Causal networks of biosphere–atmosphere interactions

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Abstract. Local meteorological conditions and biospheric activity are tightly coupled. Understanding these links is an essential prerequisite for predicting the Earth system under climate change conditions. However, many empirical studies on the interaction between the biosphere and the atmosphere are based on correlative approaches that are not able to deduce causal paths, and only very few studies apply causal discovery methods. Here, we use a recently proposed causal graph discovery algorithm, which aims to reconstruct the causal dependency structure underlying a set of time series. We explore the potential of this method to infer temporal dependencies in biosphere-atmosphere interactions. Specifically we address the following questions: How do periodicity and heteroscedasticity influence causal detection rates, i.e. the detection of existing and non-existing links? How consistent are results for noise-contaminated data? Do results exhibit an increased information content that justifies the use of this causal-inference method? We explore the first question using artificial time series with well known dependencies that mimic real-world biosphere-atmosphere interactions. The two remaining questions are addressed jointly in two case studies utilizing observational data. Firstly, we analyse three replicated eddy covariance datasets from a Mediterranean ecosystem at half hourly time resolution allowing us to understand the impact of measurement uncertainties. Secondly, we analyse global NDVI time series (GIMMS 3g) along with gridded climate data to study large-scale climatic drivers of vegetation greenness. Overall, the results confirm the capacity of the causal discovery method to extract time-lagged linear dependencies under realistic settings. The violation of the method’s assumptions increases the likelihood to detect false links. Nevertheless, we consistently identify interaction patterns in observational data. Our findings suggest that estimating a directed biosphere-atmosphere network at the ecosystem level can offer novel possibilities to unravel complex multi-directional interactions. Other than classical correlative approaches, our findings are constrained to a few meaningful set of relations which can be powerful insights for the evaluation of terrestrial ecosystem models.
1 Introduction

Understanding biosphere-atmosphere interactions is a prerequisite to accurately quantify feedbacks in the Earth system (Bonacci, 2015). On the one hand, the biosphere responds to the atmospheric drivers such as radiation intensity, temperature, vapour pressure deficit, and composition of trace gases. On the other hand, the biosphere influences the atmosphere via partitioning the incoming net radiation into sensible, latent, and ground heat fluxes as well as via controlling the exchange of trace gases and volatile organic compounds. Over the past decades, many of these processes have been identified and their physical, chemical and biological effects have been investigated (see e.g. Monson and Baldocchi, 2014; McPherson, 2007, for overviews). However, the synergistic effects of these processes and the specific cause-effect interactions underlying these effects are still not fully understood (Baldocchi et al., 2016; Miralles et al., 2018).

Today, there is a manifold of monitoring systems operating at various spatial and temporal scales. Networks of eddy covariance towers offer high temporal resolution to study the dependence between carbon, water, and energy fluxes and the atmosphere under a variety of normal and stress conditions or in response to disturbances (Baldocchi, 2014). Satellite remote sensing data complement this picture. They typically only monitor vegetation states at multi-day resolutions and some products offer nearly complete global coverages (Justice et al., 2002; Woodcock et al., 2008). The actual and future satellite missions are leading to rapid development in the field with ever higher spatial, temporal, and spectral measurements (Malenovský et al., 2012; Guanter et al., 2015; Qi and Dubayah, 2016).

The study of biosphere-atmosphere interactions using observations typically relies on correlative approaches, or is based on model-data i.e. requires a-priori knowledge. Only few attempts have been made to learn directional dependencies in a data-driven manner in biogeosciences. For instance, Ruddell and Kumar (2009) used transfer entropy, a bivariate information theoretic measure, to estimate networks of information flow. These networks constructed for a corn-soybean ecosystem under drought and normal conditions showed substantial differences in connectivity. The decoupling during drought between variables describing both land and atmospheric conditions was attributed to changes in the feedback patterns for the two conditions. In recent years, a new branch in statistics aiming for causal inference from empirical data has experienced substantial progress. The idea of causal inference emerged already in the early 20th century (Wright, 1921). Later, Granger suggested one of the first applicable formalisms (Granger, 1969); since then, several efforts in ecology and climate science have concentrated on the bivariate form of Granger causality (Elsner, 2006, 2007; Kodra et al., 2011; Attanasio, 2012; Attanasio et al., 2012).

Aiming to mitigate some of the limitations of the traditional Granger causality, Detto et al. (2012) used a conditional spectral Granger causality framework that allows to disentangle system inherent periodic couplings from external forcing. The disentanglement is enabled via decomposition into the frequency domain using wavelet theory. This method enabled the finding that soil respiration in a pine and hardwood forested ecosystem in winter is not influenced by canopy assimilation but only by temperature, a result that would not be detectable via lagged correlation or bivariate Granger causality. Green et al. (2017) used a similar approach to investigate biosphere-atmosphere feedback loops. It is found that they can explain up to 30% of variance in radiation and precipitation in certain regions. Recently, Papagiannopoulou et al. (2017a) applied a non-linear multivariate conditional Granger causality framework to study climatic drivers of vegetation at the global scale. This approach revealed...
that water limitations on plant productivity play a considerably larger role than previously expected (Papagiannopoulos et al., 2017b). These examples show that unexpected interactions patterns can be found from data only and may challenge our theoretical assumptions. In fact, in the last years the science of causal inference has developed a strong theoretical foundation and several algorithms have been proposed (Spirtes et al., 2001; Pearl, 2009; Peters et al., 2017; Runge et al., 2019). However, we only find few studies that test the latest generation of such methods in ecosystem ecology (see e.g. Shadaydeh et al., 2018; Christiansen and Peters, 2018).

Ecological and climate data is usually time ordered. Using the concept of time series graphs (Ebert-Uphoff and Deng, 2012), the time order can be exploited to create efficient causal graph discovery algorithms (Runge, 2018a). Recently, Runge et al. (2018) introduced such an algorithm, called PCMCI, a combination of the PC algorithm (named after its inventors Peter and Clark, Spirtes and Glymour, 1991) and the Momentary Conditional Independence (MCI) test. PCMCI has been applied to artificial tests (Runge et al., 2018) and climatological case studies (Runge et al., 2014; Kretschmer et al., 2016). Hence, this method could be potentially of very high relevance for learning the causal dependency structure in the complex biosphere-atmosphere system.

In this study, we explore the potential of PCMCI for disentangling and quantifying interactions and feedbacks between terrestrial biosphere state and fluxes and meteorological variables. The study is structured as follows: Firstly, we describe the performance of the method on artificial time series data with well-known dependencies that mimic some basic properties of observed land surface fluxes such as heteroscedasticity (Sect.3.1). Secondly, we explore three replicated eddy covariance measurement towers in a Mediterranean ecosystem and explore how the identified interdependencies of carbon and energy fluxes and micro-meteorological observations vary over time (Sect.3.2). Thirdly, we use global satellite data of vegetation greenness to understand the lagged dependency of ecosystems with respect to climatic drivers (Sect.3.3). Based on these results, we discuss the potentials and limitations of PCMCI for other applications in land-atmosphere studies and give recommendations for further methodological developments (Sect.4).
2 Method and Data

2.1 Causal Discovery via PCMCI

PCMCI assesses the causal structure of a multivariate dataset or process $X$ by estimating its time series graph. In short, the nodes of a time series graph $G$ represent single variables $X^i, X^2, ..., \in X$ at specific (lagged) time steps $t, t-1, ..., t-\tau_{max}$. A lag is the number of time steps $\tau$ between two nodes $X^i_t$ and $X^j_{t-\tau}$, i.e. $\tau = t_1 - t_2$ for $t_1 > t_2$. To generate a causal graph from observational data through statistical dependencies, a series of assumptions must be adopted. Here, we assume: time order, the causal Markov condition, faithfulness, causal sufficiency, and no contemporaneous causal effects (Runge et al., 2018). PCMCI is applied in combination with the ParCorr linear independence test based on partial correlations (cf. Sect. 2.1.1). This application additionally requires stationarity in mean and variance and linear dependencies. In the following, we briefly discuss these assumptions (further details in Runge, 2018a) and refer to a variable $X^i$ that is causally affecting a variable $X^j$ as ‘parent’ or ‘driver’ and the latter as ‘receiver’ or ‘target’.

The time order within the time series allows to orient directed links which are only pointing forward in time. This accounts for causal information propagating forward in time only, i.e. the cause shall precede the effect. Therefore, a directed causal link $X^i_t \rightarrow X^j_{t-\tau}$ can only exist between two nodes $X^i_t, X^j_{t-\tau}$ if $t_1 < t_2$. When a contemporaneous link is found, i.e. $t_1 = t_2$, it is considered to be undirected. The connection between the graph $G$ and its process $X$ is given by the causal Markov condition together with the faithfulness. The Markov condition says that if there is no link in the graph, there is conditional independence in the distribution, and if there is no conditional independence, there is connectedness in the graph; the reverse relations, if there is conditional independence, there is also separation in the graph and if there is connectedness in the graph, there is no conditional independence, are given by the faithfulness assumption. With these assumptions the graph is a visualisation of the conditional dependence and independence relationships among the variables including their lags. Finally, the causal sufficiency assumption implies that, every common cause of two or more variables $X^i \in X$ is included in $X$. If this is not the case, detected links may be indirect or due to an unobserved common driver. However, the absence of a link in the detected graph still implies that no direct link is present (as this only requires the assumption of faithfulness).

Several causality methods suffer from high dimensionality, i.e. multivariate transfer entropy (Runge et al., 2012). They use the whole past $X^{-}_t \backslash X^{-}_{t-\tau}$ of $X^i_t$ with $X^i_t = (X^i_{t-1}, X^i_{t-2}, ...)$ as a conditioning set $S$ in the estimation of the test statistic $I(X^{-}_t \backslash X^{-}_{t-\tau} \mid X^i_t) : S = X^i_t \backslash X^{-}_{t-\tau}$. PCMCI addresses this issue by reducing the set of conditions $S$. PCMCI first estimates $S$ for each $X^i_t$. The estimation removes all variables that are either independent or dependent due to indirect paths or common drivers only and leaves only relevant conditions converging to the true causal parents in the limit of infinite sample size. This is done using a variant of the PC algorithm (Spirtes et al., 2001). The conditioning set is subsequently used in the MCI step to test every possible link; MCI also attributes a link strength. These two steps as well as the concept of independence tests, which are at the core of both PC and MCI algorithms, are explained in more detail in the following.
2.1.1 Independence Test

At the core of PCMCI, there are conditional independence tests \( CI(X_i^{t-\tau}, X_j^t, S) \) testing for \( X_i^{t-\tau} \perp X_j^t \mid S \). Within the PCMCI software package Tigrumite (Runge, 2018b), several independence tests are implemented. Here, we focus on the linear independence test called ParCorr. The ParCorr conditional independence test is based on partial correlations and a t-test. This assumes the model

\[
X = S\beta_x + \epsilon_x, \quad Y = S\beta_y + \epsilon_y,
\]

with coefficients \( \beta \) and Gaussian noise \( \epsilon \). This leads to the residuals

\[
r^x = X - S\hat{\beta}_x, \quad r^y = Y - S\hat{\beta}_y
\]

with estimated \( \hat{\beta} \). ParCorr removes the influence of \( S \) on \( X \) and \( Y \) via ordinary least squares regression and tests for independence of the residuals using the Pearson correlation with a t-test. The independence test returns a p-value and test statistic value \( I \), i.e. the correlation coefficient in case of ParCorr.

2.1.2 PC algorithm

The PC algorithm estimates a set of parents for each variable of the process \( X_i^t \in \mathbf{X} \), which are used as low-dimensional conditions in the MCI algorithm. A comprehensive pseudo code of this procedure is given in Runge et al. (2018). The algorithm starts with a fully connected graph and iteratively removes links if conditional independence is found. At first, a preliminary set of parents \( \tilde{\mathcal{P}} \) for each \( X_i^t \) equal to the whole past of the process \( \mathbf{X} \), is initialised: \( \tilde{\mathcal{P}}(X_i^t) = \mathbf{X}_i^t = \{X_i^{t-\tau} : i = 1, \ldots, N, \tau = 1, \ldots, \tau_{\max}\} \). Next, conditional independence \( X_i^{t-\tau} \perp X_j^t \mid S \) is tested for all \( X_i^{t-\tau} \in \tilde{\mathcal{P}} \cdot S \) is a subset of \( \tilde{\mathcal{P}}(X_i^t) \setminus \{X_i^{t-\tau}\} \). The conditional independence test is embedded into two loops. The outer loop iteratively increases the cardinality \( p \) of \( S \) starting from 0, the empty set. Thus it is first tested, whether \( X^i \) and \( X^j \) are independent. Further, only the p strongest conditions are selected. The inner loop iterates through all \( X_i^{t-\tau} \in \tilde{\mathcal{P}}(X_i^t) \). The cardinality is increased until \( |S| = |\tilde{\mathcal{P}}(X_i^t) \setminus \{X_i^{t-\tau}\}| - 1 \). At the end of each iteration in the outer loop the non-significant parents are removed from \( \tilde{\mathcal{P}}(X_i^t) \). Every conditional independence test is evaluated at a significance threshold called \( \alpha_{pc} \), which is usually set to a liberal value between 0.1 and 0.4. Alternatively, one can let \( \alpha_{pc} \) unspecified. PCMCI then evaluates the best choice of \( \alpha_{pc} \in \{0.1, 0.2, 0.3, 0.4\} \) based on the Akaike information criterion which is further explained in (Runge et al., 2018).

2.1.3 MCI tests

MCI is the actual causal discovery step that ascribes a p-value and strength to each possible link. MCI iterates through all pairs \( \{X_i^{t-\tau}, X_j^t\} : i = 1, \ldots, N, \tau = 0, \ldots, \tau_{\max} \) and calculates \( CI(X_i^{t-\tau}, X_j^t, S) \). \( S \) consists of the two potentially multivariate sets of parents \( \tilde{\mathcal{P}}(X_i^t) \) and \( \tilde{\mathcal{P}}(X_i^{t-\tau}) \) obtained in the PC step. \( \tilde{\mathcal{P}}(X_i^{t-\tau}) \) is constructed by shifting the time series of \( \tilde{\mathcal{P}}(X_i^t) \) by \( \tau \). In case \( X_i^{t-\tau} \in \tilde{\mathcal{P}}(X_i^t) \), \( X_i^{t-\tau} \) has to be removed from \( \tilde{\mathcal{P}}(X_i^t) \). If \( \tau = 0 \), conditional dependence is estimated for contemporaneous nodes \( X_i^t \) and \( X_j^t \). Due to missing time order, a dependence would be left undirected. Further, as the parents \( \tilde{\mathcal{P}}(X_i^t) \) and \( \tilde{\mathcal{P}}(X_i^{t-\tau}) \) used in
each conditional dependence test are defined to lie in the past of $X^t_i$ and $X^t_j$, links, both contemporaneous and lagged, can be spurious due to contemporaneous common drivers or contemporaneous indirect paths. The absence of a link, though, means that a physical (contemporaneous) link is unlikely (assuming faithfulness, cf. Runge et al. (2018)).

The link strength is given by the effect size of the conditional independence test statistic measure CI used in combination with MCI. In case of ParCorr, the effect size is given by the partial correlation value, which is between -1 and 1. This value is shown in Runge et al. (2014) to directly depend on the receiver’s and driver’s variance as well as the coupling coefficient. Even non-linear links can be detected with ParCorr as they can often be linearly approximated. In case the linear part is even stronger than the non-linear part, ParCorr can have better detection rate than a non-linear independence test (Runge et al., 2018).
2.2 Data

2.2.1 Artificial Dataset - Test Model

We tested the algorithm on artificial datasets prior to its application to real world data. Using time series of measured global radiation ($R_G$) we created three artificial time series that conceptually represent temperature ($T$), gross primary production ($GPP$) and ecosystem respiration ($Reco$). Note that this test model is not intended to accurately represent observed land-atmosphere fluxes, but only serves to test the procedure. The model incorporates one linear auto dependence $T_{t-\tau_1} \rightarrow T_t$, one linear additive cross-dependence $R_G\rightarrow T_t$ and two non-linear dependencies, multiplicative $R_G\cdot T_{t-\tau_4}$ $\rightarrow$ $GPP_t$, and multiplicative exponential $GPP_{t-\tau_5} \cdot e^{T_{t-\tau_6}} \rightarrow$ $Reco_t$ (cf. Fig. 1) according to the equations:

$$R_{G_{mo}} = R_{G_{obs}}$$

$$T_{mo}(t) = c_1 \ T_{mo}(t-\tau_1) + c_2 \ R_{G_{mo}}(t-\tau_2) + \xi_T$$

$$GPP_{mo}(t) = c_3 \ R_{G_{mo}}(t-\tau_3) \cdot T_{mo}(t-\tau_4) + \xi_{GPP}$$

$$Reco_{mo}(t) = c_4 \ GPP_{mo}(t-\tau_5) \cdot e^{T_{t-\tau_6}} + \xi_{Reco}$$

The parameters $c_1, c_2, ..., c_5$ are referred to as coupling coefficients, and the time lags are noted as $\tau_1, \tau_2, ..., \tau_5$. The subscripts $mo$ and $obs$ abbreviate model and observation, respectively. $T_{ref}$ is set to 15°C. The term $\xi$, termed “intrinsic” or “dynamical noise”, here represents values from uncorrelated, normally distributed noise. Having dynamic noise is essential for a method utilizing conditional independence tests. It is based on the assumption that a process or state is never fully described by its deterministic part because there are unresolved intrinsic processes, summarized as $\xi$.

The model was fitted to real observational data (using radiation, temperature and land-atmosphere fluxes) of daily time resolution, measured by eddy-covariance method (Baldocchi et al., 1988; Baldocchi, 2003) from FLUXNET, by minimizing the sum of squared residuals using the gradient descent implemented in the Optim.jl package (Mogensen and Riseth, 2018).

We fitted the model to 72 sites listed in Table B1 given in the Supplementary Material section. The value range for the coupling coefficients $c_1$ to $c_4$ and $c_5$ were set to $[0.2,1]$ and $[1,2.5]$, respectively. The lags were limited to integer values in the range $[0,25]$. For the fitting, the dynamical noise $\xi$ was set to 0. The distributions of obtained lags and coupling coefficients are given in the Supplementary Material Fig. A1 to D1. The fitting process thus generated 72 sets of parameters, containing coupling coefficients and lag values, which were used for the time series generation.

From each of the 72 sets of parameters we generated four sets of time series each having a length of 500 years. The time series generation was initiated using two types of data: first, uncorrelated, normally distributed noise, and second, unprocessed radiation data as used during the fitting (the available radiation data was repeated to 500 years). The resulting datasets are called baseline dataset and seasonality dataset, respectively. In both cases, the model was run twice, once with homoscedastic (constant variance of $\xi$), once with heteroscedastic dynamical noise $\xi$. To induce heteroscedasticity, $\xi$ was multiplied with a mean daily variance that was extracted for each variable at each FLUXNET site. In Supplementary Material Fig. E1 a five year
2.2.2 Eddy Covariance Data - Majadas de Tiétar Experimental site

Data from three towers located in Majadas de Tiétar, (ES-LMa, ES-LM1, ES-LM2), a Mediterranean Savanna in central Spain, are used (coordinates of central tower: 39°56'25''N 5°46'29''W). Measurements include the exchange of CO₂ between the land surface and atmosphere at half-hourly resolution using the eddy covariance method. The three tower footprints received different fertilisation treatments in spring 2015 (El-Madany et al., 2018). We consider data from before the fertilization from April 2014 to March 2015 of shortwave downward radiation (Rg), air temperature (T), net ecosystem exchange (NEE), vapour pressure deficit (VPD), sensible heat (H) and latent heat (LE). The average temperature within this period was 17.3°C, with a total precipitation of 765mm. Most precipitation fell between October and April.

We expect the causal imprints in the data to vary between seasons and during the course of the day. To satisfy causal stationarity, we estimate networks separately for each month and consider only samples for which the potential radiation was above 4/7 of the potential daily maximum, which corresponds to midday samples. We used a mask type that limits only the receiver variable to the respective month and day time values (cf. Table A1 for PCMCI parameter settings). This setting causes time series lengths ranging from 239 datapoints in December to 372 datapoints in July. Minimal and maximal lags were set to 0 and 8, respectively. Due to the limited number of years, we left the data unprocessed, i.e. we did not subtract a seasonal mean for anomalisation. Constraining the samples to separate month and midday values reduces the effect of seasonality as a common driver that would lead to spurious links. Furthermore, to correct for multiple testing we applied the Benjamini-Hochberg false discovery rate correction (Benjamini and Hochberg, 1995). Thereby, the p-values for the whole graph obtained from the MCI step are adjusted to control the number of false discoveries (Runge et al., 2018). We chose a two-sided significance level of 0.01.

2.2.3 Gridded global data set

The second observational case study was performed on a global data set. We used data with 0.5° spatial and monthly temporal resolution from 1982 to 2008. The dataset is composed of three climatic variables, global radiation (Rg), temperature (T) and precipitation (P), and one vegetation state index, the Normalized Difference Vegetation Index (NDVI). Both temperature and precipitation datasets were taken from the Climate Research Unit (CRU), version TS3.10 (Harris et al., 2014). The radiation data stems from the Climate Research Unit and National Centers for Environmental Prediction dataset (CRUNCEP, Viovy, 2016). The used NDVI data stems from the Global Inventory Monitoring and Modeling Systems (GIMMS) in version 3g_v1 (Pinzon and Tucker, 2014).

To examine the influence of radiation, temperature, and precipitation on NDVI by means of PCMCI we used the following settings. We compute the anomalies by subtracting a smoothed seasonal mean. A maximal time lag of three months was chosen based on the largest lag with significant partial correlation among all pairs of variables, partialling out only the autocorrelation of each variable. The receiver variable was limited to the growing season defined by T>0 and NDVI>0.2, which allows good
comparison to Wu et al. (2015). The significance level ($\alpha_{pc}$) in the condition selection phase (cf. Sect. 2.1.2) was chosen based on the AIC selection criterion. A concise list of PCMCI parameters that were altered from default settings is given in Table A1.
Figure 1. The artificial datasets are generated with a prescribed interaction structure (True Network), which is obtained by fitting the test model to the FLUXNET sites. Here we show for four time series length the process graphs estimated via both lagged correlation and PCMCI. The data used stems from the homoscedastic realisation of the seasonality dataset of the Hainich site. The significance level was set to 0.01. The number of time lag labels were limited to five in the correlation networks. But for the longest time series typically the whole range of lags (0-25) was significant.

3 Results

3.1 Test Model

As motivating example, in Fig. 1 we show PCMCI and lagged correlation networks in the form of process graphs. It is clearly visible, that many more spurious links pass the significance threshold of 0.01 using lagged correlation as compared to using PCMCI. Those spurious links can complicate the analysis or lead to false assumptions and misleading hypotheses. We examined four cases of different time series lengths: 91 [183<=doy<=274] and 120 [153<=doy<=274] days, 1 and 5 years for daily data (doy: day of the year). For each time series length and each parameter set, the causal network structure was estimated for 100 realisations of the model (each based on a realization of intrinsic noise), which allowed the estimation of false positive (FPR) and true positive (TPR) detection rates. The detection rates are calculated for each tower, FPR in general and TPR link-wise. The TPR for each link is its sum of detections among 100 realisations divided by 100. The FPR is the number of falsely detected links divided by the number of all possible false links and 100. The summary of the experiments i.e. the overall false positive rate (FPR) and the distributions of the link’s true positive rate (TPR) across sites are given in Fig. 2 and 3, respectively. The blue violin plots always report the case of normal distributed (non heteroscedastic) intrinsic noise and the corresponding orange violin plots summarize the case of heteroscedastic noise. The effect of heteroscedasticity and seasonality is then assessed by comparing the distributions obtained from the baseline dataset to the results of the seasonality dataset.

The FPR of homoscedastic time series in the baseline dataset is in the expected range of 0.01, the chosen significance level, indicating a well calibrated test due to fulfilled assumptions. The assumption of stationarity is violated as soon as heteroscedastic-
Figure 2. The distribution of false positive detection rates estimated for the baseline dataset, the seasonality dataset and the anomalised seasonality dataset (mean seasonal cycle subtracted). The distributions are given for different time series length (number of datapoints). Additionally, the distributions are split to show the impact of heteroscedastic noise (orange) compared to normal distributed noise (blue). The significance level of 0.01 is given by a blue horizontal line.

ticity or seasonality is present. The effect on the FPR is an increase above 0.01 for time series length of 1 and 5 years with a much stronger increase due to seasonality (factor of 4 and 8, respectively) than for heteroscedasticity (factor of 2).

The effect of non-stationarities on the TPR differs among the links. The detection of linear links ($R_g \rightarrow T$ and $T \rightarrow T$) is not affected by seasonality and slightly improves for heteroscedastic dynamical noise. The detection of non-linear links is improved by seasonality with the strongest effects in the link $T \rightarrow GPP$. The link $T \rightarrow Reco$ has a stronger non-linearity and therefore the detection rate shows a weaker effect on seasonality. Furthermore, the coupling coefficient $c_5$, the base of $T$, can be close to or be exactly one (cf. Fig. B1). This would actually cause on the one hand the effect of $T \rightarrow Reco$ to vanish, rendering a detection impossible and on the other hand result in a linear dependence of $GPP$ on $Reco$ which improves its detection. Heteroscedasticity seems to have a slight negative effect on non-linear links. In general, the TPRs in the seasonality dataset are quite high, even for non-linear links, and predominantly above 80% and often reaching 100%.

Comparing the TPRs of the non-linear links, shows some disparity. The links $T \rightarrow GPP$ and $T \rightarrow Reco$ experience zero detection in the baseline dataset but partially considerable rates in the seasonality dataset with a strong dependence on the time series length. On the contrary, the median of the TPR of the links $R_g \rightarrow GPP$ and $GPP \rightarrow Reco$ is above 95% in the seasonality dataset, even for time series length as short as 91 days, but remains high in the baseline dataset. The removal of the seasonal cycle keeps the TPRs largely unaffected, but reduces the FPR. Nevertheless, it still remains above the significance level by a factor of four and two for five and one year time series length, respectively.
Figure 3. The distribution of the true positive detection rates for each link in the test model estimated for the baseline dataset, the seasonality dataset, and the anomalised seasonality dataset. The distributions are given for different time series length (number of datapoints). Additionally the distributions are split to show the impact of heteroscedastic noise (orange) compared to normally distributed noise (blue).

In summary, the seasonality dataset exhibits high TPR even for non-linear links. Compared to stationary time series, the detection of non-linear links actually benefits from seasonality. The high detection, though, comes at the cost of a high false positive rate for time series length of and above one year. To a certain degree, the increase in FPR can be counteracted by anomalization.

3.2 Majadas de Tiètar dataset

At first we look at the link consistency by comparing networks that were obtained for each tower within a month. The comparison is done for two months with strongly differing climate conditions: April and August. In Fig. 4 we compare the estimated link strengths (effect size estimated via partial correlation) as long as the corresponding links are significant in at least one network. The confidence intervals are overlapping for the majority of links, suggesting that the uncertainty of the fluxes is much smaller than the observed effects (El-Madany et al., 2018). Exceptions are found for only a few links (Rg→T, Rg→LE, VPD→VPD, H→H, NEE→LE, H→NEE, NEE→NEE, number above the arrow indicates the lag) where the detection rates do not or barely overlap. Cross links (a link from one variable to another) with two or more significant appearances are predomi-
Figure 4. Comparison of the networks of three eddy covariance measurement stations (LMa, LM1, LM2) located in Majadas (Spain). Links that are found to be significant in one of the three networks are included. For each link, the calculated strength of all three networks is plotted with its 90% confidence interval. The colors blue, orange, and green correspond to the towers LMa, LM1, and LM2, respectively. The significance threshold is 0.01. If a link does not pass the significance, it is marked by a black dot. The links are grouped into lag 0 (top), lag 1 (middle) and all lags greater than 1 (bottom). Negative NEE is associated with carbon uptake by the ecosystem. Links at lag 0 are left undirected (−), yet as Rg is set as main driver, links incorporating Rg at lag 0 are directed (→).
nanty at zero lag. Approximately half of the links with lag one are auto-dependent links (a link from the past of a variable to its present). Comparing the links between the month April and August, distinct differences can be noticed. First, August has slightly fewer significant links compared to April. Second, the only links remaining that are significant in two or three towers are between atmospheric variables. Third, the remaining link strengths tend to be weaker in August than in April.

The difference among the seasons is further investigated in Fig. 5 which shows process graphs for each month of the year. We combine the networks of the three towers to one process graph by plotting every link that is significant in at least one tower. The process graphs in Fig. 5 visualize clearly gradual changes within the interaction structure of the biosphere-atmosphere system during the course of a year. During the main growing season from February to May, NEE is coupled strongly to the energy fluxes latent (LE) and sensible heat (H). These connections weaken, disappear or even switch sign with start and course of the dry season. Less regularly NEE also shows connections to radiation (Rg) and temperature (T). Between the atmospheric variables, a basic network between VPD, T, Rg and H remains intact and relatively constant in strength. The dominance of contemporaneous links is found as well, as seen already in Fig. 4. Besides the decoupling of NEE from any variable in the dry period, there are additional interesting patterns. For example, the positive reappearance of the link between NEE and LE in September. Here, the onset of precipitation events (cf. Fig. G1) occurred that lead to strong respiration peaks (Ma et al., 2012). Creating such a network via lagged correlation would result in much more significant links (causing the network to be not interpretable as opposed to PCMCI) and NEE does not decouple from the atmosphere in August (cf. Fig. F1).

The above results demonstrate that PCMCI is sensitive enough to capture seasonal differences and certain physiological reasonable biosphere behaviour. Moreover, PCMCI yields a better interpretable network structure than pure correlation approaches.

3.3 Global

Subject of inspection were the significant lags and MCI values of each climatic variable on NDVI. In Fig. 6 the maximal MCI value and the corresponding lag are plotted for the links $X_{t-\tau} \rightarrow \text{NDVI}$ : $\tau \in \{0, 1, 2, 3\}$ with X being one of the climatic drivers radiation, temperature, or precipitation. The chosen significance threshold was set to 0.05. Fig. 7 shows the climatic driver with largest MCI per grid point. PCMCI detects a regionally varying influence of climatic drivers. As expected, the boreal regions are strongly driven by temperature instantaneously, while (semi-) arid regions, which correspond predominantly to grass or prairie dominated areas, respond strongest to precipitation at a time lag of one month. Radiation is found to have a comparatively low spatial effect with hot spots in south and east China, central Russia and east Canada.

The dominant lags are found to be zero and one. Just a very small fraction of the total area shows a maximal MCI value at a higher lag of two or three months. The lags are also not equally distributed among the climatic drivers. Radiation and temperature are predominantly strongest at lag zero, while precipitation has a much larger fraction of area showing the strongest response at lag one. Regions where the impact of Rg on NDVI is strongest at lag 1 tend to respond negatively to Rg but positively to precipitation at lag one. On the other hand, a large part of regions with the strongest impact of precipitation at lag zero respond negatively to it but positively to radiation.
Figure 5. To visualize the gradual changes in interaction structure the networks of the three towers are combined for each month. The number of significant occurrences of a link is given by its width. The link strength, given by the link color, is calculated by averaging the significant links of the towers. The link’s lag is shown in the centre of each arrow, sorted in descending order of link strength. The resulting graphs are shown for April 2014 till March 2015. The significance threshold is 0.01. The networks of April and August, illustrated in Fig. 4, are highlighted by a box.
Figure 6. Influence of climatic drivers on NDVI as calculated by PCMCI. The first column shows the estimated causal influence given as maximal absolute MCI value of climatic drivers on NDVI. The second column gives the time lag at which the maximal absolute MCI value occurs (in month).

Figure 7. Map of the strongest climatic driver (largest absolute MCI value) per grid point.
In summary, PCMCI estimates coherent interaction patterns which match well with anticipated behaviour based on vegetation type and prevailing climatic conditions.
4 Discussion

Causal discovery methods promise an improved understanding and can help to come up with new hypotheses about the interaction between biosphere and atmosphere (Christiansen and Peters, 2018; Runge et al., 2019). But the underlying assumptions need to be properly taken into account. The coupled biosphere-atmosphere system possesses several challenges that potentially violate the underlying assumptions of causal discovery in general and the employed method’s assumptions in particular. Here, we investigate the effect of a violation of assumptions on PCMCI network estimates.

With regard to expected non-linearities in biosphere-atmosphere interactions, using a linear independence test within the PCMCI framework may not be adequate. We motivate our choice with the following arguments: first, non-linearities are often approximated linearly. Second, a linear regression based test has a much higher power for detecting linear links than a non-parametric test (Runge, 2018a) and can, hence, detect links already at smaller sample sizes. Third, linear partial correlation is easily interpretable, for example, positive and negative MCI values. The results of the test model support our choices. In general, the derived results show a high detection power with a strong consistency in calculated effect strengths on eddy covariance data and global, regularly gridded reanalysis data that leads to well interpretable patterns. Observed drawbacks are a high FPR in case of violated assumptions, especially strong periodicity, as well as the appearance of contemporaneous lags in measurement datasets.

4.1 Lessons learned from the test model

The probability to detect a link with PCMCI depends strongly on a link’s MCI effect size, which is larger for strong variance in the driver and a low variance in the receiver (Runge et al., 2018). Several results can be explained by this observation. First, the variance of three out of five drivers of cross dependencies in the test model are either directly or indirectly (via GPP) influenced by $R_g$, which has the highest variance of all variables. Consequently, the detection power of the three links is large, almost 100%. In comparison, the other variables’ variances are weaker, since they are influenced by $T$, which results in a lower detection power. This is the origin of the disparity in detection rates of the non-linear links. Second, also the partially strong increase in TPR of non-linear links (influenced in a multiplicative way by $T$) from the baseline dataset to the seasonality dataset can be explained by this increase in variance. A multiplicative link is actually not generally expected to be found by ParCorr (Runge, 2018a), but the value of the multiplicative factor is dominated by the seasonal value, and not the dynamical noise, which might cause rather a scaling of the dynamical noise terms rather than a random distortion. Third, the dependence on the variance ratio can also explain the difference in TPR between homoscedastic (equal error variance) and heteroscedastic (error variance changing over time) time series, i.e., the variance of $R_g$ and $GPP$ exhibits a strong seasonality with its peak in summer, while the variance of $T$ is rather constant. This explains, for example, the strong decrease in TPR for the link $T \rightarrow GPP$ at 91 days time series length when comparing homoscedasticity to heteroscedasticity. The decrease in TPR is less pronounced when another season, implying a different variance, is chosen for this comparison. As links with weak driver variance and strong response variance are more likely to be missed, one may ask which effect this will have on the detection...
of feedback loops where one variable has low and the other high variance. Here lies a limitation of the test model where no feedback loops were implemented.

Seasonality and heteroscedasticity constitute violations of the stationarity assumption underlying the independence test ParCorr. As shown in Runge (2018a), including the cause of the non-stationarity as an exogenous driver in the analysis allows PCMCI to regress out its influence on the other variables. However, for ParCorr this is only valid if the dependence on the non-stationary driver is linear. Therefore, the regression on $R_g$ fails for $GPP$ and $Reco$ in the test model. With this ill-posed setting, the probability to detect false links increases with increasing time series length or when more periods are included. Stationarity in mean is obviously also not fully guaranteed when subtracting the seasonal mean. Here we observe that the FPR stays above the significance level for the anomalised seasonality dataset. One can ask whether the FPR stays above the significance threshold because subtracting the seasonal mean does not remove the heteroscedasticity. However, we attribute this high FPR to a not fully removed seasonality since the FPR of both homoscedastic and heteroscedastic time series decreases by roughly the same amount in the anomalised seasonality dataset and the effect of heteroscedasticity is rather weak in the baseline dataset. Much of the influence of heteroscedasticity is also removed when limiting the analysis to a specific time period, i.e. season, which makes the data causally stationary (cf. Sect. 2.1). For example, the link from radiation to GPP vanishes in winter as there is mostly no active plant material left. One can argue, as it is done in Peters et al. (2017), that the causality of a system is invariant even between seasons, the physical mechanism is the same in all seasons. For example, a functional dependence $f(x)$ might be ‘flat’ for small values of $x$ and linearly increasing for larger values. If only small values occur in the winter season, then the link is absent, while it ‘appears’ only in the summer season. Across all seasons, this can be considered as a nonlinear functional dependence $f(x)$. In practice, restricting an analysis to different seasons can help in interpreting the mechanism, here in a linear framework.

Summarizing the results of the test model, the different detection rates, disparity among non-linear links, and the detection of multiplicative links are largely explainable via the effect of the variance on the link detection. Yet, the discussion revealed the need for further research in several aspects. On the one hand, feedback loops are not included in the test model yet are an important aspect in natural systems. On the other hand, removing non-stationarities is essential to keep the false positive rate in the expected range, but standard procedures of subtracting the mean seasonal cycle are not sufficient. Further, the effect of non-stationarity on the causal network structure needs to be investigated.

4.2 Causal interpretation of estimated networks from observational data

In both the half-hourly time resolved eddy covariance data and the monthly global dataset the predominant type of dependence found is contemporaneous. PCMCI leaves these undirected since no time order indicating the flow of causal information is available. Further, as discussed in Sect. 2.1, contemporaneous common drivers or mediators are not accounted for. The consequence is that both spurious contemporaneous and spurious lagged links can appear, if they are due to contemporaneous variables. For interactions that are contemporaneous in nature since they occur on considerably shorter time scales than the time resolution, therefore, PCMCI is not the optimal choice regarding a causal interpretation and other methods should be applied in conjunction (Runge et al., 2019). Further, we faced a trade off between fulfilling causal assumptions and detection power.
In practice, accounting for causal stationarity (by limiting the analysis to certain periods of the dataset) means decreasing the number of available data points while accounting for causal sufficiency leads to an increase in dimensionality by adding variables and increasing the maximal lag. Both will lead to a decrease in detection power, which can affect the network structure. PCMCI alleviates the curse of dimensionality by applying a condition selection step, but still one cannot indefinitely add more variables. Another important factor that affects detection power and dimensionality is the time resolution. There are several points in favour and against increasing time resolution. On the one hand, increasing time resolution can resolve contemporaneous links and potentially increases the detection power due to an increased number of datapoints. On the other hand, the dimensionality increases if the maximal lag is adapted. Further, causal information might be split apart and distributed over more lags, rendering the links at each individual lag less detectable. This can cause links to disappear, but links can also appear if new processes are resolved at a higher time scale. At last, observational noise (measurement errors) might be larger in higher resolution data than in aggregated data, as it is averaged out in the latter and thus affects link detection less. Consequently, when comparing network structures based on different settings, i.e. maximal lag, included variables, time resolution, and considered time period, the (dis-)appearance of single links among specific variables can stem from several factors, i.e. a change in detection power, a changed (conditional) dependency, or due to a common driver. These factors together with a non-zero false positive detection rate are challenging for a causal interpretation. Therefore, detected links should be interpreted with care and can give rise to new hypotheses and analyses involving further variables. Generally, a causal interpretation is more robust regarding the absence of links (cf. Sect.2.1). In particular it does not require that all common drivers are observed.

Nevertheless, robust patterns were identified in our studies that are also consistent with other studies. Furthermore, a causal analysis has the advantage of an enhanced interpretability compared to correlative approaches. First of all, we could show that the networks’ estimated link strengths are consistent for observational data, even though measurement error affects the data. The dataset used was suitable for this analysis, as the measurement stations are located in a reasonably homogeneous ecosystem that shows only little spatial variation (El-Madany et al., 2018). Thus, also the interaction between biosphere and atmosphere is expected to change only marginally across space. Second, the gradual changes in plant activity that are taking place in the ecosystem of Majadas throughout the year do very well emerge in the coupling strength of daytime NEE to the atmospheric variables. The observed decoupling during the dry season is in accordance with the one of a soybean field during drought conditions observed by Ruddell and Kumar (2009). The gradual changes in ecosystem activity are not visible in a pure (lagged) correlation analysis or are only visible in color or density changes but the large number of significant links prevents any detailed interpretation on the physical mechanisms and changes thereof. The large number of significant links compared to the PCMCI networks stems solely from the absence of conditioning on common drivers or mediating variables, which often further leads to an overestimation of the link strength in correlation networks. As a result, processes, such as the decoupling of NEE during the dry period, stay hidden. To reduce the effect of confounding, often analyses utilize partial correlation (see e.g. Buermann et al., 2018). However, a partial correlation can introduce new dependencies (as opposed to removing them) if one conditions on causal effects of the variables under consideration (the ‘marrying parents’ effect). This issue is avoided in PCMCI by only conditioning on past variables. Additionally, PCMCI chooses only relevant variables as conditions by applying
the PC condition selection step which is especially valuable in high dimensional study cases and improves detection power and computation time (Runge, 2018a).

The global study of climatic drivers of vegetation shows a general pattern of lags and dependence strengths of vegetation on climatic variables that is easily-interpretable. The boreal regions appear energy limited and especially driven by temperature (cf. Fig 6c), while the strongest dependence of (semi-)arid regions on precipitation reflects their limitation in water supply. Two recent studies performed a similar analysis. Both Wu et al. (2015) and Papagiannopoulou et al. (2017b) investigated lagged effects and dependence strengths of NDVI on precipitation, temperature and radiation. Wu et al. (2015) estimated the lags of the strongest effects via an univariate regression of the climatic drivers on NDVI and subsequently used those lags to fit a multivariate regression model of the climatic drivers on NDVI and determined their relative effects. Papagiannopoulou et al. (2017b) applied a non-linear Granger causality framework utilising a random forest predictive model; the method was presented by Papagiannopoulou et al. (2017a). We recognize that similar patterns are observed in Wu et al. (2015), but the lags at the maximal MCI value are usually lower than the one found in Wu et al. (2015), which stems from the methodical differences. Besides having used anomaly values, PCMCI regresses both NDVI and the climatic drivers on their parents before calculating the MCI value (cf. Sect. 2.1.3). This especially removes the influence of autocorrelation. Runge et al. (2014) shows how autocorrelation affects the correlative lag causing it to be larger for stronger autocorrelation; thereby the correlative lag may become larger than the causal lag. Therefore, according to Fig. 6b, the causal information embedded in monthly resolution is predominantly received within one month. Finding the strongest causal links at a time lag up to one month appear in agreement with Papagiannopoulou et al. (2017b). Also the spatial distribution of the strongest climatic influences compares well. But there are certain noteworthy differences which not necessarily stem from masking differences, i.e. that we took only values belonging to the growing season while Papagiannopoulou et al. (2017b) took the whole time series. First, there is little significant Granger causality of water availability found in boreal regions while there are significant negative causal dependencies detected via PCMCI. Second, NDVI in arid regions is not or barely Granger caused by radiation and temperature, but in parts shows a negative PCMCI value on those variables. There might be physiological reasons that can explain the PCMCI patterns, i.e. water logging or too high temperatures. To explain the differences though, we could identify two possible reasons. First, Papagiannopoulou et al. (2017b) masked out negative influences of radiation arguing that radiation is not negatively affecting NDVI. They found, that negative influences of Rg are usually a consequence of poor conditioning on other variables. Second, a precipitation event in boreal regions coincides with a reduction in radiation and temperature. Boreal regions usually do not suffer from water shortages. Thus they respond stronger to the reduction of radiation and temperature than precipitation. As precipitation is coupled negatively to radiation and temperature at lag zero, the effect of precipitation on NDVI is found to be negative. Thus, the link P → NDVI might be an effect of the contemporaneous common driver scheme P ← Rg → NDVI and therefore would not be causal. In fact, a similar argumentation can be given for the negative impact of temperature and radiation on NDVI in arid regions.

In summary, we pointed out the need for careful interpretations in applying causal discovery methods and especially highlighted the challenges linked to the study of biosphere-atmosphere interaction via PCMCI. We demonstrated that the network structures estimated from observational data are explainable with respect to plant physiology and climatic effects. Finally, our
study shows that causal methods can deliver better interpretability and a much improved process understanding in comparison to correlation and bivariate Granger causality analyses that are ambiguous to interpret since they do not account for common drivers.

4.3 Outlook

The preceding discussion has shed light on the merits of PCMCI as well as the challenges of applying causal discovery methods. Runge et al. (2019) discuss further challenges and methods and give an outlook how multiple methods can be combined to alleviate limitations.
5 Conclusions

Here we tested PCMCI, an algorithm that estimates causal graphs from empirical time-series. We specifically explored two types of data sets that are highly relevant in biogeosciences: eddy covariance measurements of land-atmosphere fluxes and global satellite remote sensing of vegetation greenness. The causal graphs estimated from the eddy covariance data collected in a Mediterranean site confirm patterns we would expect in these ecosystems: During the dry season’s plants senescence, for instance, the ecosystem’s carbon cycle (NEE) decouples from meteorological variability. On the contrary during the main growing season with warm and humid conditions strong links between NEE, LE and H characterise the graph. Not only the strongly contrasting states emerge in the graph structure using the causal framework, but also the gradual transitions that relate to minor changes like the connectivity of sensible heat to temperature with progressing dryness. A purely correlative analysis, instead, is not able to resolve these patterns. PCMCI allows us to identify and focus on much fewer, but highly relevant dependencies only. Applying the approach to three replicated eddy covariance systems shows the robustness of the method to random errors in the fluxes measurements and confirm one of the assumption of eddy covariance: above a relatively homogeneous terrain the fluxes measured should be spatially invariant, and so the underlying causal relationship between climate and fluxes. The global analysis of NDVI in relation to climatic drivers confirms the known patterns of dependence strengths of vegetation on climatic variables: boreal regions are energy limited and especially driven by temperature and secondarily radiation, while in semi-arid regions vegetation dynamics are strongly dependent on water supply. However, obtained response times of vegetation to climatic variations are lower using PCMCI than correlation which can be attributed to a better treatment of the autocorrelation in the time-series and cross-relations among climate variables. Compared to merely correlative approaches, this leads to a interpretable pattern of driver-response relationships. In short, the new developments achieved in causal inference allow to gain well constrained insights on processes, that would otherwise be drowning in the correlation chaos. Therefore we hope that this study fosters usage of causal inference in analysing interactions and feedbacks of the biosphere-atmosphere system and furthermore exhibits our demand of further developments.
Table A1. PCMCI parameters that were used differently from default settings.

| Dataset           | significance | $\alpha_{pc}$ | $\tau_{min}$ | $\tau_{max}$ | selected_variables | mask_type | fdr_method |
|-------------------|--------------|---------------|--------------|--------------|--------------------|-----------|------------|
| Test Model        | 0.01         | 0.4           | 0            | 25           | [1,2,3]            | 'none'    | 'none'     |
| Majadas Dataset   | 0.01         | None          | 0            | 8            | [1,2,3,4,5]        | 'y'       | 'fdr_bh'   |
| Gridded global data set | 0.05       | None          | 0            | 3            | [1,2,3]            | 'y'       | 'fdr_bh'   |

Code and data availability. The eddy covariance data of the FLUXNET sites can be downloaded from the official webpage (https://fluxnet.fluxdata.org/). CRU temperature and precipitation data is available at http://badc.nerc.ac.uk/data/cru/. CRUNCEP radiation data can be downloaded via ftp://nacp.ornl.gov/synthesis/2009/frescati/temp/land_use_change/original/readme.htm. The NDVI dataset is available at http://ecocast.arc.nasa.gov/data/pub/gimms/3g/.

The TIGRAMITE software package that includes PCMCI can be found on github https://github.com/jakobrunge/tigramite/. All other code will be made available upon request.
| FLUXNET-ID | start year | end year | Data Reference       |
|-----------|------------|----------|----------------------|
| AT-Neu    | 2002       | 2012     | Wohlfahrt et al.     |
| AU-Cpr    | 2010       | 2014     | Meyer et al. (2015)  |
| AU-DaP    | 2007       | 2013     | Beringer et al. (2011a) |
| AU-DaS    | 2008       | 2014     | Hutley et al. (2011) |
| AU-Dry    | 2008       | 2014     | Cernusak et al. (2011) |
| AU-How    | 2001       | 2014     | Beringer et al. (2007) |
| AU-Stp    | 2008       | 2014     | Beringer et al. (2011b) |
| AU-Tum    | 2001       | 2014     | Leuning et al. (2005) |
| BE-Lon    | 2004       | 2014     | Moureaux et al. (2006) |
| BE-Vie    | 1996       | 2014     | Aubinet et al. (2001) |
| BR-Sa3    | 2000       | 2004     | Saleska et al. (2003) |
| CA-Man    | 1994       | 2008     | Brooks et al. (1997) |
| CA-NS2    | 2001       | 2005     | Bond-Lamberty et al. (2004) |
| CA-NS3    | 2001       | 2005     | Wang et al. (2002a) |
| CA-NS5    | 2001       | 2005     | Wang et al. (2002b) |
| CA-NS6    | 2001       | 2005     | Wang et al. (2002c) |
| CA-Qfo    | 2003       | 2010     | Chen et al. (2006) |
| CA-SF2    | 2001       | 2005     | Rayment and Jarvis (1999a) |
| CA-SF3    | 2001       | 2006     | Rayment and Jarvis (1999b) |
| CH-Cha    | 2005       | 2014     | Merbold et al. (2014) |
| CH-Dav    | 1997       | 2014     | Zielis et al. (2014) |
| CH-Fru    | 2005       | 2014     | Imer et al. (2013) |
| CH-Lae    | 2004       | 2014     | Etzold et al. (2011) |
| CH-Oe1    | 2002       | 2008     | Ammann et al. (2009) |
| CH-Oe2    | 2004       | 2014     | Dietiker et al. (2010) |
| CZ-wet    | 2006       | 2014     | Dušek et al. (2012) |
| DE-Akm    | 2009       | 2014     | Bernhofer et al. (a) |
| DE-Geb    | 2001       | 2014     | Anthoni et al. (2004) |
| DE-Gri    | 2004       | 2014     | Prescher et al. (2010a) |
| DE-Hai    | 2000       | 2012     | Knohl et al. (2003a) |
| DE-Kli    | 2004       | 2014     | Prescher et al. (2010b) |
| DE-Lkb    | 2009       | 2013     | Lindauer et al. (2014) |
| DE-Obe    | 2008       | 2014     | Bernhofer et al. (b) |
| DE-Spw    | 2010       | 2014     | Bernhofer et al. (c) |
| DE-Tha    | 1996       | 2014     | Grünewald and Bernhofer (2007) |
| DK-NuF    | 2008       | 2014     | Westergaard-Nielsen et al. (2013) |
| DK-Sor    | 1996       | 2014     | Pilegaard et al. (2011) |

| FLUXNET-ID | start year | end year | Data Reference       |
|-----------|------------|----------|----------------------|
| DK-ZaH    | 2000       | 2014     | Lund et al. (2012) |
| FI-Hyy    | 1996       | 2014     | Suni et al. (2003) |
| FI-Sod    | 2001       | 2014     | Thum et al. (2007) |
| FR-Fon    | 2005       | 2014     | Delpierre et al. (2016) |
| FR-LBr    | 1996       | 2008     | Berbigier et al. (2001) |
| FR-Pue    | 2000       | 2014     | Rambal et al. (2004) |
| GF-Guy    | 2004       | 2014     | Bonal et al. (2008) |
| IT-BCi    | 2004       | 2014     | Vitale et al. (2016) |
| IT-Col    | 1996       | 2014     | Valenti et al. (1996) |
| IT-Lav    | 2003       | 2014     | Marcolla et al. (2003) |
| IT-MBo    | 2003       | 2013     | Marcolla et al. (2011) |
| IT-Noe    | 2004       | 2014     | Spano et al.  |
| IT-Ro1    | 2000       | 2008     | Rey et al. (2002)  |
| IT-SRo    | 1999       | 2012     | Chiesi et al. (2005) |
| IT-Tor    | 2008       | 2014     | Galvagno et al. (2013) |
| JP-SMF    | 2002       | 2006     | Matsumoto et al. (2008) |
| NL-Hor    | 2004       | 2011     | Jacobs et al. (2007) |
| RU-Fyo    | 1998       | 2014     | Kurhatova et al. (2008) |
| US-ARM    | 2003       | 2012     | Fischer et al. (2007) |
| US-Blo    | 1997       | 2007     | Schade et al. |
| US-Ha1    | 1991       | 2012     | Wofsy et al. (1993) |
| US-Me2    | 2002       | 2014     | McDowell et al. (2004) |
| US-Me6    | 2010       | 2014     | Ruehr et al. (2012a) |
| US-MMS    | 1999       | 2014     | Pryor et al. (1999) |
| US-Ne1    | 2001       | 2013     | Gitelson et al. (2003) |
| US-Ne2    | 2001       | 2013     | Cassman et al. (2003a) |
| US-Ne3    | 2001       | 2013     | Cassman et al. (2003b) |
| US-SRG    | 2008       | 2014     | Ruehr et al. (2012b) |
| US-SRM    | 2004       | 2014     | Scott et al. (2008) |
| US-Ton    | 2001       | 2014     | Tang et al. (2003) |
| US-Twt    | 2009       | 2014     | Hatala et al. (2012) |
| US-UMB    | 2000       | 2014     | Rothstein et al. (2000) |
| US-Var    | 2000       | 2014     | Nave et al. (2011) |
| US-Whs    | 2007       | 2014     | Xu et al. (2004) |
| US-Wkg    | 2004       | 2014     | Scott et al. (2006) |

Table B1. List of FLUXNET sites used for the generation of artificial datasets and the time period used.
Figure A1. Distribution of coupling coefficients obtained after fitting the test model to the Fluxnet sites. Here shown are the distributions used for generation of heteroscedastic time series.
Figure B1. Distribution of coupling coefficients obtained after fitting the test model to the Fluxnet sites. Here shown are the distributions used for generation of homoscedastic time series.
**Figure C1.** Distribution of time lags obtained after fitting the test model to the Fluxnet sites. Here shown are the distributions used for generation of heteroscedastic time series.
Figure D1. Distribution of time lags obtained after fitting the test model to the Fluxnet sites. Here shown are the distributions used for generation of homoscedastic time series.
Figure E1. Observed (blue) and model (orange) time-series for Hainich Fluxnet site. The model data was produced with heteroscedastic noise.
Figure F1. Same as Fig. 5 but using simple correlation analysis to estimate the graph structures. The number of significant occurrences of a link is given by its width. The link strength, given by the link color, is calculated by averaging the significant links of the towers. Link labels indicating the lag were removed to improve link visibility. They typically ranged from 1 to 8 (full range of possible lags). The resulting graphs are shown for April 2014 till March 2015. The significance threshold is 0.01.
Figure G1. Precipitation in Majadas de Tiétar measured at the three tower sites from April 2014 to March 2015. Missing values are set to -1.
Author contributions. CK and MDM designed the study with contributions from JR and DGM. CK conducted the analysis and wrote the manuscript. All authors helped to improve the manuscript. AC, MM, TEM, OPP conducted field experiments in Majadas, processed and provided its data as well helped with interpretation.

Competing interests. The authors declare that they have no competing interests.

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