



## 1 Introduction

Modelling of in situ soil water dynamics requires knowledge of the soil hydraulic properties that characterize the soil at the site of interest. Various methods have been developed to determine the soil hydraulic properties on soil samples taken to the laboratory. Among these laboratory methods, the multi-step outflow method (van Dam et al., 1994; Durner and Iden, 2011) and the evaporation method (Šimůnek et al., 1998; Peters and Durner, 2008) have found widespread application because they allow for the determination of both the soil water retention function and the soil hydraulic conductivity function from a single experiment.

However, there is ever increasing evidence in the soil hydrological literature that the hydraulic properties determined on soil cores in the laboratory are of limited use for predictive modelling of soil water dynamics under field conditions (e.g. Mertens et al., 2005; Baroni et al., 2010). The possible reasons for this are manifold. For example, Mallants et al. (1997) suggested that the sample volume analysed in the laboratory may not be a representative elementary volume. Another likely reason was discussed by Basile et al. (2003), who stressed that laboratory experiments typically aim at the determination of the soil hydraulic functions that belong to the primary drainage curve, that is, starting from an initial condition that is as closely as possible to complete saturation. This condition, however, basically never occurs in situ, and the so derived properties may therefore not be representative for in situ conditions. Closely related to the argument of Basile et al. (2003) is the fact that hysteresis is typically neglected in modelling of in situ soil water dynamics, even though it is widely acknowledged to be an intrinsic feature of soils which affects soil water dynamics (e.g. Dane and Wierenga, 1975; Lenhard et al., 1991). Other effects, like dynamic non-equilibrium (Diamantopoulos and Durner, 2012), which may be significant in some situations but are neglected in standard modelling approaches may further limit the usefulness of the hydraulic properties derived in the laboratory to predict water dynamics in field soils.

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Given these limitations it is promising to make use of in situ observations of soil water state variables to obtain more useful estimates of the soil hydraulic properties by inverse modelling (Vereecken et al., 2008). The soil hydraulic properties derived in this way give the best possible description of the observational data. Since they are conditional on the model formulation used for parameter inference, they are also deemed particular useful for predictive modelling. However, as the soil hydraulic properties are part of the flow model, they may compensate for deficits in the model formulation and for errors in the forcing variables.

From a statistical point of view, the estimation of soil hydraulic properties by inverse modelling of in situ observations of soil water state variables is afflicted with two complicating features of this type of observational data. The first complicating feature is that in situ observations generally do not contain sufficient information to warrant accurate and precise estimation of the soil hydraulic properties (Scharnagl et al., 2011). The reason for this is that in situ data cover only a relative narrow range of soil water states. This is true even for observations taken close to the surface where soil water dynamics are highest.

To account for the limited amount of information of in situ observations, it is particularly beneficial to make explicit use of prior information about the soil hydraulic parameters in the inference process. This can be done using Bayesian statistics, which offers a formal way to combine the information on the parameters before assimilating field data and the information contained in the observational data. This formal Bayesian approach was applied by Hou and Rubin (2005), who incorporated prior information about the soil hydraulic properties in the calibration of a one-layer soil hydrological model using neutron probe and time domain reflectory (TDR) measurements. They used the principle of minimum relative entropy (Woodbury and Ulrych, 1993) to define a prior probability density function of the soil hydraulic parameters from information about the expected values, variances, as well as the lower and upper bounds of the parameters derived from the ROSETTA pedotransfer function (Schaap et al., 2001). Scholer et al. (2011) investigated the information content of ground-penetrating radar measure-

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ments for estimating the soil hydraulic properties of a four-layer profile. They used prior information about the soil hydraulic parameters either taken from Carsel and Parrish (1988) or derived from independent measurements on core samples taken at the field site. Scholer et al. (2011) concluded that prior information helped to substantially constrain the parameter space, especially when a prior density was used that includes information about the correlation structure of the soil hydraulic parameters. Concurring results were presented by Scharnagl et al. (2011) who used spatially distributed TDR measurements to estimate the effective soil hydraulic parameters of a single-layer profile. They implemented a Monte Carlo approach that allows to estimate the correlation structure of the soil hydraulic parameters predicted by ROSETTA, which is not available from the standard output of this pedotransfer function. Scharnagl et al. (2011) found that only if the full covariance matrix was used to define the prior density of the soil hydraulic parameters, the Bayesian approach was effective and robust in constraining the parameters to a small region of the parameter space.

The second complicating feature of in situ observations is that the corresponding residuals, that is, the deviations between observations and model predictions, are often strongly autocorrelated. The obvious reason for this are model errors, that is, errors in the model structure as well as errors in the forcing variables that are used to run the model, which all together result in a systematic misfit between observations and model predictions. Doherty and Welter (2010) provide an overview of the various formal and informal statistical approaches that are currently used to explicitly treat autocorrelation in the residuals.

Here, we adopted a formal statistical approach, which requires that the autocorrelation structure of the residuals is considered in the inference process. This can be done using the classical first-order autoregressive (AR(1)) model. This model has found widespread application in many fields of science, such as econometrics (e.g. Cochrane and Orcutt, 1949; Beach and MacKinnon, 1978) or catchment hydrology (e.g. Sorooshian and Dracup, 1980; Kuczera, 1983). Quite surprisingly, up to now it has received very little attention in the soil hydrological literature, where autocorrelation

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of the residuals as well as possible deviations from the assumptions of homogeneity and normality are typically neglected in the inference process. We are currently aware of only one study that considered autocorrelation of the residuals in inverse modelling of soil hydrological processes. Wöhling and Vrugt (2011) implemented the classical AR(1) scheme to estimate the soil hydraulic properties of a five-layer profile from multiple, parallel time series of soil water content and pressure head recorded at five depths. They found that the calibrated model did not provide a simultaneous fit of both pressure head and soil water content observations, with the model predicted soil water contents deviating substantially and consistently from the corresponding observations. Wöhling and Vrugt (2011) suspected that the reason for this failure are structural errors in the soil hydrological model that cannot be captured by the AR(1) scheme.

In this study, we considered a simple example of inverse modelling of in situ soil water dynamics: a bare, homogeneous soil profile for which a time series of soil water content observations near the surface was available. The inverse problem was posed in a formal Bayesian framework and solved using Markov chain Monte Carlo (MCMC) simulation. The general problem of limited information content of in situ data was taken into account by using an informative prior distribution of the soil hydraulic parameters, as was previously done by Scharnagl et al. (2011) at the same field site. We applied three different likelihood models that differ in their underlying assumptions about the statistical features of the error. The first likelihood model corresponds exactly to the ordinary least-squares approach, which is the de facto standard in inverse modelling of soil hydrological processes. The second likelihood model implements the classical AR(1) model to account for autocorrelation, which was previously used by Wöhling and Vrugt (2011) in a soil hydrological application. This second likelihood model additionally allows for heteroscedastic and non-Gaussian distributed residuals. In the third likelihood model, we introduced a modified AR(1) scheme that overcomes a critical deficiency of the classical scheme.

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The lower boundary was defined by a constant pressure head:

$$h(t) = h_{\text{LB}} \quad \text{at } z = -100 \text{ cm.} \quad (4)$$

The pressure head at the lower boundary  $h_{\text{LB}}$  was treated as an additional unknown parameter in the soil hydrological model.

5 The soil hydrological model used in this study corresponds exactly to the one used in Scharnagl et al. (2011). More detailed information about the model setup and the rationale for treating the lower boundary condition as an unknown parameter can be found in this reference.

### 2.3 Likelihood models

10 To simplify notation, let us denote the VGM parameters by vector  $\mathbf{x}_1 = [\theta_r \theta_s \alpha n K_s L]$  and the pressure head at the lower boundary by  $\mathbf{x}_2 = [h_{\text{LB}}]$ . To draw inference about the unknown soil hydrological parameters  $\mathbf{x}_1$  and  $\mathbf{x}_2$  given the observations of soil water content and meteorological forcing, the information contained in the time series of residuals  $\varepsilon_1, \dots, \varepsilon_N$  is used, where  $N$  is the number of observations. The residuals are defined as:

$$\varepsilon_i(\mathbf{x}_1, \mathbf{x}_2) = \tilde{y}_i - y_i(\mathbf{x}_1, \mathbf{x}_2) \quad i = 1, \dots, N \quad (5)$$

where  $\tilde{y}_i$  denotes the  $i$ th water content observation, and  $y_i$  is the corresponding model prediction. Formal statistical inference requires a likelihood model that describes the statistical features of the time series of residuals as closely and consistently as possible. The common first step in the development of the three likelihood models used in 20 the present study was the standardization of the residuals:

$$\underline{\varepsilon}_i(\mathbf{x}_1, \mathbf{x}_2, \mu_i, \sigma_i) = \frac{\varepsilon_i - \mu_i}{\sigma_i} \quad i = 1, \dots, N \quad (6)$$

where  $\mu_i$  and  $\sigma_i$  denote the expectation and SD of the  $i$ th residual, respectively. A common assumption about the general behaviour of the expectation of the residuals, which

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was also adopted in this study, was that  $\mu_i$  equals zero for all observations:

$$\mu_i = 0 \quad i = 1, \dots, N. \quad (7)$$

In the following, we refer to Eq. (7) as the zero expectation assumption. It reflects our prior belief that the applied model is a useful approximation of reality. If provided with 5 sufficient information about the distribution of material properties within the soil volume of interest, appropriate initial and boundary conditions, as well as model parameters, such a model should be able to describe the observations without bias. This basic assumption was adopted throughout this study.

To make the inference process feasible, we need to make three other basic assumptions 10 on the statistical nature of the residuals. The first relates to the SD of the  $i$ th residual  $\sigma_i$ , which we assume either to be homoscedastic or heteroscedastic, that is, to have either a constant or a variable SD. The second assumption relates to the serial independence of the residuals or its counterpart, autocorrelation. Last but not least, an assumption must be made on which statistical distribution appropriately represents 15 the actual distribution of the residuals. In the remainder of this subsection, we give a detailed description of the three different likelihood models.

**Likelihood 1:** The first likelihood model corresponds exactly to the ordinary least-squares approach. In this approach, it is assumed that the residuals are homoscedastic, that is,  $\sigma_i$  takes on some constant value:

$$\sigma_i = \sigma_{\text{const}} \quad i = 1, \dots, N. \quad (8)$$

It is common in applied Bayesian statistics to eliminate the constant SD  $\sigma_{\text{const}}$  from the inference equations (e.g. Kavetski et al., 2006; Scharnagl et al., 2011). This is done because the primary interest is often not in this parameter of the likelihood model but rather in the parameters of the process model. In this study, 25 however, we treated  $\sigma_{\text{const}}$  as an additional parameter that needs to be estimated simultaneously with the soil hydrological parameters  $\mathbf{x}_1$  and  $\mathbf{x}_2$ .

By assuming that the standardised residuals follow a Gaussian distribution, the likelihood of an individual standardised residual  $\underline{\varepsilon}_i$ , and hence, of the corresponding observation  $\tilde{y}_i$  is:

$$\rho(\tilde{y}_i|\mathbf{x}_1, \mathbf{x}_2, \sigma_{\text{const}}) = \frac{1}{\sqrt{2\pi}\sigma_{\text{const}}} \exp\left(-\frac{\varepsilon_i^2}{2}\right) \quad i = 1, \dots, N. \quad (9)$$

5 Based on the additional assumption that the residuals are independent, the joint likelihood of all observation  $\tilde{y}_1, \dots, \tilde{y}_N$  is:

$$\rho(\tilde{y}_1, \dots, \tilde{y}_N|\mathbf{x}_1, \mathbf{x}_2, \sigma_{\text{const}}) = \prod_{i=1}^N \rho(\tilde{y}_i|\mathbf{x}_1, \mathbf{x}_2, \sigma_{\text{const}}). \quad (10)$$

The ordinary least-squares approach is the de facto standard in inverse modelling of soil hydrological processes (e.g. Jacques et al., 2002; Sonnleitner et al., 2003; Wollschläger et al., 2009; Schelle et al., 2012). However, its underlying assumptions are rarely tested in practice. In this study, the assumptions of homoscedastic, independent, and Gaussian distributed residuals did not hold in the case of inverse modelling of in situ soil water dynamics and good statistical practice then requires that the likelihood model is revised accordingly.

15 **Likelihood 2:** The second likelihood model allows for heteroscedastic and non-Gaussian distributed residuals and implements the classical AR(1) scheme to account for serial dependence of the residuals. To parametrise heteroscedasticity, we modelled the SD of the  $i$ th residual as a function of the simulated water content,  $\sigma_i(y_i)$ , using a piecewise cubic Hermite interpolating polynomial (PCHIP, Fritsch and Carlson, 1980). The PCHIP approach offers great flexibility in modelling the underlying, but a priori unknown variation of the SD of the residuals as

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a function of the simulated water content. It was inspired by the “free-form” approach (Bitterlich et al., 2004; Iden and Durner, 2007) to model the soil hydraulic properties.

5 The PCHIP is calculated on four points, defined by matrix  $\Omega$  (size  $4 \times 2$ ). The relationship  $\sigma_i(y_i)$  can then formally be written as:

$$\text{PCHIP: } \Omega = \begin{bmatrix} \theta_a & \sigma_a \\ \theta_b & \sigma_b \\ \theta_c & \sigma_c \\ \theta_d & \sigma_d \end{bmatrix} \rightarrow \sigma_i(y_i) \quad i = 1, \dots, N \quad (11)$$

10 where  $\theta_{a, \dots, d}$  and  $\sigma_{a, \dots, d}$  represent the  $x$  and  $y$  ordinates of the four points, respectively. The  $x$  ordinates of the first and last of these points were defined by  $\theta_a = 0.9 \tilde{y}_{\min}$  and  $\theta_d = 1.1 \tilde{y}_{\max}$ , respectively, where  $\tilde{y}_{\min}$  denotes the minimum and  $\tilde{y}_{\max}$  the maximum of the soil water content observations. This definition of  $\theta_a$  and  $\theta_d$  was chosen to avoid extrapolation of the PCHIP. The remaining two  $x$  ordinates,  $\theta_b$  and  $\theta_c$ , were set equidistantly between the minimum and maximum values. The corresponding  $y$  ordinates,  $\sigma_{a, \dots, d}$ , are unknown parameters of the likelihood model.

15 In a next step, we consider the autocorrelation of the residuals using the classical AR(1) scheme. Its application presumes homoscedasticity of the residuals (e.g. Box et al., 2008; Evin et al., 2013) and has therefore to be applied to the standardised residuals. The classical AR(1) scheme is defined by:

$$\eta_i(\mathbf{x}_1, \mathbf{x}_2, \Omega, \phi) = \underline{\varepsilon}_i - \phi \underline{\varepsilon}_{i-1} \quad i = 2, \dots, N \quad (12)$$

20 where  $\eta_i$  represents the decorrelated standardised residuals and  $\phi$  is the autocorrelation coefficient. The decorrelated residuals  $\eta_i$  can be considered as the remaining, unexplained deviations between observations and corresponding model predictions after removal of the systematic deviations that can be explained by

autocorrelation. The autocorrelation coefficient  $\phi$  is an additional parameter of the likelihood model. In general, for a AR(1) process to be stationary, the autocorrelation coefficient must satisfy  $|\phi| < 1$  (e.g. Box et al., 2008).

The classical AR(1) scheme in Eq. (12) was successfully applied to inverse modelling in catchment hydrology (e.g. Sorooshian and Dracup, 1980; Kuczera, 1983; Bates and Campbell, 2001; Vrugt et al., 2009b). It failed, however, in the soil hydrological application reported by Wöhling and Vrugt (2011). To understand the reason for this failure, it is instructive to look at the expectation of the decorrelated and restandardised residuals  $\underline{\eta}_i$ . Taking expectations on both sides of Eq. (12) gives:

$$\begin{aligned} E(\eta_i) &= E(\varepsilon_i) - E(\phi)E(\varepsilon_{i-1}) \\ &= (1 - E(\phi))E(\varepsilon_i) \quad i = 2, \dots, N \end{aligned} \quad (13)$$

where we used  $E(\varepsilon_i) = E(\varepsilon_{i-1})$ , which follows directly from the basic assumption given in Eq. (7). The expectation of the decorrelated residuals shows that the classical AR(1) scheme may introduce bias in the decorrelated residuals because the expectation of  $\eta_i$  will always be closer to zero than the expectation of  $\varepsilon_i$ . This disadvantage is most pronounced if  $E(\phi) \rightarrow 1$ , in which case  $E(\eta_i)$  approaches zero even if  $E(\varepsilon_i)$  differs markedly from zero. In fact, as we illustrated in this study, this bias may distort the inference process and may lead to meaningless results, very similar to those obtained by Wöhling and Vrugt (2011).

The transformation of the residuals implied by the AR(1) scheme in Eq. (12) needs to be accounted for in the formulation of the likelihood model. To do so, we need to restandardise the decorrelated residuals as:

$$\underline{\eta}_i(x_1, x_2, \Omega, \phi) = \frac{\eta_i}{\sqrt{1 - \phi^2}} \quad i = 2, \dots, N \quad (14)$$

where  $\sqrt{1 - \phi^2}$  is just the SD of the decorrelated residuals (e.g. Box et al., 2008).  
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If we further assume that the restandardised decorrelated residuals  $\underline{\eta}_i$  follow a Student distribution with unit variance, the likelihood of a single observation is given by:

$$p(\tilde{y}_i | x_1, x_2, \Omega, \phi, \nu) = \frac{\Gamma\left(\frac{\nu+1}{2}\right) \sqrt{\frac{\nu}{\nu-2}}}{\Gamma\left(\frac{\nu}{2}\right) \sqrt{\pi\nu} \sqrt{1 - \phi^2} \sigma_i} \left(1 + \frac{\underline{\eta}_i^2}{\nu-2}\right)^{-\frac{\nu+1}{2}} \quad i = 2, \dots, N \quad (15)$$

where  $\Gamma$  denotes the gamma function and  $\nu > 2$  is a kurtosis parameter that controls the tailing behaviour of the distribution. For  $\nu \rightarrow 2$ , the tailing of the Student distribution becomes increasingly heavy, while for  $\nu \rightarrow \infty$  the Student distribution converges to the Gaussian distribution. This is illustrated in Fig. 1a. The Student distribution has been proposed for the use in parameter inference by West (1984) and Lange et al. (1989), among others, because it has the ability to adequately fit heavy-tailed distributions, which also makes the inference process more robust against possible outliers in the observations.

In a next step, we applied the method of Fernández and Steel (1998) to introduce skewness in the symmetric Student distribution shown in Eq. (15). The skewed Student distribution is given by:

$$\begin{aligned} p(\tilde{y}_i | x_1, x_2, \Omega, \phi, \nu, \kappa) &= \frac{2c_2 \Gamma\left(\frac{\nu+1}{2}\right) \sqrt{\frac{\nu}{\nu-2}}}{\left(\kappa + \frac{1}{\kappa}\right) \Gamma\left(\frac{\nu}{2}\right) \sqrt{\pi\nu} \sqrt{1 - \phi^2} \sigma_i} \\ &\quad \times \left(1 + \frac{1}{\nu-2} \left(\frac{c_1 + c_2 \underline{\eta}_i}{\kappa \operatorname{sgn}(c_1 + c_2 \underline{\eta}_i)}\right)^2\right)^{-\frac{\nu+1}{2}} \quad i = 2, \dots, N \end{aligned} \quad (16)$$

where  $\operatorname{sgn}$  represents the signum function,  $c_1$  and  $c_2$  are correction terms that account for the shift in expectation and variance of the decorrelated and restandardised residuals caused by the skewing mechanism, respectively, and  $\kappa > 0$





**2.4 Bayes’ theorem**

The joint likelihood summarizes in a probabilistic way what we can learn about the estimated parameters given the observations. However, sometimes it is desirable to make explicit use of prior knowledge about the parameters in the inference process. Bayes’ theorem provides a formal way of combining information from observations with prior information about the estimated parameters. In case of Likelihood 1, Bayes’ theorem can be written as:

$$\rho(\mathbf{x}_1, \mathbf{x}_2, \sigma_{\text{const}}|\tilde{\mathbf{y}}) \propto \rho(\mathbf{x}_1) \rho(\mathbf{x}_2) \rho(\sigma_{\text{const}}) \rho(\tilde{\mathbf{y}}|\mathbf{x}_1, \mathbf{x}_2, \sigma_{\text{const}}) \tag{22}$$

where the term  $\rho(\mathbf{x}_1, \mathbf{x}_2, \sigma_{\text{const}}|\tilde{\mathbf{y}})$  on the left hand side is the posterior density of the estimated parameters given the observations.  $\rho(\mathbf{x}_1)$ ,  $\rho(\mathbf{x}_2)$ , and  $\rho(\sigma_{\text{const}})$  represent the prior densities of the parameters, and  $\rho(\tilde{\mathbf{y}}|\mathbf{x}_1, \mathbf{x}_2, \sigma_{\text{const}})$  is just Likelihood 1, given in Eq. (10). Similar expressions for the posterior density are obtained for Likelihood 2 and 3.

For computational reasons, it is often not the prior, likelihood, and posterior that is used in practice, but rather its natural logarithm. This is because the prior, likelihood, and posterior may take on values larger or smaller than the largest or smallest number that can be represented by a double-precision floating-point format. This is especially true if the joint likelihood of a large number of observations is calculated, as it was the case in the present study. The log-prior, log-likelihood, and log-posterior was therefore used in all calculations, and it is these numbers that are reported here as model diagnostics.

**2.5 Prior densities**

In order to apply Bayes’ theorem, the prior densities of the parameters need to be defined. As a prior of the VGM parameters we chose a truncated multivariate normal distribution  $\rho(\mathbf{x}_1) \sim \mathcal{T}\mathcal{N}(\boldsymbol{\mu}_{x_1}, \boldsymbol{\Sigma}_{x_1}, a_{x_1}, b_{x_1})$ , where  $\boldsymbol{\mu}_x$  denotes the mean vector of the VGM parameters,  $\boldsymbol{\Sigma}_x$  is the corresponding covariance matrix, and  $a_{x_1}$  and  $b_{x_1}$  represent

the upper and lower bounds, respectively. The mean vector and covariance matrix was derived in Scharnagl et al. (2011) using the ROSETTA pedotransfer function (Schaap et al., 2001). Here, we make use of the full covariance matrix (Prior 3 in Scharnagl et al., 2011), which also contains information about the correlation structure of  $\mathbf{x}_1$ . In addition, we used the lower and upper bounds on the VGM parameters reported in Scharnagl et al. (2011) and listed in Table 1 to truncate the Gaussian distribution. The use of the truncated multivariate normal distribution resulted in slightly faster convergence of the Markov chains (Sect. 2.6) to the region of the parameter space with highest posterior density compared to using the conventional multivariate normal distribution as a prior.

The priors of the remaining parameters were given by a uniform distribution, which assigns a constant probability density within a specified interval and zero else. This so-called noninformative prior reflects our knowledge about these parameters before the inference process. In case of the pressure head at the lower boundary, the uniform prior can formally be written as  $\rho(x_2) \sim \mathcal{U}(a_{x_2}, b_{x_2})$ , where  $a_{x_2}$  and  $b_{x_2}$  denote the corresponding lower and upper bound of that parameter, respectively. Similar expressions can be given for the unknown parameters in the likelihood model. The boundaries for all these parameters are also listed in Table 1.

We used a  $\log_{10}$ -transformation for parameters  $\alpha$ ,  $n$ , and  $K_s$  in the VGM model (Scharnagl et al., 2011). The kurtosis parameter in Likelihood 2 and 3 was transformed to  $1/v$  following the suggestion of Lange et al. (1989). These transformations aim at reducing nonlinear correlations of the estimated parameters, which helps to improve the performance of the MCMC scheme.

**2.6 Markov chain Monte Carlo simulation**

The posterior distribution in Eq. (22) cannot be calculated by analytical means. Instead, we used MCMC simulation (e.g. Brooks, 1998) to generate samples from the posterior distribution. We used the Differential-Evolution-Markov-chain (DE-MC) framework (ter Braak, 2006; ter Braak and Vrugt, 2008) to run the MCMC simulations. The algorithm we applied generates candidate points from an archive of past states (DE-MC<sub>(Z)</sub>,

ter Braak and Vrugt, 2008) and decides upon their acceptance by the Metropolis rule Metropolis et al. (1953). We did two minor changes to the standard DE-MC<sub>(Z)</sub> scheme, which aimed at improving the performance of this MCMC scheme. Without these modifications, the exploration of the parameter space by the various Markov chains run in parallel was quite poor if Likelihood 2 or 3 was used. In fact, standard DE-MC<sub>(Z)</sub> even failed to locate the region of the parameter space with highest posterior density in some of the test runs. The two modifications improved the performance in these cases substantially. The two modifications are as follows:

1. We implemented a block updating scheme, similar to what was used in ter Braak (2006). The idea behind block updating is that the convergence to a stationary distribution can sometimes be increased substantially if only a subset of the estimated parameters is updated in a single step. This general finding motivated Vrugt et al. (2009a) to implement a self-adaptive randomized updating scheme into DE-MC, resulting in the DiffeREntial Evolution Adaptive Metropolis (DREAM) algorithm. This updating strategy did not work well in our application, presumably because it ignores the correlation structure of the VGM parameters induced by the informative prior of these parameters. Instead of the randomized subspace sampling used in DREAM, we therefore used a block updating scheme, as follows. For the first 2000 steps in each of the parallel chains, each parameter block was updated with probability 0.8. The VGM parameters were considered a block, whereas the remaining parameters were considered a block each. This simple block updating scheme increased the exploration capabilities of the Markov chains and the convergence to a stationary distribution substantially. The block updating was restricted to the initial phase of the MCMC simulation because it is not particularly efficient any more once the chains have approached the stationary distribution.
2. Roberts and Rosenthal (2001) suggested that acceptance rates between 0.15 and 0.5 can generally be considered efficient, with an optimal value being 0.23.

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Within standard DE-MC, the jumping distance of a proposal is scaled such that optimal acceptance rates are obtained when sampling from Gaussian and Student posteriors (Roberts and Rosenthal, 2001; ter Braak, 2006). Here, we introduce a factor  $r$  (dimensionless) that reduces this reference scaling of the jumping distance. The reasoning for this reduction is that efficient sampling from posterior distributions that differ from the multivariate Gaussian or Student distribution generally require smaller jumps than those taken by the reference scaling used in the standard DE-MC scheme. In the present study, the reduction factor was set to  $r = 1.0$  in the case of Likelihood 1, that is, no reduction, and  $r = 0.5$  in the cases of Likelihood 2 and 3. With these settings, acceptance rates near the optimal value of 0.23 were obtained in all three cases.

Apart from these two changes to the DE-MC<sub>(Z)</sub> scheme, we used standard settings for the algorithmic parameters reported by ter Braak (2006) and ran five chains in parallel. Convergence of the Markov chains to the stationary distribution was monitored using the interval diagnostic of Brooks and Gelman (1998). After convergence, we ran each of the five chains for another 10 000 steps. Parameter inference was based on 5000 samples selected from this final sampling stage by thinning the chains, that is, by retaining every 10th sample. The calculation of the uncertainty bounds of the model prediction is described in Schoups and Vrugt (2010), and is therefore not repeated herein.

### 3 Results

Figure 2 shows the results of inverse modelling of the time series of in situ soil water content observations using Likelihood 1 together with precipitation and potential evaporation fluxes that define the upper boundary condition in HYDRUS-1-D. The parameter estimates that belong to the prediction with maximum posterior density are listed in Table 1. This table also lists the corresponding mean error, root mean square error,

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model efficiency (Nash and Sutcliffe, 1970), as well as the log-prior, log-likelihood, and log-posterior, which serve as model diagnostics to rate the goodness of the model prediction. Based on visual inspection of Fig. 2, the ordinary least-squares approach implemented in the first likelihood model results in a good overall description of the observational data. This impression is also corroborated by the various model diagnostics listed in Table 2.

To check the assumptions of homoscedastic, independent, and Gaussian distributed standardised residuals  $\underline{\varepsilon}_i$  used in ordinary least-squares, we made use of three diagnostic plots. Figure 3a depicts the standardised residuals plotted against the simulated soil water content. This diagnostic plot indicates that the residuals are heteroscedastic. The SD of the residuals increases with increasing values of the predicted soil water content. Figure 3b shows the autocorrelation function of the residuals. This diagnostic plot shows that the residuals are strongly autocorrelated. The autocorrelation of the residuals at lag 1, representing a time interval of 3 h, is close to one. The autocorrelation decreases rather slowly with increasing lag in an approximately exponential manner, which points to an autoregressive process of first order (e.g. Box et al., 2008). At lag 60, corresponding to a temporal separation of 7.5 days, the autocorrelation is still significant, as indicated by the theoretical 95 % uncertainty bounds for a time series of uncorrelated random variables. Figure 3c depicts a quantile–quantile plot of the standardised residuals. This diagnostic plot indicates that the actual distribution of the standardised residuals differs from that of a Gaussian distribution. The actual distribution appears to be bimodal and also differs in its tailing behaviour from that of a Gaussian distribution. In summary, the three diagnostic plots show that neither of the three assumptions used in the formulation of Likelihood 1 is fulfilled. In this case, good statistical practice requires that the likelihood model is revised and that the inference process is repeated with the revised likelihood function.

The results for Likelihood 2, which allows for heteroscedastic, autocorrelated, and non-Gaussian distributed residuals, are presented in Fig. 4. The predicted soil water content deviates substantially and consistently from the observations. Clearly, the pa-

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parameter estimates obtained with this likelihood model are rather meaningless. They violate the fundamental assumption of zero expectation of the residuals given in Eq. (7). Essentially the same results were obtained by Wöhling and Vrugt (2011), who also used the classical AR(1) scheme in their case study. We explain this repeated failure with the fact that the classical AR(1) scheme introduces bias in the expected value of the residuals, as shown in Eq. (13). Figure 4 shows that the decorrelated and restandardised residuals  $\underline{\eta}_i$  are on average very close to zero even though the standardised residuals  $\underline{\varepsilon}_i$  deviate substantially from this value. This is reflected by a large mean error and root mean square error of the prediction listed in Table 2. As briefly discussed in Sect. 2.3, the bias of the classical AR(1) scheme increases with increasing values of the autocorrelation coefficient  $\phi$ . Indeed, the estimated autocorrelation coefficient  $\phi$  obtained with Likelihood 2 is very close to the upper bound of this parameter (Table 1). For  $\phi \rightarrow 1$ , the AR(1) process becomes instationary, which is clearly not a meaningful result in a parameter inference setting.

In contrast, the modified AR(1) scheme used in Likelihood 3 does not suffer from the deficiency of the classical scheme, as shown in Eq. (21). The results obtained with this likelihood model are depicted in Fig. 5. The prediction gives a good overall description of the observations, which by visual inspection appears to be as good as the prediction obtained with Likelihood 1. Using the modified AR(1) scheme implemented in Likelihood 3, both the standardised residuals  $\underline{\varepsilon}_i$  and the decorrelated and restandardised residuals  $\underline{\eta}_i$  have an expectation close to zero and a SD close to one. This result is obtained with an estimated autocorrelation coefficient  $\phi$  that does not touch the upper bound of this parameter and has a significant smaller value than that obtained with Likelihood 2 (Table 1). The mean error and root mean square error of the model predicted soil water content are slightly larger for Likelihood 3 compared to values obtained with Likelihood 1 (Table 2), while the model efficiency indicates an equally good performance of both models. From a statistical point of view, however, the superiority of the parameter estimates obtained with Likelihood 3 over the ones obtained with Likelihood 1 becomes evident when comparing the corresponding posterior densities. The

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log-likelihood as well as the log-posterior obtained with Likelihood 3 are substantially larger than the corresponding values obtained with Likelihood 1 (Table 2). The larger log-likelihood value already indicates that Likelihood 3 better describes the statistical features of the residuals.

5 Figure 6 illustrates the standardization of the residuals using the PCHIP approach and also shows the diagnostic plots used to check the underlying assumptions of the likelihood model. In Fig. 6a, the raw residuals  $\varepsilon_i$  are plotted as a function of the predicted soil water content. It becomes visible here that the variance of the raw residuals varies as function of the predicted soil water content, that is, that the  $\varepsilon_i$  are heteroscedastic. However, what becomes also clear from this figure is that the expectation of the raw residuals also varies as a function of the predicted soil water content. The soil hydrological model systematically overestimates larger observations, which is an obvious expression of bias of the process model. The results of the PCHIP approach to estimate the SD of the raw residuals as a function of the predicted soil water content is shown in Fig. 6b. The corresponding standardised residuals  $\underline{\varepsilon}_i$  are depicted in Fig. 6c. The standardised residuals appear to be homoscedastic, even though the systematic deviation of the expectation of  $\underline{\varepsilon}_i$  persists. We explain this by the fact that the PCHIP approach partially compensates for model bias by assigning larger SDs to the more biased residuals.

20 The three assumptions of homoscedasticity, independence, and skewed Student distribution of the decorrelated and restandardised residuals  $\eta_i$  that underlie Likelihood 3 are checked in the three diagnostic plots shown in Fig. 6d to f. These plots show that the three underlying assumptions are approximately fulfilled in the case of Likelihood 3. The decorrelated and restandardised residuals  $\eta_i$  used in this likelihood model appear to be approximately homoscedastic, independent, and skewed Student distributed. From a statistical point of view, the parameter estimates obtained with Likelihood 3 can therefore be considered statistically valid. It is also noteworthy that the estimated values of the kurtosis parameter  $\nu$  and the skewness parameter  $\kappa$  indicate that the actual distribution of the residuals deviates substantially from that of a Gaussian dis-

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tribution. The parameter estimates given in Table 1 show that the actual distribution is much more heavy tailed than the Gaussian distribution and that it is slightly left-skewed.

In the following, we compare the estimated soil hydrological parameters and the resulting soil hydraulic functions obtained with Likelihood 1 and 3. Figure 7 depicts pairwise scatter plots of the posterior samples obtained with these two likelihood models together with the prior 95 % uncertainty regions. The figure illustrates that all model parameter were identified uniquely. Although some of the soil hydrological parameters correlated strongly, all parameter estimates generally showed little variance. Another important result of this comparison was that the parameter estimates obtained with Likelihood 1 and 3 differ systematically. The pairwise scatter plots show almost no overlap of the two posterior samples. It is noteworthy that the shape of the posterior sample obtained with Likelihood 3 deviates somewhat from that of a multivariate normal distribution. Even this slight deviation, however, adversely affected the performance of the MCMC scheme and gave reason for the use of a reduced reference scaling of the jumping distance described in Sect. 2.6. The uniqueness of the estimated parameters was not restricted to the hydrological parameters, but was also observed for the parameters of the likelihood model. The posterior samples for these parameters generally showed little variance and no strong correlations with any of the other parameters, neither with those of the likelihood model nor with those of the soil hydrological model (not shown).

25 The soil hydraulic properties computed from the posterior samples obtained with Likelihood 1 and 3 are depicted in Fig. 8. In this figure, we display the 95 % posterior uncertainty bounds for the water retention and hydraulic conductivity function together with the corresponding 95 % prior uncertainty bounds. The posterior uncertainty bounds are generally very tight, reflecting the high precision of the soil hydraulic parameters which is illustrated in Fig. 7. In general, the posterior uncertainty bounds are tightest in the intermediate soil water content range, which corresponds to the range of actual soil water content observations. This is also where the soil hydraulic functions obtained with the two likelihood models match each other most.

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#### 4 Discussion

The application of Likelihood 1, which makes use of the ordinary least-squares assumptions, resulted in parameter estimates that are invalid from a statistical point of view because the residuals violate all underlying assumptions of this likelihood model.

5 Neglecting autocorrelation in the residuals means to overestimate the information content of the data and therefore leads to an underestimation of parameter uncertainty. This becomes evident in the parameter sample from the posterior shown in Fig. 7: the variance of the posterior obtained with Likelihood 1 is generally smaller than that obtained with Likelihood 3. If residuals are correlated, the amount of information provided by each observation is smaller than in the case of uncorrelated observations. In fact, using a likelihood model which neglects autocorrelation essentially implies to treat the soil hydrological model as if it was perfect. This is overoptimistic because environmental models are – almost by definition – at best good approximations of real-world processes: they cannot be perfect.

15 We showed that the classical AR(1) scheme implemented in Likelihood 2 resulted in unrealistic parameter estimates and, correspondingly, to biased model predictions. This corroborates the results reported by Wöhling and Vrugt (2011), who used the classical AR(1) scheme in a soil hydrological application. We showed analytically that the AR(1) scheme introduces bias, which explains the failure of this autoregressive model in the present study as well as in the application of Wöhling and Vrugt (2011). Obviously, inverse modelling of in situ soil water dynamics seems to be much more susceptible to the bias problem induced by the classical AR(1) scheme than, for example, inverse problems in catchment hydrology, where it has been applied successfully many times (e.g. Sorooshian and Dracup, 1980; Bates and Campbell, 2001). We conjecture that this is because the systematic misfit between observations and model predictions depicted in Fig. 4 can easily be compensated for by an estimated value of the autocorrelation coefficient that is larger than actually necessary. In a recent study, Evin et al. (2014) showed that the results obtained with the classical AR(1) model applied

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to inverse modelling of stream flow can be nonrobust, indicated by strong correlations between parameters in the hydrological model and the likelihood model. We suspect that the reason for this nonrobustness might also be explained, at least partially, with the bias that is introduced by the classical AR(1) scheme.

5 The residuals obtained with Likelihood 3 clearly satisfied all assumptions of this likelihood model and furthermore demonstrate that Likelihood 3 solves the bias-problem inherent to Likelihood 2. Since the log-likelihood and log-posterior values for Likelihood 3 were substantially larger than those obtained with Likelihood 1 (Table 2), we consider this likelihood function and the parameter estimates obtained with it as clearly superior and statistically valid. The model predictions obtained with Likelihood 3 slightly violate the zero-expectation assumption, which becomes apparent from the non-constant expectation of the residuals shown in Fig. 6a. This model bias can in principle be accounted for in the formulation of the likelihood model by introducing a term for bias correction, which can for example be estimated as a function of the predicted soil water content (e.g. Kennedy and O'Hagan, 2001; Erdal et al., 2012). From our point of view, however, such an approach would not be meaningful in the present application. This is because the bias correction would be valid only for the depth within the soil profile where the observations were taken, but it remains unclear how this bias correction affects the predicted soil water contents at other depths within the profile and the fluxes to adjacent compartments within the water cycle. This is a fundamental difference between the Bayesian approach adopted in this study and other statistical approaches that are designed to account for model bias, such as particle filtering (e.g. Montzka et al., 2011; Vrugt et al., 2013). Within the the present statistical framework, we think that the model bias can therefore only be meaningfully corrected for by a thorough revision of the process model, the assumption of a homogeneous profile, the parametrisation of the soil hydraulic properties, and the inclusion of hysteresis.

Likelihood 3, which allows for heteroscedastic, autocorrelated, and non-Gaussian residuals, may be equally useful in other soil hydrological applications. In general, autocorrelation of the residuals becomes increasingly apparent if observations are taken

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at a sufficiently high temporal frequency, which is typically the case for automated measurement setups either in the field or in the laboratory. For example, autocorrelation of the residuals is particularly pronounced in inverse modelling of multi-step outflow experiments (e.g. Diamantopoulos et al., 2012), evaporation experiments (e.g. Dettmann et al., 2014), or lysimeter data (e.g. Schelle et al., 2012). Based on the findings of the present study, we conjecture that a likelihood model that allows for heteroscedastic, autocorrelated, and non-Gaussian residuals would also prove useful in these applications. The main advantages are a more accurate quantification of the model parameters and their uncertainty. In the case of multiple time series used for parameter inference, Likelihood 3 must be extended to the multivariate case by using a first-order vector autoregressive (VAR(1)) model (e.g. Box et al., 2008). In contrast to the AR(1) model treated in this study, the VAR(1) model additionally accounts for the cross-correlation between the multiple, parallel time series of observations.

## 5 Summary and conclusions

In this case study, we estimated the soil hydraulic properties of a homogeneous, bare soil profile from in situ observations of soil water content taken at a single location near the soil surface. This type of observational data often contain too little information to warrant accurate and precise estimates of all of the parameters in the soil hydrological model. We posed the inverse problem in a formal Bayesian approach, which allowed us to use prior information of the soil hydraulic parameters in terms of an informative prior distribution for these parameters. The prior distribution was taken from Scharnagl et al. (2011). We then applied three likelihood models, which differ in their underlying assumptions about the salient statistical features of the time series of residuals.

The first likelihood model corresponds to the ordinary least-squares approach, which is the de facto standard in inverse modelling of soil hydrological processes. However, the results obtained with Likelihood 1 showed that the underlying assumptions of this likelihood model of homoscedastic, independent, and Gaussian residuals were all vio-

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lated. From a statistical point of view, the parameter estimates obtained with this first likelihood model can therefore be considered invalid.

The second likelihood model makes use of the classical AR(1) scheme to account for autocorrelated residuals, allows for heteroscedasticity, and supports heavily-tailed, non-Gaussian distributions of the residuals. To describe the heteroscedastic nature of the residuals, we devised a flexible model for the relationship between model predicted soil water content and the SD of the residuals that uses cubic Hermite interpolation. Deviations from a Gaussian distribution were modelled with a skewed Student distribution. A theoretical analysis of this parameter inference scheme and the results obtained with Likelihood 2 demonstrated that the classical AR(1) model introduces bias into the inference process. Due to this bias, the results obtained with Likelihood 2 were essentially meaningless, which corroborates the findings of Wöhling and Vrugt (2011) obtained in a similar soil hydrological study and using the same classical AR(1) scheme.

In the third likelihood model, we introduced a modified AR(1) scheme, which does not suffer from the bias problem induced by the classical scheme. With this likelihood model, we obtained parameter estimates that differed significantly from those obtained with Likelihood 1, which implements the ordinary least-squares assumptions. Furthermore, parameter uncertainty was higher as compared to Likelihood 1 because the decreased information content of the calibration data caused by autocorrelation was accounted for. Since the residuals obtained with Likelihood 3 fulfilled the underlying statistical assumptions, and because Likelihood 3 lead to a much larger log-posterior than Likelihood 1, we concluded that the parameter estimates obtained with Likelihood 3 are statistically valid, and therefore, superior to the ones obtained with Likelihood 1.

The results of this study demonstrate that the ordinary least-squares approach may lead to statistically invalid estimates of the soil hydraulic parameters, because the underlying assumptions about the statistical features of the time series of residuals are not fulfilled. The current practice in inverse modelling of soil hydrological processes should therefore be revised. We introduced a new, highly flexible likelihood model that accounts for autocorrelated, heteroscedastic, and non-Gaussian residuals, which per-

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formed very well in the present test study. Based on our findings, we advocate the use and further extension of the new likelihood model in inverse modelling of soil hydrological processes.

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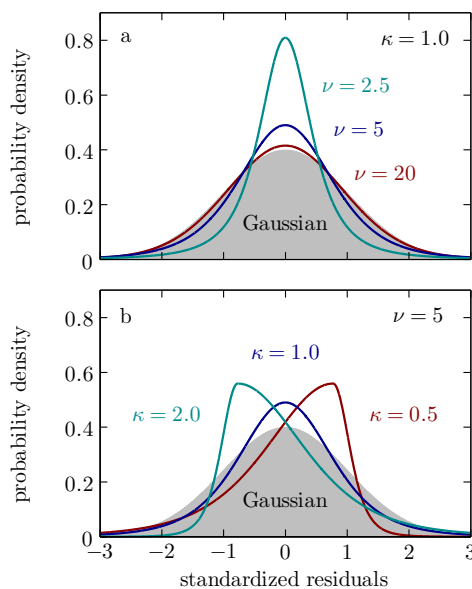
**Table 1.** Estimated parameters in the soil hydrological and likelihood model, respectively, as well as parameter estimates with maximum posterior density obtained using the three different likelihood models.

	parameter	unit	transformation	lower bound	upper bound	parameter estimates with maximum posterior density		
						likelihood model 1	likelihood model 2	likelihood model 3
soil hydrological model	$\theta_r$	$\text{cm}^3 \text{cm}^{-3}$	–	0.043	0.091	0.051	0.061	0.053
	$\theta_s$	$\text{cm}^3 \text{cm}^{-3}$	–	0.409	0.481	0.461	0.460	0.436
	$\alpha$	$\text{cm}^{-1}$	$\log_{10}(\alpha)$	0.0028	0.0085	0.0045	0.0051	0.0037
	$n$	–	$\log_{10}(n)$	1.51	1.85	1.77	1.70	1.77
	$K_s$	$\text{cm h}^{-1}$	$\log_{10}(K_s)$	0.006	0.832	0.534	0.254	0.091
	$L$	–	–	–5.49	6.27	4.20	1.76	3.67
	$h_{LB}$	cm	–	–250	–50	–170	–51	–178
likelihood model	$\sigma_{\text{const}}$	$\text{cm}^3 \text{cm}^{-3}$	–	0.000	0.025	0.0121		
	$\sigma_a$	$\text{cm}^3 \text{cm}^{-3}$	–	0.000	0.050		0.0183	0.0092
	$\sigma_b$	$\text{cm}^3 \text{cm}^{-3}$	–	0.000	0.050		0.0190	0.0131
	$\sigma_c$	$\text{cm}^3 \text{cm}^{-3}$	–	0.000	0.050		0.0316	0.0218
	$\sigma_d$	$\text{cm}^3 \text{cm}^{-3}$	–	0.000	0.050		0.0498	0.0360
	$\phi$	–	–	0.000	1.000		0.996	0.957
	$\nu$	–	$1/\nu$	2.000	$\infty$		3.32	2.89
	$\kappa$	–	–	0.500	2.000		1.27	0.85

**Table 2.** Summary statistics of model residuals and the values of the prior, likelihood and posterior computed for the parameter estimates with maximum posterior density. Results are given for the three different likelihood models.

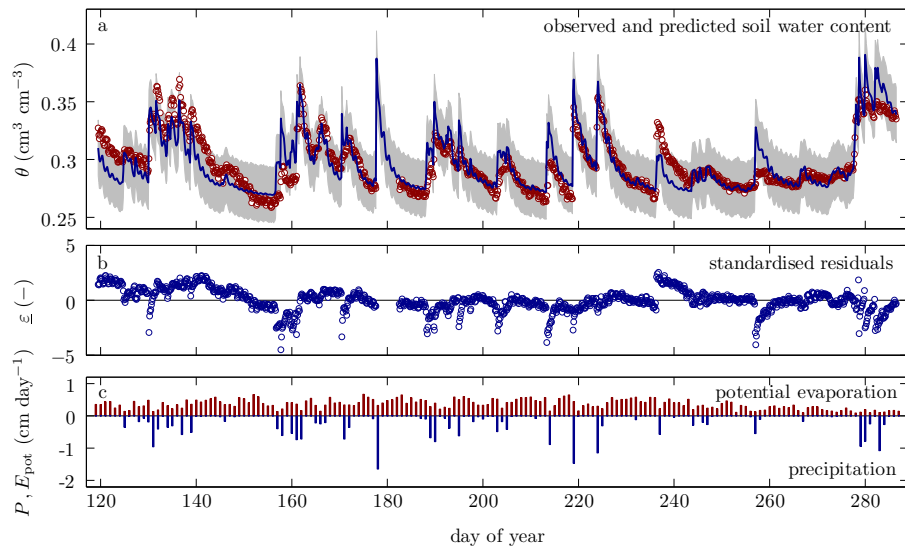
model diagnostic	likelihood model 1	likelihood model 2	likelihood model 3
mean error	-0.0001	-0.0738	-0.0012
root mean square error	0.0121	0.0754	0.0132
model efficiency	0.74	0.01	0.74
log-prior	23	28	27
log-likelihood	3891	5880	5761
log-posterior	3914	5908	5788

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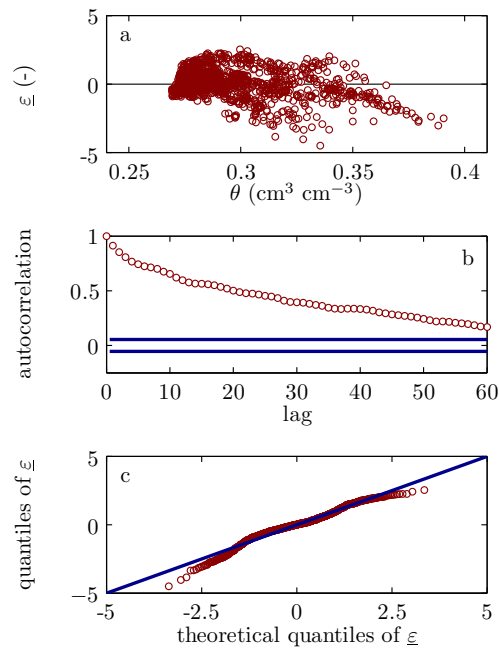
**Figure 1.** Illustration of the skewed Student distribution. In **(a)**, the skewness parameter is kept constant ( $\kappa = 1$ ) while the kurtosis parameter is varied. In **(b)**, the skewness parameter is varied while the kurtosis parameter is kept constant ( $\nu = 5$ ). To allow for a direct comparison, the corresponding Gaussian distribution is also shown. All distributions have an expectation of zero and a SD of one.

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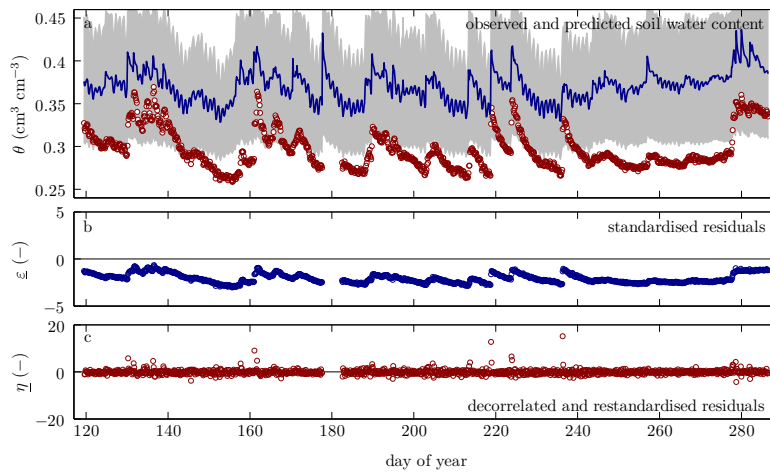
**Figure 2.** Model prediction of the time series of soil water content observations obtained using Likelihood 1. In (a), the model prediction with maximum posterior density and the 95% uncertainty bounds of the prediction are shown together with the observations of soil water content  $\theta$ . (b) displays the standardised residuals  $\underline{\varepsilon}_i$ , corresponding to the prediction with maximum posterior density. (c) shows the precipitation  $P$  (negative) and potential evaporation  $E_{\text{pot}}$  (positive) fluxes used as input to the HYDRUS-1-D model. For graphical reasons, daily fluxes are shown here whereas hourly values were used to run the model.

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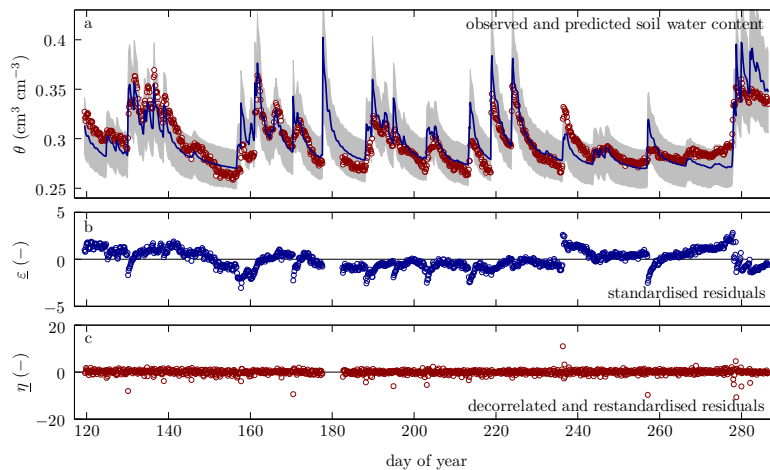
**Figure 3.** Diagnostic plots of the standardised residuals  $\underline{\varepsilon}_i$  of the model prediction with maximum posterior density using Likelihood 1. In (a), the standardised residuals  $\underline{\varepsilon}_i$  are plotted over the predicted water content  $\theta$ . (b) depicts the autocorrelation function with theoretical 95% uncertainty intervals of a time series of uncorrelated random variables, and (c) shows a quantile–quantile plot of the standardised residuals.

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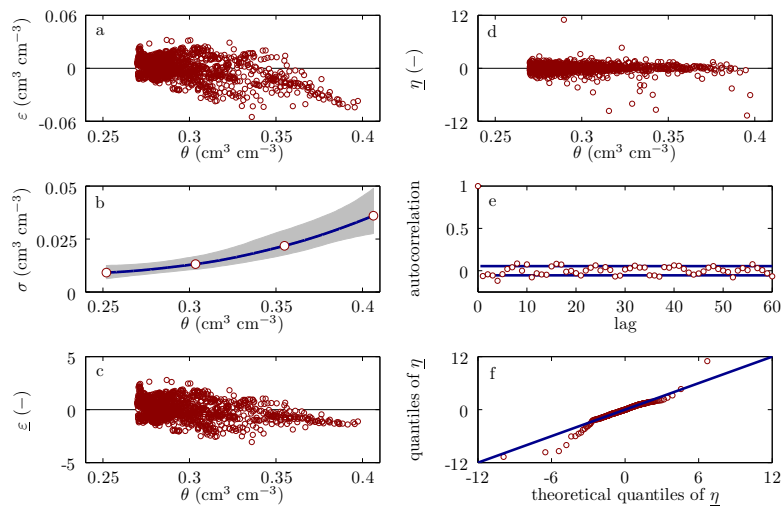
**Figure 4.** Model prediction of the time series of soil water content observations obtained using Likelihood 2. In (a), the model prediction with maximum posterior density and the 95% uncertainty bounds of the prediction are shown together with the observations of soil water content  $\theta$ . (b) displays the standardised residuals  $\underline{\epsilon}_i$ , corresponding to the prediction with maximum posterior density. (c) shows the decorrelated and restandardised residuals  $\underline{\eta}_i$ .

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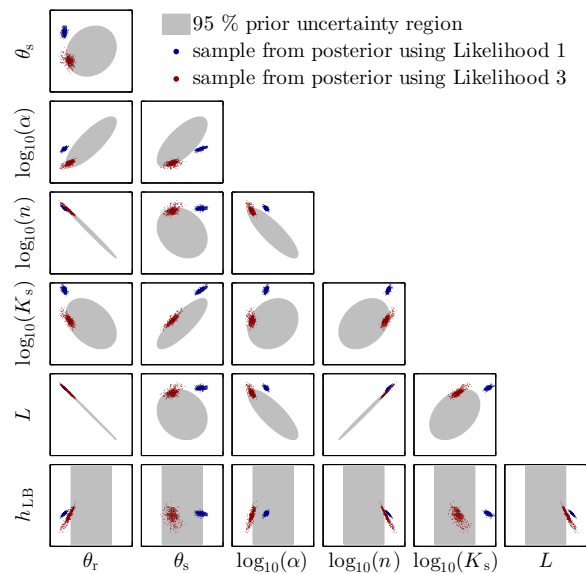
**Figure 5.** Model prediction of the time series of soil water content observations obtained using Likelihood 3. In (a), the model prediction with maximum posterior density and the 95% uncertainty bounds of the prediction are shown together with the observations of soil water content  $\theta$ . (b) displays the standardised residuals  $\underline{\epsilon}_i$ , corresponding to the prediction with maximum posterior density. (c) shows the decorrelated and restandardised residuals  $\underline{\eta}_i$ .

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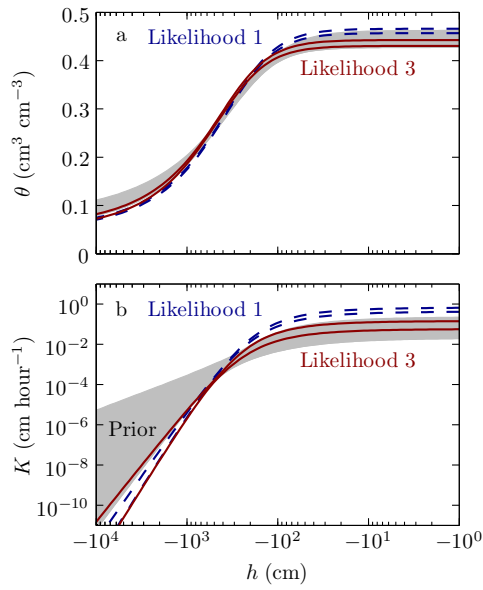
**Figure 6.** Performance of the PCHIP approach used for standardization of the residuals and diagnostic plots of the decorrelated and restandardised residuals corresponding to the model prediction with maximum posterior density using Likelihood 3. **(a)** shows the residuals  $\varepsilon_i$  as a function of the predicted soil water content  $\theta$ . **(b)** gives the estimated SD as a function of the predicted soil water content, showing the estimated points  $\sigma_{a,\dots,d}$  and the resulting function given by the PCHIP. The grey area represents the corresponding 95 % uncertainty. In **(c)** the standardised residuals  $\underline{\varepsilon}_i$  are plotted over the predicted water content, while **(d)** shows the decorrelated and restandardised residuals  $\underline{\eta}_i$ . **(e)** depicts the autocorrelation function of  $\underline{\eta}_i$  with theoretical 95 % uncertainty intervals of a time series of uncorrelated random variables, and **(f)** shows a quantile–quantile plot of  $\underline{\eta}_i$ .

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**Figure 7.** Pairwise scatter plots of the posterior samples of the parameters of the soil hydrological model obtained with Likelihood 1 and 3 together with the corresponding 95 % prior uncertainty regions of these parameters. The axes cover the parameter bounds listed in Table 1.

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**Figure 8.** 95% posterior uncertainty bounds of **(a)** the water retention function and **(b)** the hydraulic conductivity function obtained with Likelihood 1 and 3. Additionally, the corresponding 95% prior uncertainty regions are shown as a grey area.