized model explaining only 41% of the global variability of SOC (Hararuk et al., 2014). On the other hand, the use of a structurally different model that accounted for microbial biomass was shown to produce a rather realistic large-scale SOC variability, but very different soil carbon dynamics in response to future climate change (Wieder et al., 2013). This illustrates that model structure matters a lot for the simulation of the current distribution of soil carbon and its future evolution in response to climate and CO$_2$ changes.

Discrepancies between global land biosphere model predictions and observations are partially due to models lacking key mechanisms controlling SOC dynamics (Schmidt et al.,
One example is the interactions with the N cycle. The majority of the ESMs used for the IPCC-AR5 CMIP5 Earth System simulations did not represent explicitly the nitrogen cycle, but the two ESMs with an explicit nitrogen cycle did not result either in a better simulations of current SOC (Todd-Brown et al., 2013). Another example is the role of microorganisms. The first order kinetics used in most models obviates the role that microbial decomposers are known to play in controlling SOC mineralization (Cleveland et al., 2007; Garcia-Pausas and Paterson 2011), but their activities is controlled by physical and chemical drivers (Kemmit et al., 2008). Therefore, ESMs have significant gaps in reproducing the mechanisms related to microbial dynamics such as priming (see definition below), the object of this study.

Soil C priming is defined as a modification of SOC decomposition rates when fresh organic C (FOC) is added (Kuzyakov et al., 2000). Priming is almost ubiquitously observed in ecosystem studies where organic matter inputs are altered in laboratory incubations (reviewed by Blagodatskaya and Kuzyakov 2008) or directly on the field (Boone et al., 1998; Borken et al., 2002; Chemidlin-Prévost-Bouré et al., 2010; Subke et al., 2004; Sulzman et al., 2005; Xiao et al., 2015). Priming can occasionally be negative but most commonly has a stimulative effect on the decomposition of organic matter that decomposes. Several mechanisms may be involved in controlling priming (Fontaine et al., 2003; Blagodatskaya and Kuzyakov 2008, Guenet et al., 2010b), and conceptual models of priming can have substantial number of parameters making their parameterization quite complex at large scales (Wutzler and Reichstein, 2013). Wutzler and Reichstein (2008) proposed conceptual models summarized into different equations to introduce priming without using too many parameters, but in all cases an explicit representation of microbial biomass was required. Recently, Guenet et al., (2013a) modified the equation proposed by Wutzler and Reichstein (2008) to represent priming without an explicit representation of microbial biomass, assuming that microbial biomass is always at equilibrium with FOC. This assumption is suitable for being
incorporated into ESMs since it adds only one more free parameter compared to the first order kinetic models. This priming scheme was incorporated into the global land biosphere model ORCHIDEE, with the priming parameters statistically calibrated to reproduce the same equilibrium state (in terms of C stocks, after spin up of the model) than the standard version based on CENTURY (Guenet et al., 2013b). Despite its calibration ensuring the same initial state of SOC for England and Wales, the version of ORCHIDEE with priming resulted in a loss of SOC during the late 20th Century, in better agreement with inventory data (Bellamy et al., 2005) than the standard version which produced a continuous SOC gain. In that study, however, the parameters of the priming model were not based on observations but tuned instead to equilibrium SOC values. The objectives of this study are therefore:

- To derive optimal parameter values of a priming model (PRIM) with C inputs forced by data by using a Bayesian method (Tarantola, 1987) with priors and data from 20 different soil incubations.
- To introduce the calibrated PRIM model into the ORCHIDEE ecosystem model version AR5 and evaluate the new version ORCHIDEE-PRIM against independent in situ litter manipulation experiments at ecosystem scale.
- To assess if the priming model significantly improves the simulation of SOC mineralization compared to the standard first order decay model used in ORCHIDEE, on time scales of months to years.
2. **Materials and Methods**

The material and methods section is summarized in Fig. 1.

2.1 **Models presentation**

2.1.1 Soil carbon priming model PRIM

To represent priming, we used the ORCHIDEE soil decomposition module, which is based on the carbon-related modules of CENTURY (Parton et al., 1988). It has three carbon pools (active, slow and passive) and two litter pools (metabolic and structural). SOC decomposition is modulated by soil temperature and moisture functions. Active SOC decomposition is further modulated by a clay function. These functions are the same as in CENTURY but they are driven by soil physical variables calculated at a daily time step by the soil physics of ORCHIDEE (Krinner et al., 2005). The transfers among pools are described using the CENTURY equations with similar parameters (Parton et al., 1988). In the PRIM model, we replaced the CENTURY decomposition equations by those developed by Guenet et al. (2013a) to simulate a priming effect:

$$
\frac{dSOC_{\text{Active}}}{dt} = I - k_{SOC_{\text{Active}}} \times SOC \times (1 - e^{-c \times (Litter_C)}) \times \theta \times \tau \times \gamma
$$

$$
\frac{dSOC_{\text{Slow}}}{dt} = I - k_{SOC_{\text{Slow}}} \times SOC \times (1 - e^{-c \times (Litter_C + SOC_{\text{Active}})}) \times \theta \times \tau
$$

$$
\frac{dSOC_{\text{Passive}}}{dt} = I - k_{SOC_{\text{Passive}}} \times SOC \times (1 - e^{-c \times (Litter_C + SOC_{\text{Active}} + SOC_{\text{Slow}})}) \times \theta \times \tau
$$

with $I$ being the input of C into the pool considered, $k_{SOC}$ the SOC decomposition rate for the active, the slow and the passive pool, $Litter_C$, the sum of all the litter pools of the model. $\theta$, $\tau$, and $\gamma$ are the soil moisture function, the temperature function and the clay function modulating decomposition, respectively. $c$ is a parameter controlling the impact of the fresh organic carbon (FOC) pool on the SOC mineralization rate. Here, we considered that FOC represents all the carbon from pools more labile than the pool being affected as shown in equation (1) to (3). Therefore, FOC is only litter for the active SOC pool, but for the slow SOC pool, FOC is the sum of the litter and the active SOC pool. Finally, for the passive SOC
pool, FOC is the litter and the active and slow carbon pools. The decomposition of the first donor litter pool is described using first order kinetics (4):

$\frac{dLitter - C}{dt} = I - k_{Litter} \times Litter - C \times \theta \times \tau$

In the Wutzler and Reichstein (2008) equation, the SOC mineralization was described by:

$\frac{dSOC}{dt} = I - k_{SOC} \times SOC \times (1 - e^{-c \times MB})$

with $MB$ being the microbial biomass. Unlike Wutzler and Reichstein (2008), our model does not explicitly simulate $MB$ but assumes that MB equilibrates with FOC thus the relationship between MB and FOC is linear. Consequently, we represent priming using a direct relationship between FOC and SOC mineralization. Finally, the moisture, temperature and clay functions are described by equation (6), (7) and (8), respectively with $soil\_moisture$ in m$^3$ H$_2$O m$^{-3}$ of soil, $soil\_temperature$ in Kelvin and $clay$ in %wt:

$\theta = \max(0.25, \min(1, -1.1 \times soil\_moisture^2 + 2.4 \times soil\_moisture + 0.29))$

$\tau = \exp(0.69 \times (soil\_temperature - 303)/10)$

$\gamma = 1 - 0.75 \times clay$

2.1.2 ORCHIDEE and ORCHIDEE-PRIM

ORCHIDEE is a process-based global land biosphere model that calculates the fluxes of CO$_2$, H$_2$O, and heat between the terrestrial land and the atmosphere. The time step of the model is 1/2-hour, and the variations of H$_2$O and C pools are calculated on a daily basis. The model has been evaluated at different scales (sites, regions, globes) and under different climates from the tropics to northern boreal zones (Krinner et al., 2005; Ciais et al., 2005;
Santaren et al., 2007; Piao et al., 2006). ORCHIDEE results from the coupling of three different sub-models. The first one is called SVAT SECHIBA and describes soil water budget and turbulent fluxes of energy and water between the atmosphere and the biosphere (Ducoudré et al., 1993; de Rosnay and Polcher, 1998). The second one is derived from the dynamic global vegetation model LPJ (Sitch et al., 2003) and deals with vegetation dynamics (fire, sapling establishment, light competition, tree mortality, and climatic criteria for the introduction or elimination of plant functional types). The last, called STOMATE (Saclay Toulouse Orsay Model for the Analysis of Terrestrial Ecosystems) deals with phenology and carbon dynamics of the terrestrial biosphere. Twelve plant functional types (PFT) are used to classify the vegetation. Each PFT dynamic is controlled by similar set of governing equations but using different parameter values. Only the leafy season onset and offset, are PFT-specific (Krinner et al., 2005).

The simulation of SOC in ORCHIDEE version is based on CENTURY (Parton et al., 1988) as described above. No vertical description of the SOC is included in the ORCHIDEE version used here. In ORCHIDEE-PRIM we replaced CENTURY by the PRIM model described in section 2.1.1.

2.2 Data description

2.2.1 Incubation experiments to calibrate the priming model

We optimized the PRIM parameters and the ORCHIDEE soil module parameters using data from soil incubation experiments where FOC was added and the priming effect was measured by comparing a control study without FOC with a perturbation study with FOC (table 1). The data come from 20 incubations (from nine studies) of duration going from one week to 10 months. The incubated soil samples have very different characteristics (table 1) and came from different ecosystems (grassland, cropland, broadleaf forest, needleleaf forest,
savannah). However, the great majority of the data used to optimize the model were obtained from temperate soils. In the incubation experiments, added FOC was labeled with $^{13}$C or $^{14}$C and therefore the respired CO$_2$ fluxes coming from either SOC already present before the FOC amendments or from the FOC induced priming of SOC pools was estimated separately. We used only incubations performed during at least 7 days to eliminate all studies that potentially observed apparent priming effects. Apparent priming is a replacement of the $^{12}$C in microbial biomass with labeled carbon isotopes, a short-term artifact due to the amendment of labeled material to an unlabelled soil (Blagodatskaya and Kuzyakov, 2008). Moreover, we used only studies that reported cumulative respired CO$_2$ fluxes in order to optimize the priming parameters against the extra CO$_2$ fluxes obtained at the end of the experiment and not those resulting from short-term priming dynamics, since cumulative mineralization integrates the different processes occurring during incubation. Finally, several treatments might be performed in the studies used to optimize the model (different soils, different types and amount of FOC). When the treatments performed differed on aspects reproducible by the model (amounts of FOC added, different clay content in the soils used, etc.) we considered all the treatments. In the opposite case we averaged the results of the different treatments to perform the optimization except in case where the treatments clearly impact the results without the possibility to reproduce the experimental design with the model (addition of mineral N for instance).

We also use the control incubations without FOC amendments to evaluate both models. We extracted data from the figures of original publications (Table 1) using GraphClick version 3.0. Several input variables are needed to run the soil model, as described in section 2.1.1. When data were not available from the surveyed publications, we obtained them from the databases normally used for running ORCHIDEE, except for the C:N ratio of FOC and for clay content where data came from Rodal et al., (1960) and from USDA (http://soils.usda.gov/technical/classification/osd/index.html), respectively. The three
carbon pools of CENTURY are not measurable (Six et al., 2002), so we cannot estimate how much C of in each pool is present in the incubated samples. To calculate the distribution of C among the three pools of the model we ran ORCHIDEE until equilibrium was reached at the sites where soil samples were taken and calculated the percentage of each pool.

### 2.2.2 Incubation data used for evaluation of the priming model

A first evaluation of the soil carbon model with and without priming is performed at the scale of soil samples against independent data from the large database of soil incubations (300 in total) published by Moyano et al., (2012). Within this database we selected the experiments where all the inputs necessary to run the two soil carbon models were available (clay, content, moisture, temperature, SOC content at the beginning of the incubation) and where cumulative mineralization or mineralization rates associated to the time step between two measurements were reported. We removed all the studies without information on the location since geographical coordinates are necessary to run ORCHIDEE and thus estimate the initial fraction of each pool. We selected only data coming from experiments without important soil manipulation (e.g. compaction, litter amendments). The model evaluation was performed against a set of 164 independent incubation experiments.

### 2.2.3 Ecosystem-level data used for evaluation of the priming model

A second evaluation of the ORCHIDEE-PRIM model was performed at ecosystem scale against observations of four litter manipulation experiments (Boone et al., 1998; Chemidlin-Prévost-Bouré et al., 2010; Subke et al., 2004; Sulzman et al., 2005) and one compost amendment experiment (Borken et al., 2002). In the litter experiments, two treatments and a control are generally performed. The treatments are total exclusion of above ground litter using nets to prevent fresh litter from falling onto the soil, often transplanting the collected fresh litter to create a second treatment with doubled aboveground litter inputs (Boone et al., 1998; Chemidlin-Prévost-Bouré et al., 2010; Sulzman et al., 2005). For the compost
amendment experiment by Borken et al. (2002), 1.4 kg C m$^{-2}$ (and a zero-addition control) of compost was added to the soil. These studies are presented in table 3. When information about soil clay content was not available in the original study, we extracted it from Zobler (1986). The data measured at field scale are the soil CO$_2$ efflux including the heterotrophic respiration but also root respiration in the same flux without clear separation of the two components.

2.3 Optimization procedure

For PRIM, the 6 parameters optimized are turnover rate ($k_{SOC}$) and priming parameters $c$ for each of the three pools (table 2). For the ORCHIDEE soil module, only the three $k_{SOC}$ values are optimized. The same parameters are optimized against the priming incubations dataset described in 2.2.1. Since optimizations were performed using soil incubations data obtained at optimal temperature and soil moisture, we did not optimize the parameters related to the eq. (6) and (7) because the range of observations was quite limited. Optimization was performed in the framework of the Bayesian inversion method with priors (Tarantola, 1987) as described by Santaren et al., (2007) using assimilating all data streams in the same cost function. Assuming that all uncertainties follow Gaussian distributions (parameter error, measurement error, model error), the optimized parameters correspond to a set minimizing the following quadratic cost function:

\[
J(x) = \frac{1}{2} \left[ (y - H(x))^T R^{-1} (y - H(x)) + (x - x_b)^T P_b^{-1} (x - x_b) \right]
\]

The cost function defined by equation (9) contains both the mismatch between model outputs and observed data, and the mismatch between optimized parameters and the prior values. The mismatch is weighted by errors of each quantity. $x$ is the of unknown parameters vector, $x_b$ the prior values, $y$ the observations vector and $H(x)$ the model outputs. $P_b$ is the
prior parameter error variances/covariances, and $R$ contains the observational error variances/covariances which represents both measurement uncertainty and model uncertainty.

To minimize the cost function, we used a gradient-based iterative algorithm, called L-BFGS-B (Zhu et al., 1995). A range of values for all the parameters is prescribed by called L-BFGS-B. At each iteration, the cost function $J(x)$ gradient is calculated, with respect to the six parameters. When $J(x)$ is minimized, using a classic finite difference method, we further calculated the posterior error covariance matrix on the parameters $P_a$ from the prior error covariance matrices and the Jacobian of the model at the minimum of the cost function, using the linearity assumption (Tarantola, 1987). When error correlations are close to 1 it suggests that the observations do not permit to clearly separate the effect of two parameters.

The model $H(x)$ is non linear and therefore the approach to minimize the cost function is sensitive to potential local minima. We get around by performing 30 optimizations with different sets of prior parameter randomly distributed within their variation range. We then used the case providing the lowest cost function. This approach reduces drastically the sensitivity to potential local minima as illustrated in Santaren et al. (2014).

We defined the prior ranges of decomposition rates using literature data (Parton et al., 1988; Gignoux et al., 2001). However, only two studies already estimated the $c$ parameter before (Guenet et al, 2013a, Guenet et al., 2013b), its prior value is therefore considered as non-informative and we set a large error on the prior (50%). As for the variance of the model-data mismatch term in the cost function of equation (9), note that with our formalism this error should include both the model error (for instance the model capability to represent the measurement) and the measurement error. Given that the error on the measurements was difficult to estimate precisely for each study, we fixed it to 5% of the mean observed CO$_2$ flux assuming that all incubation data were independent. At its minimum, $J(x)$ should be close to
half the number of observations (reduced $\chi^2$ of one). We assumed that all errors (the observations and on the a priori parameters) are uncorrelated.

2.4 Simulations protocol

2.4.1 Simulation protocol for the soil priming model PRIM

Simulations were performed for each incubation experiment presented in 2.2.1 (table 1) as well as for the evaluation sites in 2.2.2. The simulations of the stand-alone PRIM carbon model (i.e. unplugged from the ORCHIDEE full ecosystem model) were run at a daily time step using FOC inputs from table 1 or from the Moyano et al., (2012) database. No spin-up was performed. We started the simulation by prescribing to the soil carbon models with and without priming an initial amount of SOC equal to that measured in the study considered, distributed among active, slow and passive pools as explained in section 2.2.1 At each time step we increment the cumulative heterotrophic respiration coming from SOC mineralization, so that this cumulative simulated CO$_2$ flux can be compared to data from the end of the incubation experiment. Simulations were performed using R 3.0.2.

2.4.2 Simulation protocol for ORCHIDEE-PRIM and ORCHIDEE

We ran ORCHIDEE and ORCHIDEE-PRIM at each litter manipulation site presented in table 3 using, 6 hourly climate data obtained from the combination of two existing datasets: the Climate Research Unit (CRU) (Mitchell et al. 2004) and the National Centers for Environmental Prediction (NCEP) (Kalnay et al., 1996). Both models were run using the first ten years of the climate forcing (1901-1909) repeated in a loop, and an atmospheric CO$_2$ value corresponding to the year 1901. When the simulated relative yearly change of the SOC stock was less than 0.01%, we considered that SOC equilibrium was reached. Once pre-industrial equilibrium was reached in each grid point, we run transient simulations from 1901 until the
beginning of the manipulation experiment assuming no land use change driven by
reconstructed climate and observed CO$_2$. Then when the simulation reached the year at which
the litter manipulation experiment began, we modified the input of above-ground litter in the
same proportion than in the actual manipulation experiments. Finally, we ran the model for
each treatment during a period corresponding to duration of each experiment.

2.5 Model evaluation

The model evaluation was performed in two steps. First, we evaluated separately
PRIM and the standard first order decay model with their optimized parameters, as stand
alone decomposition models, i.e. unplugged from the ORCHIDEE ecosystem model. To
evaluate the stand-alone soil models, we used incubation data coming from Moyano et al.,
(2012) as described in 2.2.2. Secondly, we evaluated ORCHIDEE and ORCHIDEE-PRIM,
against litter manipulation experiments (see 2.2.3).

To compare model outputs with data we used different metrics. First a linear mixed
effect model with intercept value forced to zero using model outputs as the variable to
explain, and data as the fixed effect and the study where data came from as random effect.
This approach aimed to take into account the fact that incubations performed within the same
study are not independent because they were performed and analyzed by the same team. The
linear-mixed effect model gives the slope of the relationship as output. A slope close to one
indicates that the model reproduces the data well. Then, we used the Normalized Standard
Deviation (NSD) or ratio of model to observed standard deviations ; $NSD = 1$ means that the
model perfectly reproduces the observed standard deviations across experiments:

$$NSD = \sqrt{\frac{\frac{1}{n} \times \sum_{i=1}^{n} (x_i - \bar{x})^2}{\frac{1}{n} \times \sum_{i=1}^{n} (o_i - \bar{o})^2}}$$
where \( x \) refers to the model value, \( o \) to the observed value and \( n \) the number of samples. Finally, we compared model performance using the Bayesian Information Criterion \((BIC)\) to take into account that the PRIM soil model has three more priming parameters (one per pool) than the standard model:

\[
BIC = \log(MSD) \times n + \log(n) \times p
\]

with \( MSD \) being the mean squared deviation derived from equation (12), \( n \) the number of data used to evaluate the model, and \( p \) the number of parameters of the soil model.

\[
MSD = \frac{\sum (m-o)^2}{n}
\]

with \( o \) the observed values, \( m \) the values calculated by the model and \( n \) the number of observations. The lowest is the BIC the better the model is.

3. Results

3.1 Optimized parameters of the priming model

The parameters obtained after optimization using incubation data described in section 2.2.1 are given in Table 2. The turnover times ranged from a few months \((0.30 \pm 0.15 \text{ year})\) for the active pool to \(462.0 \pm 233.8 \text{ years}\) for the passive pool, the slow pool being intermediate with \(1.12 \pm 0.01 \text{ years}\). The priming parameters indicated a decreasing sensitivity with increasing turnover time. The parameter \( c \) values were \(493.7 \pm 246.8, 194.0 \pm 97.0 \text{ and } 136.5 \pm 68.3 \text{ for the active, slow and passive pools, respectively. Errors correspond to the estimates from the linear assumption at the minimum of } J(x). \text{ For both, the correlation between parameters was low (data not shown).}

After optimization, both models with and without priming parameterization were able to reproduce the cumulative mineralization measured in the different incubations where FOC was added well (Fig. 2, top panel). The slope of the linear regression between optimized model output and incubation measurements was 1.13 for PRIM and 0.93 for the ORCHIDEE...
soil module. The NSD value (1.80 and 1.52 for PRIM and the standard soil module, respectively) showed that the models overestimated the variance after optimization. When both models were evaluated against the same incubation experiments but without FOM addition, the PRIM model slightly over-estimated accumulated mineralization (Fig. 2 middle panel), as indicated by the value of the slope (1.05). Nevertheless, it performed better than the standard soil module, which underestimated the soil mineralization as indicated by the value of the slope (0.72). The PRIM soil model reproduced quite well the observed priming effect (section 2.2.1) as shown in Fig. 2 (lower panel) with a slope value (1.07). PRIM largely overestimated however the variance of data as indicated by the NSD value (3.14). As expected, the standard soil module was totally unable to reproduce priming (Fig. 2, lower panel).

### 3.2 Standard soil module vs. PRIM against incubations data

To evaluate the performance of PRIM we tested it against data from soil incubation experiments independent from those used for optimization (see section 2.2.2). We did the same with the standard soil module (Fig. 3). The standard soil module tended to overestimate accumulated mineralization as indicated by a slope value of 1.32 and to underestimate the cross-experiments variance by more than 50% (NSD=0.44). PRIM performed slightly better, but underestimated accumulated mineralization (slope 0.80). The optimized PRIM underestimated the variance by 29%, but the NSD value (0.71) was closer to 1 compared to the standard model. Using the BIC index, which takes into account the higher number of parameters of PRIM, this model still performed better (BIC values of 546.2 vs. 347.4 for standard and PRIM, respectively).

### 3.3 ORCHIDEE vs. ORCHIDEE-PRIM comparison using in situ datasets
When tested at ecosystem-level against litter manipulation experiments, 4 studies x 3 treatments and 1 study with 2 treatments. Both ORCHIDEE and ORCHIDEE-PRIM performed generally well to reproduce the soil CO$_2$ efflux (Fig. 4). Generally, both versions showed similar performance as indicated by the values of slopes and NSD presented in table 4. The mean slopes are 0.98 for ORCHIDEE-PRIM against 0.97 for ORCHIDEE, and the mean NSD are 1.26 and 1.27, respectively. It must be noted that slope values were generally lower for the treatments excluding litter compared to control and double litter inputs (Table 4). No particular differences of the NSD values were observed between the different litter input regimes. Nevertheless, the BIC index was always higher for ORCHIDEE-PRIM because three more parameters were used by this version compared to ORCHIDEE.

ORCHIDEE-PRIM was able to reproduce the priming observed defined as the difference of CO$_2$ efflux coming from SOC only with or without litter (Fig. 5), but tended to underestimate its intensity as indicated by the slope value lower than one (0.55). The variance between experiments calculated for priming was overestimated as shown by the NSD value of 1.29. It must be noted that priming was not calculated for ORCHIDEE since the structure of its soil decomposition model does not include a priming mechanism.

### 4. Discussion

#### 4.1 PRIM in the context of other soil priming conceptual models

Priming is a complex phenomenon controlled by several mechanisms, such as N mining by microbial communities with different growth strategies, competition between microbial groups for substrate, energy limitations, etc. (Kuzyakov et al., 200; Fontaine et al., 2003; Guenet et al., 2010b). Priming may have important consequences on the feedbacks between climate and C cycle (Schmidt et al., 2011) and it is therefore crucial to better quantify the C fluxes due to priming, especially at large scale (i.e, continental to global). Several models have been developed to describe soil C mineralization with a representation of priming (Gignoux et al., 2001; Fontaine and Barot, 2005; Neill and Gignoux, 2006; Moorhead...
and Sinsabaugh, 2006; Wutzler and Reichstein, 2008; Neill and Guenet, 2010; Blagodatsky et al., 2010) and such models generally succeeded at reproducing short-term data, mainly incubation. However, to our knowledge, they have never been tested in a range of contrasted situations (different soil types, different FOC amount and chemical composition, different temperature and soil moisture, etc.). Here, we used most of the available incubation data respecting the criteria described in the material and method section. Moreover, previous priming models all needed a high number of parameters compared to PRIM. For these two reasons, the conceptual soil models accounting for soil priming were thus far not included in global land biosphere models (Wutzler and Reichstein, 2008) and very few studies of soil priming at global scale have been performed (Foereid et al., 2014). Here, using a simple scheme with only three additional parameters than the standard soil module of ORCHIDEE, we were able to reproduce priming but also soil mineralization data coming from very different incubation studies performed with different soils at different temperature and moisture, with different time length, etc. The PRIM soil model, which is a microbial steady-state model, might not be able to reproduce short-term response to abrupt change of FOC inputs but with negligible bias over the long term (Wutzler and Reichstein 2013). However, it might have similar performances than more complex models to reproduce long-term trends of FOC inputs (Wutzler and Reichstein 2013). PRIM performed better than the standard soil module to reproduce soil incubation data used to optimize, but it must be noted that the BIC values indicate that the improvement observed with PRIM may be simply due to a higher number of parameters. Nevertheless, when using independent soil incubations data from the one used to optimize the model the improvement is quite clear with BIC values much lower with PRIM than with the standard soil module (347.4 and 546.2, respectively). Furthermore, PRIM was not able to fully catch the observed variability of priming. As discussed above, priming is a complex phenomenon resulting from the interactions of different mechanisms that we summarized in a very simple equation. Therefore, PRIM is probably good in representing a general trends but not all the complexity of the phenomenon. Nevertheless, the
use of the PRIM soil model seems justified since it increases only slightly the number of
parameter of a global land biosphere model and since the parameter values were obtained
after optimization on data coming from incubations performed in a range of soils and
conditions (different soil types, different ecosystems, different temperatures, different
moistures, different amount and type of FOC amended, etc.).

4.2 ORCHIDEE vs. ORCHIDEE-PRIM

4.2.1 Cross sites evaluation

ORCHIDEE-PRIM exhibited similar performance than ORCHIDEE when simulating
litter manipulation experiments. It must be noted that both versions share the same scheme for
primary production (controlling soil C input by litter), soil temperature and moisture function.
The similar performance obtained by the two versions may be due to a model bias for these
quantities as well as poorly constrained site histories and climate forcing errors. Since primary
production is the main driver of the C input into the soil, the soil CO₂ efflux calculated by the
models was largely driven by the capacity of the model to reproduce the observed primary
production. In particular, both models largely underestimated the soil CO₂ efflux when litter
was removed (Table 4), but obtained good results when litter was kept or when litter was
added. This suggests that both models performed quite well when reproducing soil CO₂
efflux, but this was due to bias compensation, meaning that the fraction of CO₂ coming from
soil mineralization and root respiration was underestimated and the fraction of CO₂ coming
from litter mineralization was overestimated. Moreover, the modification of the litter cover
may change the soil humidity and temperature and these effects were not represented in the
models.

Finally, the use of microbial steady state model like ORCHIDEE-PRIM present some
advantages compared to explicit microbial models. Wieder et al., (2015) identified several
challenges related to the incorporation of explicit microbial models in ESMs. In particular, it
needs much more parameter than the classical approach. With ORCHIDEE-PRIM these
difficulty is resolved since we only add three more parameters.

5. Conclusion

Regarding the several processes that may lead to priming, the satisfactory performance
of ORCHIDEE-PRIM compared to observations from both laboratory incubation and field
litter manipulation experiments suggests that the simple PRIM conceptual model simulates
well the magnitude of observed priming. Consequently, ORCHIDEE-PRIM has the potential
to quantify the impact of priming on the soil C cycle at large scales. Nevertheless,
ORCHIDEE-PRIM underestimates the priming intensity as shown by the slope value (0.55),
indicating that the model still misses important mechanisms explaining the observations. In
particular, N availability is an important driver of priming, inducing higher priming when N
availability is reduced (Fontaine et al., 2004; Blagodatskaya et al., 2007). The role of N in the
priming intensity as well as the extra N mineralization induced by priming and its effect on
primary production may represent the next addition to the soil representation in a land surface
model by adding a control on the $c$ parameter depending on the mineral N availability and on
the C:N ratio of the considered pool. Nevertheless, some detailed information on the N
dynamic in priming effect experiments would be necessary to do so and very few authors
reported the impact of priming effect on N dynamic after FOC additions.

Code availability

For ORCHIDEE, the main part of the code was written by Krinner et al., (2005). The
version used here is the 1.9.5.2 version. In this version, compared to the one presented in
Krinner et al., (2005), the albedo representation was improved (Hourdin et al., pers. com.), a
routing scheme controlling the flux of water from land surface to the ocean was added (Ngo-
Duc et al., 2007) and the dynamic of vegetation was modified (Viovy et al., pers. com.).
Furthermore, since 2005 the code has been parallelized. A detailed documentation and the
code can be provided upon request to the corresponding author.

ORCHIDEE-PRIM is derived from ORCHIDEE with the modifications presented in
the section 2.1.2. A detailed description can be found in Guenet et al., (2013). The code is
available upon request to the corresponding author.
References


Cleveland CC, Nemergut DR, Schmidt SK, Townsend AR (2007) Increases in soil respiration following labile carbon additions linked to rapid shifts in soil microbial community composition. Biogeochemistry 82: 229-240


Conde, E., Cardenas, M., Poncemendoza, a, Lunaguido, M., Cruzmondragon, C., Dendooven, L., 2005. The impacts of inorganic nitrogen application on mineralization of C-labelled maize and glucose, and on priming effect in saline alkaline soil. Soil Biology and Biochemistry 37, 681–691.


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Figure legends

**Figure 1:** Summarizing scheme of the methods

**Figure 2:** Scatter plot between data and the PRIM model outputs for the incubations with FOC amendment (a), without FOC amendment (b) and for priming effect (c). The dataset used here are the similar to those used for optimization (a) or are the control incubations (b) and are described in section (2.2.1). Red lines indicate the 1:1 line. Different symbol indicate different studies.

**Figure 3:** Scatter plot between independent data from optimization (dataset describes in section 2.2.2) and the soil module of ORCHIDEE outputs (a) or between data and the PRIM model outputs (b). Red lines indicate the 1:1 line.

**Figure 4:** Soil CO$_2$ efflux calculated by ORCHIDEE on the left side and by ORCHIDEE-PRIM on the right side for the data coming from Boone et al., (1998) (a), from Borken et al., (2002) (b), from Chemidlin-Prévost-Bourré et al., (2010) (c), from Subke et al., (2004) (d) and from Sulzman et al., (2005) (e). Red lines indicate the 1:1 line, black, dashed and dotted lines correspond to control, litter exclusion and litter amendment situations respectively.

**Figure 5:** Scatter plot between the priming effect measured and the priming effect calculated by ORCHIDEE-PRIM. Red line indicate the 1:1 line and different symbol indicate different studies.
Figure 1

Incubation data with FOC addition (n = 20)

Incubation data without FOC addition

Other incubation data (n = 164)

Litter manipulation control (n = 5)

Difference of respired CO$_2$ flux = “priming”

Optimized PRIM soil decomposition model

ORCHIDEE soil module (first order decay)

Bayesian optimization

Optimized PRIM soil decomposition model

ORCHIDEE soil module (first order decay)

Optimized PRIM + ORCHIDEE

Standard soil decomposition model + ORCHIDEE

Respired CO$_2$ flux

Time scale $\approx$ 100 days
Space scale: soil unit
Processes: decomposition

Time scale $\approx$ 100 days
Space scale: soil unit
Processes: decomposition

Time scale $\approx$ few years
Space scale: ecosystem
Processes: all modeled ecosystem processes, soil physics, decomposition

Litter manipulation (n = 9)

Other incubation data (n = 164)

Respired CO$_2$ flux

Difference of respired CO$_2$ flux = “priming”

Optimized PRIM soil decomposition model

ORCHIDEE soil module (first order decay)
Figure 2

(a) Cumulative mineralization in FOC amended incubations (PRIM) (gC-CO$_2$.kg$^{-1}$ dried soil)
Slope = 1.13
NSD = 1.80
Pearson's corr. = 0.66
BIC = 31.48

(b) Cumulative mineralization in control incubations (PRIM) (gC-CO$_2$.kg$^{-1}$ dried soil)
Slope = 1.05
NSD = 1.51
Pearson's corr. = 0.73
BIC = 11.55

(c) Cumulative priming (PRIM) (gC-CO$_2$.kg$^{-1}$ dried soil)
Slope = 1.07
NSD = 3.14
Pearson's corr. = 0.24
BIC = 1.05

(d) Cumulative mineralization in FOC amended incubations (ORCHDEE soil module) (gC-CO$_2$.kg$^{-1}$ dried soil)
Slope = 0.93
NSD = 1.47
Pearson's corr. = 0.68
BIC = 19.09

(e) Cumulative mineralization in control incubations (ORCHDEE soil module) (gC-CO$_2$.kg$^{-1}$ dried soil)
Slope = 0.72
NSD = 1.02
Pearson's corr. = 0.72
BIC = 6.91

(f) Cumulative priming (ORCHDEE soil module) (gC-CO$_2$.kg$^{-1}$ dried soil)
Slope = -0.97
NSD = 9.41
Pearson's corr. = -0.33
BIC = 9.50
Figure 3
Soil CO$_2$ fluxes modelled (gC-CO$_2$.m$^{-2}$.day$^{-1}$)

Soil CO$_2$ fluxes measured (gC-CO$_2$.m$^{-2}$.day$^{-1}$)

Figure 4
Figure 5

- Measured priming (% of soil respiration without aboveground litter)
- Modelled priming (% of soil respiration without aboveground litter)

Slope = 0.55
NSD = 1.29
Pearson’s corr. = 0.54
<table>
<thead>
<tr>
<th>Study</th>
<th>Incubations</th>
<th>Study site</th>
<th>FOC types</th>
<th>Amount of FOC amended (g C kg(^{-1}) dry soil)</th>
<th>Lignin:C ratio of FOC</th>
<th>C:N ratio of FOC</th>
<th>Soil clay content (%)</th>
<th>Temperature (K)</th>
<th>Moisture (% of Field Capacity)</th>
<th>Incubation length (days)</th>
<th>SOC content (g C kg(^{-1}) dry soil)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bell et al., (2003)</td>
<td>Experience 1 averaged over the 4 soils tested</td>
<td>Ralston, Washington State, USA</td>
<td>Wheat straw</td>
<td>1.54</td>
<td>0.22*</td>
<td>128*</td>
<td>0.08</td>
<td>298.15</td>
<td>0.2</td>
<td>31</td>
<td>10.1</td>
</tr>
<tr>
<td>Blagodatskaya et al., (2007)</td>
<td>GL treatment</td>
<td>Ramon, Voronezh region, Russia</td>
<td>Glucose</td>
<td>0.0487</td>
<td>0</td>
<td>0</td>
<td>0.2*</td>
<td>295.15</td>
<td>0.6</td>
<td>14</td>
<td>50</td>
</tr>
<tr>
<td>Conde et al., (2005)</td>
<td>Soil A+ Maize</td>
<td>Former lake Texcoco in the valley of Mexico City (Mexico)</td>
<td>Maize</td>
<td>1</td>
<td>0.575</td>
<td>39.8</td>
<td>0.22</td>
<td>295.15</td>
<td>0.55</td>
<td>28</td>
<td>53</td>
</tr>
<tr>
<td></td>
<td>Soil A + Glucose</td>
<td>former lake Texcoco in the valley of Mexico City (Mexico)</td>
<td>Glucose</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0.22</td>
<td>295.15</td>
<td>0.55</td>
<td>28</td>
<td>53</td>
</tr>
<tr>
<td></td>
<td>Soil B + Maize</td>
<td>former lake Texcoco in the valley of Mexico City (Mexico)</td>
<td>Maize</td>
<td>1</td>
<td>0.575</td>
<td>39.8</td>
<td>0.054</td>
<td>295.15</td>
<td>0.55</td>
<td>28</td>
<td>38.8</td>
</tr>
<tr>
<td></td>
<td>Soil B + Glucose</td>
<td>former lake Texcoco in the valley of Mexico City (Mexico)</td>
<td>Glucose</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0.054</td>
<td>295.15</td>
<td>0.55</td>
<td>28</td>
<td>38.8</td>
</tr>
<tr>
<td>De Nobili et al., (2001)</td>
<td>Experiment 1 with Soil 2</td>
<td>Rothamsted experimental station, UK</td>
<td>Cellulose</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0.24</td>
<td>298.15</td>
<td>0.5</td>
<td>11</td>
<td>14.8</td>
</tr>
<tr>
<td>Falchini et al., (2003)</td>
<td>Average over the tree treatments</td>
<td>Grassland in Tuscany, Italy</td>
<td>Oxalix acid/ Glutamic acid/ Glucose</td>
<td>0.1815</td>
<td>0</td>
<td>0</td>
<td>0.14</td>
<td>298.15</td>
<td>0.5</td>
<td>7</td>
<td>16.6</td>
</tr>
<tr>
<td>Fontaine et al., (2004)</td>
<td>Only one incubation</td>
<td>Lamto experimental station, Ivory Coast</td>
<td>Cellulose</td>
<td>0.495</td>
<td>0</td>
<td>0</td>
<td>0.1</td>
<td>231.15</td>
<td>0.032</td>
<td>70</td>
<td>10.5</td>
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<tr>
<td>Guenet et al., (2010)</td>
<td>S1 without N amendment</td>
<td>La cage experimental station, France</td>
<td>Wheat Straw</td>
<td>1.5</td>
<td>0.22*</td>
<td>44</td>
<td>0.167</td>
<td>293.15</td>
<td>0.17</td>
<td>80</td>
<td>10.4</td>
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</tr>
<tr>
<td>S2 without N amendment</td>
<td>La cage experimental station, France</td>
<td>Wheat Straw</td>
<td>2.2</td>
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<td>0.167</td>
<td>293.15</td>
<td>0.17</td>
<td>80</td>
<td>10.4</td>
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<tr>
<td>S3 without N amendment</td>
<td>La cage experimental station, France</td>
<td>Wheat Straw</td>
<td>3.2</td>
<td>0.22*</td>
<td>44</td>
<td>0.167</td>
<td>293.15</td>
<td>0.17</td>
<td>80</td>
<td>10.4</td>
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<tr>
<td>Guenet et al., (2012)</td>
<td>Arable soil with high cellulose input</td>
<td>Closeaux experimental station, France</td>
<td>Cellulose</td>
<td>5</td>
<td>0</td>
<td>0</td>
<td>0.167</td>
<td>293.15</td>
<td>0.19</td>
<td>209</td>
<td>19.9</td>
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<td>Arable soil with high wheat straw input</td>
<td>Closeaux experimental station, France</td>
<td>Wheat Straw</td>
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<td>0.22*</td>
<td>98</td>
<td>0.167</td>
<td>293.15</td>
<td>0.19</td>
<td>209</td>
<td>19.9</td>
<td></td>
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<tr>
<td>Arable soil with low cellulose input</td>
<td>Closeaux experimental station, France</td>
<td>Wheat Straw</td>
<td>0.5</td>
<td>0</td>
<td>0</td>
<td>0.167</td>
<td>293.15</td>
<td>0.19</td>
<td>209</td>
<td>19.9</td>
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<td>Arable soil with low wheat straw input</td>
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<td>0.167</td>
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<td>0.19</td>
<td>209</td>
<td>19.9</td>
<td></td>
</tr>
<tr>
<td>Harmer &amp; Marschner (2005)</td>
<td>Dystric cambisol (A horizon) + Alanine</td>
<td>Steigerwald, Baviaria, Germany</td>
<td>Alanine</td>
<td>13.3</td>
<td>0</td>
<td>3</td>
<td>0.14</td>
<td>293.15</td>
<td>0.6</td>
<td>26</td>
<td>44</td>
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<tr>
<td>Dystric cambisol (A horizon) + Fructose</td>
<td>Steigerwald, Baviaria, Germany</td>
<td>Fructose</td>
<td>13.3</td>
<td>0</td>
<td>0</td>
<td>0.14</td>
<td>293.15</td>
<td>0.6</td>
<td>26</td>
<td>44</td>
<td></td>
</tr>
<tr>
<td>Haplic podzol (EA horizon) + Alanine</td>
<td>Fichtelgebirge, Baviaria, Germany</td>
<td>Alanine</td>
<td>13.3</td>
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<td>0.6</td>
<td>26</td>
<td>32</td>
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</tr>
</tbody>
</table>

*estimated values
Table 2: Model parameters summary for PRIM and the ORCHIDEE soil module

<table>
<thead>
<tr>
<th>Model parameter</th>
<th>Meaning</th>
<th>SOC pools</th>
<th>Prior range</th>
<th>Posterior modes ± s.d. (prior modes) for PRIM</th>
<th>Posterior modes ± s.d. (prior modes) for the ORCHIDEE soil module</th>
</tr>
</thead>
<tbody>
<tr>
<td>( k_{SOC} )</td>
<td>Turnover rate of SOM (d)</td>
<td>Active</td>
<td>10^{-3}-0.5</td>
<td>0.30 ±0.15 (0.31)</td>
<td>0.43 ±0.22 (0.43)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Slow</td>
<td>0.5-5</td>
<td>1.12 ±0.01 (4.51)</td>
<td>0.50 ±0.09 (2.39)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Passive</td>
<td>5-500</td>
<td>462.0 ± 233.8 (467.55)</td>
<td>40.17 ± 22.19 (44.39)</td>
</tr>
<tr>
<td>( c )</td>
<td>Influence of the FOM carbon pool in the SOM mineralization (priming parameter)</td>
<td>Active</td>
<td>2.10^{-4}-500</td>
<td>493.7 ± 246.8 (493.7)</td>
<td>NA</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Slow</td>
<td>2.10^{-4}-500</td>
<td>194.0 ± 97.0 (194.0)</td>
<td>NA</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Passive</td>
<td>2.10^{-4}-500</td>
<td>136.5 ± 68.3 (136.5)</td>
<td>NA</td>
</tr>
<tr>
<td>Study</td>
<td>Treatments performed</td>
<td>Ecosystems</td>
<td>Sites Names (Coordinates)</td>
<td>Treatment performed in:</td>
<td>CO₂ monitored between:</td>
</tr>
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<td>---------------------------</td>
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</tr>
<tr>
<td>Borken et al. (2002)</td>
<td>Compost amendment/ Control</td>
<td>Needleleaf forest</td>
<td>Solling, Norway (51°46’N, 9°34’E)</td>
<td>August 1997</td>
<td>September 1997-December 1999</td>
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*estimated values
Table 4: Model performances for each evaluation sites

<table>
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</thead>
<tbody>
<tr>
<td></td>
<td>All data</td>
<td>No litter</td>
<td>Control</td>
<td>Double litter</td>
<td>All data</td>
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<tr>
<td><strong>ORCHIDEE</strong></td>
<td>slope</td>
<td>0.56</td>
<td>0.45</td>
<td>0.65</td>
<td>0.66</td>
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<tr>
<td></td>
<td>NSD</td>
<td>1.43</td>
<td>1.86</td>
<td>1.37</td>
<td>1.48</td>
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<td></td>
<td>BIC</td>
<td>103.4</td>
<td>57.9</td>
<td>49.3</td>
<td>53.8</td>
</tr>
<tr>
<td><strong>ORCHIDEE-PRIM</strong></td>
<td>slope</td>
<td>0.55</td>
<td>0.45</td>
<td>0.65</td>
<td>0.61</td>
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<tr>
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<td>NSD</td>
<td>1.53</td>
<td>1.85</td>
<td>1.37</td>
<td>1.59</td>
</tr>
<tr>
<td></td>
<td>BIC</td>
<td>116.3</td>
<td>64.9</td>
<td>56.5</td>
<td>63.46</td>
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</table>
Table 5: Correlation between optimized parameters

<table>
<thead>
<tr>
<th></th>
<th>k_{soc}</th>
<th></th>
<th></th>
<th>c</th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Active</td>
<td>Slow</td>
<td>Passive</td>
<td>Active</td>
<td>Slow</td>
<td>Passive</td>
</tr>
<tr>
<td>k_{soc}</td>
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<td></td>
<td></td>
</tr>
<tr>
<td>Active</td>
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<td>0.00</td>
<td>0.00</td>
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