1 Data-worth Analysis through Probabilistic Collocation-based Ensemble Kalman

- 2 Filter
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Abstract

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We propose a new and computationally efficient data-worth analysis and quantification framework keyed to the characterization of target state variables in groundwater systems. We focus on dynamically evolving plumes of dissolved chemicals migrating in randomly heterogeneous aquifers. An accurate prediction of the detailed features of solute plumes requires collecting a substantial amount of data. Otherwise, constraints dictated by the availability of financial resources and ease of access to the aquifer system suggest the importance of assessing the expected value of data before these are actually collected. Data-worth analysis is targeted to the quantification of the impact of new potential measurements on the expected reduction of predictive uncertainty based on a given process model. Integration of the Ensemble Kalman Filter method within a data-worth analysis framework enables us to assess data worth sequentially, which is a key desirable feature for monitoring scheme design in a contaminant transport scenario. However, it is remarkably challenging because of the (typically) high computational cost involved, considering that repeated solutions of the inverse problem are required. As a computationally efficient scheme, we embed in the data-worth analysis framework a modified version of the Probabilistic Collocation Method-based Ensemble Kalman Filter proposed by Zeng et al. (2011) so that we take advantage of the ability to assimilate data sequentially in time through a surrogate model constructed via the polynomial chaos expansion. We illustrate our approach on a set of synthetic scenarios involving solute migrating in a two-dimensional random permeability field. Our results demonstrate the computational efficiency of our approach and its ability to quantify the

- 37 impact of the design of the monitoring network on the reduction of uncertainty
- associated with the characterization of a migrating contaminant plume.
- 39 Keywords: Data worth; Ensemble Kalman Filter; Probabilistic Collocation;
- 40 Contaminant migration;

1. Introduction

Groundwater resources constitute a remarkable reserve of multipurpose (including
potable, agricultural, and industrial) water and are critical for ecosystem and society
sustainability and development (Foster and Chiltion, 2003). The severe challenges
posed by the need to obtain accurate predictions of contaminant dynamics in natural
aquifers hinge on the diverse sources of uncertainty associated with a selected
predictive model. These uncertainties may originate from the (typically) unknown
spatial distribution of subsurface geomaterials and the intrinsic heterogeneity of their
hydrogeological properties, as well as from the insufficient level of knowledge of the
key processes governing contaminant transport at the scale of interest. Stochastic
inverse-modeling methods and data assimilation approaches have been developed to
characterize uncertainties through model calibration against available measurements
(e.g., Zimmerman et al., 1998; Alcolea et al., 2006; Fu and Gomez-Hernandez, 2009;
Hendricks-Franssen et al., 2009; Riva et al., 2009; Rubin et al., 2010; Zhang et al., 2013;
Zhou et al., 2014). In this context, the ensemble Kalman filter (EnKF) has gained
significant popularity for the purpose of sequential (in time) data assimilation due to its
relatively straightforward operational implementation and capability of quantification
of predictive uncertainty (Anonsen et al., 2009; Chen and Zhang, 2006; Oliver et al.,
2008; Xie and Zhang., 2010; Liu et al., 2012). Applications of the EnKF to contaminant
transport settings can be found in (Liu et al., 2008; Li et al., 2012).

All inverse-modeling and data assimilation methods require the collection of suitable types of data for system characterization. While some inverse approaches, such

as those analyzed by Zimmerman et al. (1998) and Hendricks-Franssen et al. (2009) and recently reviewed by Zhou et al. (2014), use all available data simultaneously, the fast growing ability of setting monitoring networks with the capability of delivering high-resolution real-time measurements has somehow increased the interest towards assimilation methods capable of sequentially updating models as soon as data become available. Even as sequential data assimilation tends to be computationally more efficient than traditional batch inverse approaches, in practice, the high cost associated with the extension of an existing monitoring setting suggests the need to develop robust methodologies conducive to the identification of optimal strategies for the collection of future data which are potentially valuable for a specific environmental goal considered. Following Back (2007), such goal oriented data sets can assist to improve one's ability to understand the system behavior and minimize uncertainty while considering budget constraints. Applications of data-worth analysis in the context of groundwater-related problems include the works of James and Freeze (1993), Abbaspour et al. (1996), Rada and Schultz (1998), Russell and Rabideau (2006), and Dausman et al. (2010). Nowak et al. (2010) and Leube et al. (2012) introduced the preposterior data impact assessor (PreDIA) in the data-worth analysis framework and applied it to a (late time) steadystate solute transport scenario taking place in a random hydraulic conductivity field. The use of PreDIA to assist the optimal design of a monitoring scheme by minimizing the probability of making incorrect decisions during Bayesian hypothesis testing is then introduced by Nowak et al. (2012) and illustrated with reference to the prediction of contaminant arrival times at a sensitive location in an aquifer, under a variety of

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The aforementioned rapid development of in-situ monitoring technologies motivates us to explore methodologies to quantify data-worth sequentially, to allow for an optimal and flexible dynamic (in space and time) adjustment of the monitoring scheme. To the best of our knowledge, data-worth analysis in the context of a dynamically evolving subsurface transport setting has not been studied in little literatures, Kollat et al. (2011) propose a framework for the design of a monitoring network upon combining the EnKF and multi-objective evolutionary optimization. In this context, Zhang et al. (2015) employ the relative entropy, or the Kullback-Leibler divergence, as a global metric according to which they study the issue of proposing a monitoring network for contaminant source identification. Here, we employ in our dataworth analysis framework an approach which relies on the EnKF concept because: (a) it is a non-intrusive method, which can be integrated in a straightforward manner in available computational systems, (b) it allows the flexibility of providing the uncertainty associated with the estimated system states at each assimilation step, thus facilitating the data-worth analysis; and (c) it can be extended to enable us to handle the challenges posed by the typically large number of parameters that are required to characterize hydrogeological systems under uncertainty.

A critical element that might hamper the efficiency of the approach is the large number of system replicates that are required to ensure the accuracy of the EnKF-based results and minimize filter inbreeding (see, e.g., Panzeri et al., 2014 and references therein). Therefore, considering that the planned additional measurements are not yet

collected and (at best) only estimates of these can be obtained from prior information, the EnKF can still be computational demanding when incorporated in a data-worth analysis. Panzeri et al. (2013, 2014) proposed to embed stochastic groundwater flow moment equations (MEs) in the EnKF in a way that obviates the need for Monte Carlo simulation. These authors demonstrated the computational feasibility and accuracy of the methodology on a moderate size problem and showed that the approach mitigates issues of filter inbreeding and spurious covariances often plaguing standard Monte Carlo based EnKF. While theoretically and operationally elegant and effective for small to medium size problems, the ME-based EnKF approach can be classified as an intrusive method at the current stage of development, because it requires solving equations satisfied by (conditional) statistical moments (ensemble means and covariances) of hydraulic heads and fluxes in randomly heterogeneous media, the structure of these equations being typically different from that of the equations governing the dynamics of the actual (random) state variables. Here, we rely on a nonintrusive approach which aims at improving computational efficiency through the construction of a surrogate model (or proxy) to replace the original system model in the standard Monte Carlo based EnKF. The use of surrogate models within a Bayesian inference (data assimilation) framework has been explored in petroleum engineering by Amudo et al. (2006) and Schaaf (2006). Carrer et al. (2007) relied on a variety of algorithms, such as polynomial regression and Kriging, to construct proxies of the target system response. Li et al. (2011) considered model proxies constructed by polynomial regression for subsurface flow related problems and showed that Kriging

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and neural network approaches may not yield accurate statistical moments of target state variables. As a consequence, results of an EnKF approach based on these types of proxies may be inaccurate.

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In recent years, there has been increasing interest in the use of techniques based on Polynomial Chaos Expansion (PCE) for the construction of surrogate models of subsurface flow and transport processes. The method was first introduced by Ghanem and Spanos (2003) and has then been employed in a variety of areas and for diverse purposes, including optimization and global sensitivity analysis in the context of uncertainty quantification of selected model outputs (e.g., Ghanem, 1998; Reagan et al., 2005; Sudret, 2008; Fajraoui et al., 2011; Hays et al., 2011; Oladyshkin and Nowak, 2012; Oladyshkin et al., 2012; Ciriello et al., 2013a, b; Formaggia et al., 2013; Dai et al., 2014; Wu et al., 2014, and references therein). PCE-based surrogate models have also been combined with a variety of Bayesian updating methods, including, e.g., Markov chain Monte Carlo method (MCMC) (Marzouk, 2007; Jin, 2008), the EnKF method (Saad and Ghanem, 2009; He et al., 2011) and Bootstrap filter (Oladyshkin et al. 2012a, b) to alleviate computational burden. In essence, the PCE method relies on: (a) constructing a representation of the system state of interest in terms of a polynomial expansion expressed as a function of a set of uncertain system parameters; and (b) deriving appropriate discretized equations for the (deterministic) coefficients of the expansion through a Galerkin technique. The solution of these (typically coupled) equations can be computationally demanding in the presence of a large number of uncertain parameters and/or high-order terms in the polynomial approximation. In this

context, the Probabilistic Collocation Method (PCM) is an efficient non-intrusive approach that can be employed to construct PCE-based proxies of groundwater flow models. Li and Zhang (2007, 2009) explored the ability and efficiency of the PCM to quantify uncertainty for single- and two-phase flow in randomly heterogeneous porous media by combining the approach based on the Karhunen-Loeve (KL) expansion with the PCE representation. Their results suggest that accurate estimates of key statistics of variables of interest, such as pressure heads or saturations, can be obtained with a limited number of runs of the original full system model. A PCM-based EnKF (PCKF) has been employed by Li and Xiu (2009), Zeng (2010), Zeng et al. (2011), and Li et al. (2014) for parameter estimation in flow settings typical of groundwater hydrology and petroleum engineering applications. In this work, we embed the PCKF into the data-worth analysis framework to assess the worth of dynamically monitored data in a contaminant transport setting taking place in a randomly heterogeneous aquifer. Doing so is consistent with the way we approach the data worth challenge from a theoretical standpoint, according to which only expected values (i.e., ensemble moments) of quantities of interest are required (see Section 2). Due to the need for repeated solutions of the inverse problem at the

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in a randomly heterogeneous aquifer. Doing so is consistent with the way we approach the data worth challenge from a theoretical standpoint, according to which only expected values (i.e., ensemble moments) of quantities of interest are required (see Section 2). Due to the need for repeated solutions of the inverse problem at the preposterior stage which can lead to high computational cost, we propose to increase computational efficiency in the data-worth analysis context by modifying the way PCKF is employed. In essence, unlike the PCKF introduced by Li and Xiu (2009), Zeng (2010), Zeng et al. (2011) and Li et al. (2014), in which the purpose of the PCKF was limited to improve the computational efficiency of the EnKF-based data assimilation in

the context of system parameter estimation and the statistics of uncertain geological system parameters are estimated on the basis of the updated PCE coefficients, our proposed modification to the PCKF method yields marked increase of computational efficiency in the repeated EnKF process required during the preposterior data worth analysis. To achieve this objective, we follow Zeng et al. (2012) and Marzouk and Xiu (2014) and update the system parameters by adjusting the random quantities in terms of which the PCE is constructed. As compared with the PCKF method originally illustrated by Zeng et al. (2011), our modified and adapted PCKF requires prior knowledge on the geostatistical descriptors characterizing the random parameter field, such as mean, covariance and integral scale, so that these do not need to be updated during the inverse modeling process. Through this modification, the entire simulation process relies directly on the model proxy which has been constructed prior to actual data collection, thus resulting in an alleviation of the computational burden associated with performing repeated inversing modeling, as required at the preposterior stage.

Section 2 illustrates the theoretical bases and the workflow of our PCKF-based data-worth analysis. Section 3 is devoted to the presentation of the set of examples that we employ to demonstrate the ability of the proposed framework to quantify the impact of the design of the monitoring network for solute concentrations (in terms of spatial location, temporal sampling frequency, and prior data content) on the reduction of uncertainty associated with the characterization of a contaminant plume migrating in a randomly heterogeneous aquifer. Conclusions are presented in Section 4.

2. Methodology

2.1 Data-worth analysis

We rely on the data-worth analysis framework proposed by Neuman et al. (2012) and Xue et al. (2014). The analysis is performed according to three stages. The first (or prior) stage relies on the prior data, i.e., the data which are available at the outset of the analysis. The second (or preposterior) stage relies on statistics of potential additional data conditional on the prior data. In this stage, the worth of the potential additional data from a given monitoring scheme is estimated. The third (or posterior) stage utilizes joint statistics of prior data and new data made available following the preposterior stage. The posterior statistics enable us to assess the quality of their associated preposterior estimates. Although we do not explore it in this work, our analysis framework can be extended to consider conceptual model uncertainty with the aid of the newly developed multimodel EnKF method (Xue and Zhang, 2014) which embeds uncertainty quantification associated with the use of diverse conceptual models of the system behavior.

Consider a set of discrete values of a target variable, representing, e.g., solute concentrations at a number of points distributed in space and time in an aquifer, and collected as the entries of a random vector, Δ . Here, we start by considering the mean, $E(\Delta|\mathbf{D})$, and covariance, $Cov(\Delta|\mathbf{D})$, of Δ conditioned on a discrete set of prior available data forming the entries of vector \mathbf{D} and predicted through a model characterized by a set of random variables grouped in vector ξ . In the preposterior stage, we assume that the prior data set \mathbf{D} is augmented by a set of data collected in vector \mathbf{C}' . The entries of \mathbf{C}' are yet to be observed and are planned to be collected in a

future monitoring campaign. At the preposterior stage, random estimates \mathbf{C} of \mathbf{C}' can be obtained on the basis of the results of the prior data-worth analysis, according, e.g., to the strategy illustrated in the following. Predictive statistics of Δ , i.e., mean, $E(\Delta | \mathbf{D}, \mathbf{C})$, and covariance, $Cov(\Delta | \mathbf{D}, \mathbf{C})$, are then calculated by joint conditioning on $\{\mathbf{D}, \mathbf{C}\}$. Note that the predictive statistics in the prior and preposterior data-worth analyses are theoretically related to each other by:

$$E(\Delta \mid \mathbf{D}) = E_{\text{CID}} E(\Delta \mid \mathbf{D}, \mathbf{C})$$
 (1)

$$Cov(\Delta \mid \mathbf{D}) = E_{CID}Cov(\Delta \mid \mathbf{D}, \mathbf{C}) + Cov_{CID}E(\Delta \mid \mathbf{D}, \mathbf{C})$$
 (2)

Here, $E_{C|D}E(\Delta|\mathbf{D},\mathbf{C})$ and $E_{C|D}Cov(\Delta|\mathbf{D},\mathbf{C})$, respectively, are the expectation of $E(\Delta|\mathbf{D},\mathbf{C})$ and $Cov(\Delta|\mathbf{D},\mathbf{C})$ over all \mathbf{C} vectors generated, conditional on \mathbf{D} ; and $Cov_{C|D}E(\Delta|\mathbf{D},\mathbf{C})$ is the covariance of $E(\Delta|\mathbf{D},\mathbf{C})$ calculated over all \mathbf{C} vectors generated, conditional on \mathbf{D} . Following Neuman et al. (2012) and Xue et al. (2014), a scalar measure quantifying the data-worth can be introduced through the trace operator:

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$$Tr \Big[Cov(\mathbf{\Delta} \mid \mathbf{D}) \Big] = Tr \Big[E_{C|D}Cov(\mathbf{\Delta} \mid \mathbf{D}, \mathbf{C}) \Big] + Tr \Big[Cov_{C|D}E(\mathbf{\Delta} \mid \mathbf{D}, \mathbf{C}) \Big]$$
(3)

where Tr indicates the trace (sum of diagonal entries) of a matrix. Note that $Trigl[E_{C|D}Cov(\Delta|\mathbf{D},\mathbf{C})igr]$ is given by the difference between the scalar measure of prior predictive uncertainty, $Trigl[Cov(\Delta|\mathbf{D})igr]$, and the scalar measure of the estimated posterior predictive uncertainty, $Trigl[E_{C|D}Cov(\Delta|\mathbf{D},\mathbf{C})igr]$. This quantity is to be compared against the reference data-worth, i.e., $Trigl[Cov(\Delta|\mathbf{D})igr] - Trigl[Cov(\Delta|\mathbf{D},\mathbf{C}')igr]$, which can be calculated at the posterior data-worth analysis stage, after \mathbf{C}' has been observed.

The conditional moments introduced above need to be obtained through an

appropriate inverse-modeling method. For example, Xue et al. (2014) employed the geostatistical inversion method of Hernandez et al. (2003, 2006) based on equations satisfied by (ensemble) moments of the target state variable in a steady-state groundwater flow setting. The dynamic nature of the process of contaminant transport suggests evaluating the data-worth sequentially, in a data assimilation framework. To this end, our approach takes advantage of the flexibility of the Monte Carlo based EnKF method (Chen and Zhang, 2006; Liu et al., 2008) to allow for sequential (in time) and simultaneous updating of a collection of state vector realizations.

For completeness, we recount here the key theoretical elements underlying the typical EnKF approach. We start by considering a collection of N_e realizations of the state vector \mathbf{S} :

$$\mathbf{S} = \left\{ \mathbf{s}^{1}, \mathbf{s}^{2}, ..., \mathbf{s}^{N_{e}} \right\}$$
 (4)

Superscripts in (4) refer to the number identifying the realization associated with each vector \mathbf{s} whose entries are given by the random quantities that characterize the model, $\mathbf{\xi}$, the dynamic state variables, \mathbf{u} , and the observation data, \mathbf{d}^{obs} . Observations at time t and their true values are related by:

$$\mathbf{d}_{t}^{obs} = \mathbf{H}\mathbf{s}_{t}^{true} + \boldsymbol{\varepsilon}_{t} \tag{5}$$

Here, the superscripts *obs* and *true* respectively stand for the observation data and the true (usually unknown) system state; measurement errors collected in vector $\mathbf{\epsilon}_t$ are assumed to be zero-mean Gaussian with covariance matrix \mathbf{R}_t ; matrix \mathbf{H} is the observation operator, which relates the state and observation vectors. The EnKF entails two stages, i.e., the forecast and assimilation stage. In the forecast step, each state vector

in the collection (4) is projected from time step (t-1) to time t via:

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$$\mathbf{s}_{t}^{f,i} = F(\mathbf{s}_{t-1}^{a,i}); \qquad i = 1, 2, ..., N_{e}$$
 (6)

- The operator $F(\bullet)$ in (6) represents the forward numerical/analytical model of choice;
- superscripts f and a indicate the forecast and assimilation stage, respectively. In the
- assimilation stage, the Kalman gain, G_t , is calculated as:

$$\mathbf{G}_{t} = \mathbf{C}_{t}^{f} \mathbf{H}_{t}^{T} \left(\mathbf{H}_{t} \mathbf{C}_{t}^{f} \mathbf{H}^{T} + \mathbf{R}_{t} \right)^{-1}$$
 (7)

- 267 \mathbb{C}_t^f being the covariance matrix of the system state. This matrix is approximated
- 268 through the N_e model realizations as:

$$\mathbf{C}_{t}^{f} \approx \frac{1}{N_{e}} \sum_{n=1}^{N_{e}} \left\{ \left[\mathbf{s}_{t}^{f,n} - \left\langle \mathbf{s}_{t}^{f} \right\rangle \right] \left[\mathbf{s}_{t}^{f,n} - \left\langle \mathbf{s}_{t}^{f} \right\rangle \right]^{T} \right\}$$
(8)

270 Each state vector in the collection is then updated as

$$\mathbf{s}_{t}^{a,i} = \mathbf{s}_{t}^{f,i} + \mathbf{G}_{t} \left(\mathbf{d}_{t}^{obs,i} - \mathbf{H} \mathbf{s}_{t}^{f,i} \right)$$
(9)

- 272 It is worth noting that the EnKF method is characterized by a linear updating step due
- to the first-order-second-moment approximation. As such, the results based on this
- approach are optimal in the presence of moderate non-linearity of the processes
- 275 governing the system dynamics.
- 276 The updated ensemble mean and covariance respectively are:

$$E\left(\mathbf{s}_{t}^{a} \mid \mathbf{d}_{1:t}^{obs}\right) = \frac{1}{N_{e}} \sum_{i=1}^{N_{e}} \mathbf{s}_{t}^{a}$$

$$\tag{10}$$

$$Cov\left(\mathbf{s}_{t}^{a} \mid \mathbf{d}_{1:t}^{obs}\right) = \frac{1}{N_{e} - 1} \sum_{i=1}^{N_{e}} \left\{ \left[\mathbf{s}_{i,t}^{a} - E\left(\mathbf{s}_{t}^{a}\right)\right] \left[\mathbf{s}_{i,t}^{a} - E\left(\mathbf{s}_{t}^{a}\right)\right]^{T}\right\}$$
(11)

where $\mathbf{d}_{1:t}^{obs}$ are is the vector of observations collected up to time t,

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$$d_{1:t}^{obs} = \left[d_1^{obs}, ..., d_t^{obs} \right]^T$$
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Incorporating the EnKF into the data-worth analysis framework enables one to

- quantify the worth of additional data in a sequential way, after assessing the uncertainty
- associated with data currently available. The workflow that we propose for assessing
- data-worth is depicted in Figure 1 and a synopsis of the various steps involved is
- detailed in the following.
- 286 1. At the prior stage, the EnKF is implemented on the basis of the observations
- available; a set of N_{e1} system realizations of state vectors is updated via (9) to obtain
- the ensemble mean $E(\mathbf{\Delta} | \mathbf{D})$ and covariance $Cov(\mathbf{\Delta} | \mathbf{D})$, conditioned on the
- prior data set **D**.
- 290 2. A number N_{e1} of random realizations of additional hypothetical data C are
- synthetically generated by assuming these are described through a multivariate
- Normal distribution, i.e., $\mathbf{C} \sim N \left[E(\mathbf{C} | \mathbf{D}), Cov(\mathbf{C} | \mathbf{D}) \right]$. Note that \mathbf{C} is a subset of
- 293 Δ obtained in step 1.
- 3. At the preposterior stage, EnKF is implemented through a set of N_{e2} Monte Carlo
- realizations for each of the N_{e1} hypothetical data vectors, \mathbb{C} , obtained at step 2; this
- allows calculating updated ensemble mean $E(\Delta | \mathbf{D}, \mathbf{C})$ and covariance
- 297 $Cov(\Delta \mid \mathbf{D}, \mathbf{C})$, jointly conditioned on $\{\mathbf{D}, \mathbf{C}\}$. Quantities $E_{C|D}Cov(\Delta \mid \mathbf{D}, \mathbf{C})$ and
- 298 $Cov_{CID}E(\Delta \mid \mathbf{D}, \mathbf{C})$ appearing in (3) are then calculated by averaging over the
- collection of $N_{e1} \times N_{e2}$ realizations. The corresponding scalar measure of the
- worth of additional data, $Tr \lceil Cov_{C|D} E(\Delta \mid \mathbf{D}, \mathbf{C}) \rceil$, can be calculated as the trace of
- 301 the resulting covariance matrix.
- 4. At the posterior stage, the actual measurement vector, C', becomes available. The
- ensemble mean $E(\Delta | \mathbf{D}, \mathbf{C}')$ and covariance $Cov(\Delta | \mathbf{D}, \mathbf{C}')$, jointly conditioned

on \mathbf{D} and $\mathbf{C'}$, can be obtained via EnKF in a way similar to the procedure outlined in step 1. The scalar measure of the reference data-worth in light of the additional measurements is then calculated as $Tr[Cov(\Delta|\mathbf{D})] - Tr[Cov(\Delta|\mathbf{D},\mathbf{C'})]$. This step (a) is employed to assess the quality of the preposterior estimates obtained at step 2, and (b) can be regarded as the prior analysis stage for a subsequent round of sequential data-worth assessment.

The above procedure shows that, for a given monitoring strategy, a total number of $N = N_{e1} \times N_{e2}$ numerical solutions of the system model are required to assess the data-worth in the preposterior analysis step. Since N_{e1} and N_{e2} need to be sufficiently large to ensure the accuracy of the estimated covariance matrix (*Li et al.*, 2014; *Zeng et al.*, 2010), we introduce the PCKF to address the challenge posed by the computational efficiency of the analysis.

2.2 Data-worth analysis via PCKF

The PCKF has been recently developed as an alternative to the traditional Monte Carlo based EnKF. It combines the polynomial chaos expansion (PCE) introduced by Wiener (1938) with EnKF.

According to Ghanem (1998), a target random variable can be expressed by a PCE with a set of deterministic coefficients. These coefficients are typically calculated through the Galerkin method (Ghanem, 1998; Mathelin et al., 2005). The latter entails computing the PCE coefficients upon solving a set of coupled equations. This somehow hampers a routine application of the method to groundwater flow and contaminant transport problems in which the relationship between the uncertain system parameters

and state variables is typically nonlinear. The probabilistic collocation method (PCM) enables one to compute the PCE coefficients by solving a set of uncoupled equations at the so-called collocation points in the parameter space (Tatang et al., 1997). A few key details of PCE and PCM are reported in Appendix A for completeness. In a previous study on the performance of PCKF (Zeng and Zhang 2010; Zeng et al., 2011), both the parameters and state variables collected in the state vector (3) are approximated by the PCE, and the coefficients of the PCE need to be subsequently updated for each system realization at the selected collocation points as new data become available and are assimilated in the model. When translated to preposterior data-worth analysis, this approach would imply multiple reconstructions/updates of the PCE, consistent with each realization of **C**, and would lead to unsustainable computational cost.

Here, we introduce a modified version of the PCKF to improve the computational efficiency of the proposed data-worth analysis scheme. As we illustrate in the following, our approach to PCKF relies on constructing the PCE-based model proxy only once by (a) evaluating the PCE coefficients and (b) considering the (uncertain) random quantities upon which the PCE-based model proxy relies as the quantities to be updated during the data assimilation process (rather than the PCE coefficients). This approach to PCKF enables us to employ the proxy model directly in the context of the required repeated model inversions. Thus, it obviates the need for multiple evaluations of the original (full system) model, which is critical to improve computational efficiency in the preposterior data worth stage. The procedure requires having at our disposal as prior information the key geostatistical descriptors of the parameter field, such as mean,

covariance structure and integral scale. If the prior information content is insufficient to this end, one can in principle resort to an iterative PCE-based bootstrap filter, along the lines proposed by Oladyshkin et al. (2013) to enhance the reliability of the proxy model. Given the demonstration-oriented nature of our test cases described in Section 3.2, we do not pursue an intensive study of the latter strategy.

At each time step, the random vector Δ is characterized by a set of N_p (statistically independent) random variables (collected in vector). The following PCE approximation of Δ can be constructed:

$$\hat{\boldsymbol{\Delta}} = \sum_{n \le d} a_n \Gamma_n \left(\boldsymbol{\xi}_1, \boldsymbol{\xi}_2, ..., \boldsymbol{\xi}_{N_p} \right)$$
 (12)

Here, $\Gamma_n(\xi_1, \xi_2, ..., \xi_{N_p})$ is an *n*-th order multivariate orthogonal polynomial of the variables $\xi_1, \xi_2, ..., \xi_{N_p}$; a_n is a deterministic PCE coefficient; and d is the highest order of the expansion. At each time step, the coefficients in (12) can be obtained via the PCM. Before data are collected (that is, in the absence of conditioning on the future data), we compute the coefficients in (12) by running the system model for a simulation time encompassing all future time steps of interest and constructing a proxy for the whole simulation period. To do so, the PCM requires one to perform only a number:

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$$M = \frac{(N_p + d)!}{N_p!d!}$$
 (13)

of runs of the full system model, which corresponds to the number of selected collocation points for $(\xi_1, \xi_2, ..., \xi_{N_p})$.

After the construction of the PCE surrogate model for the whole target simulation period, data assimilation is then applied to sequentially update on the constructed proxy the uncertain model parameters collected in vector $\boldsymbol{\xi}$. For each updated $\boldsymbol{\xi}$, the

corresponding entries of Δ can be obtained by sampling (12) rather than running the full model simulation, which renders data assimilation performed on the PCE proxy much more efficient than relying on the full model. The statistics of Δ , i.e., mean and variance at various space and time points can then be evaluated by sampling a collection of realizations of ξ .

The selection of the order of PCE is a core step in the application of PCM. Note that: (a) increasing the order of the PCE through the data assimilation process might improve the accuracy of the surrogate model but can also require a considerably larger number of collocation points, at the expense of computational efficiency; and (b) the accuracy of the statistics obtained at a given PCE order cannot be assessed until different orders are actually implemented. As suggested by Dai et al. (2014), we assess the quality of the constructed proxy through a blind test. The latter is based on the comparison between the results obtained through the proxy and the original (full) system model with the same parameter sets, the latter being randomly selected in the parameter space at locations which differ from those of the collocation points.

The data-worth analysis based on the PCKF is essentially developed according to the workflow illustrated in Section 2.1. The only difference lies in the appearance of an additional step at the prior data-worth analysis stage, in which the coefficients of the PCE surrogate model are calculated through PCM, in addition to the evaluation of hypothetical additional data on the N_{e1} system realizations. The preposterior stage can then be implemented efficiently with the constructed surrogate model. As a result, the total number of full model runs to be performed in the data-worth analysis via the PCKF

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$$N = N_{el} + M = N_{el} + \frac{\left(N_p + d\right)!}{N_p! d!}$$
 (14)

394 3. Illustrative Examples

3.1 Governing equations

A two-dimensional contaminant transport scenario taking place in a synthetic heterogeneous two-dimensional aquifer is considered to: (a) illustrate the feasibility of the proposed PCKF-based data-worth analysis; and (b) analyze key elements of alternative sampling strategies and conditions. We assume that conservative solute transport can be described by:

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$$\frac{\partial(\theta C)}{\partial t} = \nabla(\theta \mathbf{D}_d \nabla C) - \nabla(\theta \mathbf{v}C) + q_s C_s \delta(\mathbf{x} - \mathbf{x}_0)$$
 (15)

with appropriate initial and boundary conditions. In (15), θ is porosity; C is solute concentration (ML⁻³); t is time (T); $\mathbf{x} = (x_1, x_2)$ is vector location in two-dimensional space (L); $\mathbf{v} = (v_1, v_2)$ (LT⁻¹) is seepage velocity vector; q_s (L³T⁻¹) and C_s (ML⁻³) respectively are volumetric flux and solute concentration of a point source positioned at \mathbf{x}_0 ; δ is the Dirac function; and \mathbf{D}_d (L²T⁻¹) is the dispersion tensor, which is defined as:

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$$\begin{cases} \mathbf{D}_{d11} = (\alpha_L v_1^2 + \alpha_T v_2^2) / |\mathbf{v}| + D_m \\ \mathbf{D}_{d11} = (\alpha_L v_1^2 + \alpha_T v_2^2) / |\mathbf{v}| + D_m \\ \mathbf{D}_{d12} = \mathbf{D}_{d21} = (\alpha_L - \alpha_T) v_1 v_2 / |\mathbf{v}| \end{cases}$$
(16)

where D_m (L²T⁻¹) is the diffusion coefficient; |v| (LT⁻¹) is magnitude of velocity; and α_L and α_T (L) are the longitudinal and transverse dispersivity, respectively.

In our study, we consider that solute migrates within a steady-state saturated flow,

412 described as:

$$\nabla \left(\mathbf{k} \frac{\rho}{\mu} \nabla h \right) = 0 \tag{17}$$

$$\mathbf{v} = -\mathbf{k} \frac{\rho}{\theta \mu} \nabla h \tag{18}$$

subject to boundary conditions:

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$$h(\mathbf{x}) = H(\mathbf{x}), \quad \mathbf{x} \in \Gamma_D; \quad -\mathbf{q}(\mathbf{x}) \cdot \mathbf{n}(\mathbf{x}) = Q(\mathbf{x}), \quad \mathbf{x} \in \Gamma_N$$
 (19)

Here, \mathbf{k} (L²) is permeability tensor, which we assume to be isotropic; ρ and μ are fluid density and viscosity, respectively; $\mathbf{q}(x) = \mathbf{v}(x)/\theta$ is the Darcy flux vector at x; H(x) is prescribed head on Dirichlet boundary segment Γ_D ; Q(x) is prescribed flux across Neumann boundary segment Γ_N ; and $\mathbf{n}(x)$ is the outward unit normal to the boundary $\Gamma_D \cup \Gamma_N$. Numerical solution of (15)-(19) is performed through a finite difference method.

3.2 Model setup

The key parameters of the problem are listed in Table 1, where all quantities are given in consistent length and time units. We consider an aquifer with a constant porosity θ = 0.15, which is discretized along the horizontal plane into 40 × 40 grid cells of uniform size. Flow conditions are designed upon setting prescribed head boundary values of 12 and 7, respectively, on the left and right sides of the domain, while keeping impervious upper and bottom boundaries. The (natural) logarithm of permeability is considered to be a (second-order stationary) Gaussian (correlated) random field. In this study, we assume that the key statistical attributes of the random log permeability field are known. We take the covariance function of log permeability (Zhang, 2002) as:

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$$Cov(\mathbf{x}; \mathbf{y}) = \sigma^2 \exp \left[-\frac{|x_1 - y_1|}{\eta_1} - \frac{|x_2 - y_2|}{\eta_2} \right]$$
 (20)

Here, $\mathbf{x} = (x_1, x_2)$ and $\mathbf{y} = (y_1, y_2)$ are two spatial locations; σ^2 is log permeability 434 variance, which is set to unity in our demonstration study (see Table 1); and η_1 and 435 η_2 are the correlation lengths parallel to x_1 and x_2 directions, respectively, which we 436 set as $\eta_1 = \eta_2 = 80$ (Table 1), thus resulting in a computational domain which spans 437 5 correlation scales along the x_1 and x_2 directions. We parameterize the random 438 permeability field through the Karhunen-Loeve (KL) expansion: 439

$$\ln(k(x)) = \langle \ln(k) \rangle + \sum_{n=1}^{\infty} \sqrt{\lambda_n} f_n(x) \xi_n$$
 (21)

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where $\langle \ln(k) \rangle$ is the mean of log permeability, which is set to 5 in this study (Table 1); λ_n and $f_n(x)$ respectively are deterministic eigenvalues and eigenfunctions of 442 the covariance function; and ξ is the vector of (zero-mean and unit variance) Gaussian 443 random variables, ξ_n , which constitutes the set of uncertain model parameters that we 444 consider. In practice, expansion (21) is often truncated up to the first N_p dominant terms. 445 The true (or reference) log permeability field is an unconditional random realization 446 obtained via the geostatistical software library (GSLIB) (Deutsch and Journel, 1992) 447 and is depicted in Figure 2. A point-wise contaminant source with a uniform (and 448 constant in time) concentration $C_s = 250$ and $q_s = 100$ is placed at $x_0 = (20, 200)$. 449 Diverse arrays of monitoring wells are introduced in the system corresponding to the 450 different scenarios that we investigate, as illustrated in the following. A uniform time 451 step $\Delta T = 30$ is used for the transport simulations. Concentrations in the reference field 452 are sampled at monitoring wells at equally spaced time intervals of duration ΔT . We set 453

the final observation time at $T_{60} = 60 \Delta T$. The goal of the alternative monitoring/sampling strategies that we analyze in our tests is to accurately predict the spatial distribution of the contaminant plume at this final observation time.

We illustrate our approach on a set of test cases, listed in Table 2. Test case TC1 is designed to validate the accuracy of the proposed PCKF through a comparison against the results obtained with the standard EnKF method. Concentration values are sampled at all sixteen monitoring wells depicted in Figure 2 and are used for sequential updates of the uncertain model parameters. The observation time T_{60} is subdivided into 60 temporal intervals of uniform unit length ΔT .

Assimilation of concentration values sampled at the monitoring wells is performed at the end of each time interval ΔT , starting at time $T_0 = 0$ and up to $T_{60} = 60 \Delta T$. Prior concentration data in all of our test cases are assumed to be associated with errors as large as 10% of their observed values.

Test cases TC1-TC8 are designed to investigate the effect of diverse factors on data-worth analysis. TC1 is set as the base case. It relies on prior data collected at all wells for the first 24 assimilation times (i.e., up to time $T_{24} = 24 \Delta T$). The worth of additional data which are collected at the same wells with uniform (unit) frequency for the remaining assimilation intervals (up to $T_{48} = 48 \Delta T$) is then quantified both at the preposterior and posterior steps.

Test cases TC2 and TC3 differ from TC1 in the sampling frequencies employed for the data-worth analysis. In these cases, concentration samples are planned to be collected at all 16 wells up to time $T_{60} = 60 \Delta T$ and every $3 \Delta T$ (TC3) or $6 \Delta T$ (TC4).

Test cases TC4, TC5, and TC6 consider augmenting the prior data set through observations taken at diverse sets of wells pertaining to the monitoring network of TC1 (i.e., Wells in Column 1 and 2, Column 3 and 4, or Column 2 and 3, respectively for TC4, TC5 and TC6). Similar to TC1, the prior dataset is formed by the concentrations collected during the first 24 assimilation times. For each of these test cases, the worth of additional measurements observed at the corresponding selected well network with uniform (unit) frequency for the remaining assimilation intervals (up to $T_{60} = 60 \Delta T$) is then quantified both at the preposterior and posterior steps. Test cases TC7 and TC8 differ from TC1 in that they are designed to investigate the influence of the size of the prior database on the worth of additional data collected at the same well locations up to the target final time.

3.3 Results and Discussion

3.3.1 Validation of the PCKF (Test case TC1)

We start by considering test case TC1, which is devoted to the assessment of the performance of the proposed PCKF by comparison against the standard EnKF. The random log permeability is parameterized via the KL expansion (21). To obtain a balance between computational accuracy and cost, we only retain the leading terms in the KL expansion, i.e., these terms with the largest eigenvalues. Figure 3 depicts the magnitude of the eigenvalues associated with the covariance matrix as a function of the number of terms retained in (21). These results indicate a rapid decay of the magnitude of the eigenvalues with increasing number of expansions terms. In this study, the first

30 terms are retained to parameterize the random log permeability field. This modeling choice enables us to account approximately for 70% of the energy of the target spatial random field and to achieve an appropriate balance between computational efficiency and accuracy of the overall procedure in the settings we analyze. A discussion on how to select the number of the retained terms can be found in Chang and Zhang (2009). We then select a second-order PCKF, rendering a number of M = (30 + 2)! / (30! 2!) = 496collocation points at which the full system model is required to be evaluated for the construction of the PCE of the concentration, which is considered as the target system state. We select second-order PCKF for two reasons: (a) Li and Zhang (2007) conclude that PCEs of even order have a superior performance when compared against PCEs of the subsequent (odd) order when the unknown system parameters are Gaussian; and (b) employing a fourth-order PCKF would require a total of 46,376 full model evaluations to construct the PCE approximation, thus rendering this option unfeasible to pursue. Concentration measurements are generated upon solving the flow and transport problems on the reference permeability field and sampling at all the wells from time T_0 = 0 to T_{60} = 60 ΔT with uniform sampling spacing equal to ΔT .

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For the purpose of our comparison, both the PCKF and the EnKF are implemented with a number of realizations equal to 300. Figure 4 depicts the true concentration field, and the (ensemble) mean plume obtained via the PCKF and the EnKF at time T_{60} . These results suggest that the PCKF and the EnKF render mutually consistent predictions of the spatial distribution of the reference concentrations at the target time in our setting, even as some small differences can be noted between estimated and true distributions,

which can also be related to the lack of information associated with the region within which the contaminant has not been observed. Figure 5 depicts the spatial distribution of the predictive variance of concentrations obtained via the PCKF and the EnKF at time T_{60} .

We note that the overall quality of the comparison might be influenced by issues related to filter inbreeding, which can be associated with the limited number of realizations that we employ. However, employing a considerably larger collection of realizations would render unfeasible the application of the standard EnKF from a computational standpoint. For example, the estimated CPU time of the implementation of EnKF with 3,000 realizations is 78,300 s, i.e., 21.75 h. With this in mind, and noting that repeated runs of the full system model are required at the preposterior stage, we consider that the result that we obtain can imbue us with a relatively high confidence in our selection of the reduced PCKF approach.

3.3.2 Effect of the sampling frequency of planned future measurements (Test cases TC1, TC2, and TC3)

Sampling frequency is a critical factor in a monitoring strategy design. While high sampling frequency can provide an increased flux of data that may help to reduce predictive uncertainty, it will also yield an increased workload and financial demands. The purpose of the comparison of TC1, TC2, and TC3 is to identify the relative worth of diverse sampling frequencies. In these cases, the prior data set **D** comprises the measurements taken at the monitoring wells in Column 1 to 4 during the first 24

assimilation times (i.e., from time $T_1 = \Delta T$ up to $T_{24} = 24 \Delta T$). Three hundred realizations of the additional data set are used at the prior stage of the analysis, while the preposterior analysis is performed through the PCKF due to its notable efficiency.

As stated above, Figure 5a depicts the spatial distribution at time T_{60} of the predictive variance of the contaminant concentration, conditional only on \mathbf{D} , i.e., $Var(\Delta | \mathbf{D})$. We note that the predictive uncertainty is larger at the front of the plume than at locations in the vicinity of the source. This result is consistent with the high spatial concentration gradients in the proximity of the plume front (*Rubin*, 1991).

Figure 6 depicts comparisons between concentration values observed at four wells located at (5, 5), (15, 15), (25, 15) and (35, 25) at times $T_i = i \Delta T$ (i = 25, 26, 27, ..., 48) and corresponding estimates of mean concentrations computed on the basis of the prior **D** values. Intervals of width corresponding to ± 1 standard deviations about the mean are depicted to complete the picture.

These results show that, even as in some cases the true concentration values fall within the limits of the considered uncertainty bounds, predicted values tend to systematically deviate from their true counterparts (albeit with different degrees of discrepancy, depending on the monitoring well location) with increasing time, as expected. We recall that estimates of mean concentration together with the associated predictive variance constitute the bases for the construction of the additional data sets \mathbf{C} to be employed in the preposterior analysis step for each of these test cases. A posterior analysis is then performed after \mathbf{C}' becomes available.

Data-worth analysis via the PCKF is applied to each test case to assess the worth of the additional data sets C which are built as illustrated in Section 3.2. Figures 7a, c, and e depict the spatial distribution of the expected concentration variance reduction $Var_{CD}E(\Delta \mid \mathbf{D}, \mathbf{C})$, respectively for TC1, TC2, and TC3, calculated at the preposterior stage. Figures 7b, d, and f depict the spatial distribution of concentration variance reductions, $Var(\Delta | \mathbf{D}) - Var(\Delta | \mathbf{D}, \mathbf{C}')$, calculated at the posterior stage. It can be noted that the spatial distributions of the expected variance reductions resemble closely their reference (posterior) counterparts. The scalar measures of the expected and reference data-worth are compared in Figure 8. For ease of reference, these values are also listed in Table 3 together with the corresponding results associated with all test cases examined. From these results, we conclude that the expected data-worth can estimate its reference counterpart with high accuracy. The preposterior data-worth metric $Tr \lceil Cov_{CID} E(\Delta \mid \mathbf{D}, \mathbf{C}) \rceil$ of TC1 is 215,742 and is larger than that associated with TC2 and TC3, these being equal to 125,292 and 200,672, respectively. These results support the idea that an increased sampling frequency should lead to uncertainty reduction. However, our results also reveal that the rate of increase of the preposterior data-worth somehow decreases with increasing sampling frequency. This observation, in turn, implies that the effect of an increased sampling frequency becomes less significant when the frequency has reached a certain threshold level. The latter should depend on the spatial location of the planned monitoring well, as well as on the underlying aquifer heterogeneity and transport setting.

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A remarkable advantage of our PCKF-based data-worth analysis is the

computational efficiency. Each of the test cases here analyzed requires only 796 forward runs of the full system model. These include: (a) 300 simulations for the estimation of the statistics (mean and covariance) of the additional data set to be included in \mathbb{C} ; and (b) 496 simulations to construct the surrogate model. The CPU time for the data-worth analysis with PCKF for TC1, TC2, and TC3 respectively are 9,884 s, 3,884 s, and 5,084 s. Otherwise, employing EnKF based on a collection of $N_{e1} = N_{e2} = 300 \,\mathrm{MC}$ realizations for data-worth analysis requires a number of full model solutions equal to 90,000 for each assimilation step, with estimated CPU times of 1,879,200 s, 1,096,200 s, and 1,252,800 s, respectively for TC1, TC2, and TC3.

3.3.3 Effect of the monitoring network location (Test cases TC4, TC5, and TC6)

Test cases TC4, TC5, and TC6 are identical to TC1 except that additional sampling after collection of the prior data set (i.e., the data set composed by observations collected for the first 24 time intervals) is performed at locations listed in Table 2 and identified in Figure 2. The additional data set C is then formed by the data observed at the wells in Column 1 and 2, Column 3 and 4, and Column 2 and 3 for TC4, TC5 and TC6, respectively, from assimilation times $T_{25} = 25 \Delta T$ to $T_{48} = 48 \Delta T$ with sampling frequency equal to ΔT . Figure 9 depicts the spatial distributions of the expected preposterior concentration variance reductions (Figure 9a, c, and e) and the reference posterior concentration variance reduction (Figures 9b, d, and f) at the latest time T_{60} . Only minute deviations are observed between the patterns of preposterior estimations and their posterior counterparts. Scalar measures of preposterior and posterior dataworth are also depicted in Figure 10. As compared to TC1 (for which

 $Tr[Cov_{CID}E(\Delta \mid \mathbf{D}, \mathbf{C})] = 215,742$), reducing the monitoring well network brings in an overall decrease of the predictive concentration variance reduction, as expected, the scalar measures of preposterior data-worth for TC4, TC5, and TC6 respectively being 126,698, 190,141, and 207,585. These results suggest that acquisition of additional data from monitoring wells in Column 2 and 3 has the largest potential for providing valuable information in the setting we analyze. This is consistent with the observation that wells in Columns 2 and 3, and particularly those in Column 3, are located in a region where the prior predictive uncertainty is considerably high, as opposed to wells in Column 1 and 4, which are placed at locations with smaller predictive uncertainties. Even as a generalization of these results is hard to propose and the specific choices associated with the selection of new monitoring wells can be associated with additional constraints of diverse nature (including, e.g., political and socio-economical elements), our results support the idea that an effective choice would be to acquire new data at wells located within a region where the prior predictive uncertainty is relatively large.

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3.3.4 Effect of prior data content (Test cases TC1, TC7, and TC8)

Prior data content is a critical element that may impact the results of data-worth analysis. In this sense, TC7 and TC8 are identical to TC1 except that the prior data vector \mathbf{D} is formed by diverse amounts of sampled data. In TC7 we collect prior data from time T_1 to time $T_{12} = 12 \Delta T$, while TC8 considers prior observations sampled from time T_1 to time $T_{18} = 18 \Delta T$. The spatial distributions of prior predictive concentration variance at time T_{60} are depicted in Figure 11a and b, respectively for TC7 and TC8. As expected, the prior predictive concentration variances of TC7 and TC8 are generally

larger than that of TC1 because of the diminished prior information content. Figures 12a and b, and Figures 12c and d depict the comparison of spatial patterns of the preposterior and reference posterior predictive uncertainty reductions for TC7 and TC8, respectively. The preposterior patterns approximate their posterior counterparts with a sufficiently high quality in both cases. Figure 13 depicts the scalar measures of (preposterior and posterior/reference) data-worth for these test cases. The scalar measures of preposterior data-worth are 530,745 and 330,082, respectively for TC7 and TC8, to be compared against the value of 215,742 for TC1.

As expected, a reduced number of prior data yields an increased benefit arising from collecting additional data. Due to the limited amount of prior information, additional data appear to be associated with a relatively high value.

4. Conclusions

Our work leads to the following major conclusions.

- 1. Integrating a Probabilistic Collocation Method within EnKF allows assessing dataworth through a surrogate of the full system model. This leads to a remarkable improvement of the computational efficiency of the data-worth procedure. This is particularly relevant considering that, even as incorporating the EnKF into dataworth analysis frameworks enables one to quantify the worth of additional data in a sequential way, routine applications are still challenging due to large computational costs involved.
- 2. We illustrate our approach on a suite of synthetic scenarios involving conservative solute migration in a two-dimensional random permeability field. Our test cases are

designed to: (a) test the accuracy of our approach when compared to an EnKF dataworth analysis based on Monte Carlo solutions of the full system model (TC1); and (b) investigate the effect of diverse factors on data-worth analysis (TC1-TC8), including design components of an envisioned monitoring scheme, such as spatial location of monitoring wells, temporal sampling frequency, and prior data content for solute concentrations. When compared against results obtained via the full system model, our Probabilistic Collocation EnKF (PCKF) renders significantly accurate results at a considerably reduced computational time in the setting considered, in terms of predicted concentration values and associated uncertainty quantification.

- 3. In our examples (TC1, TC2, and TC3), augmenting the temporal sampling frequency associated with planned acquisitions of additional concentration data leads to increased benefits in terms of uncertainty reduction. The effects of the increased sampling frequency tend to become less significant when the assimilation frequency has reached a certain threshold level.
- 4. Foreseeing the design of the location of a monitoring well within a region where prior predictive uncertainty is highest is more likely to provide valuable data (TC4, TC5, and TC6). Our examples (TC1, TC7, and TC8) suggest that the value of additional data is highest when these are supplemented to the smallest sets of prior data. The scalar measure of data-worth shows a sharp decrease with the increase of the number of measurements forming the prior dataset.

Appendix A

The PCE approximation $\hat{\Delta}$ of the random vector Δ that is characterized by a set of N_p (statistically independent) random variables (collected in vector ξ) relies on the spectral expansion:

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$$\hat{\Delta} = \sum_{n \le d} a_n \Gamma_n \left(\xi_1, \xi_2, ..., \xi_{N_p} \right)$$
 (A. 1)

Here, d is the highest order of the expansion; a_n are deterministic coefficients; and $\Gamma_n\left(\xi_1,\xi_2,...,\xi_{N_p}\right)$ are the n-th order multi-dimensional orthogonal polynomials in terms of random variables $\xi_1,\xi_2,...,\xi_{N_p}$ (collected in vector $\boldsymbol{\xi}$), which constitute the input to the system model. If these uncertain parameters are Gaussian, the Hermite polynomials form the appropriate orthogonal basis for Gaussian random variables (Xiu and Karniadakis, 2002).

The number of unknown coefficients in (A.1) is N_c :

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$$N_{c} = \frac{\left(N_{p} + d\right)!}{N_{p}!d!}$$
 (A. 2)

Hence, N_c linear independent equations are required to solve the coefficients.

Calculation of the PCE coefficients can be accomplished through the Probabilistic collocation method (PCM). Let us define the residual R between the real output Δ and its approximation $\hat{\Delta}$ as:

$$R(\lbrace a_i \rbrace, \xi) = \hat{\Delta} - \Delta \tag{A. 3}$$

where $\{a_i\}$ is the set of the PCE coefficients.

The residual (A. 3) should satisfy the following integral equation:

$$\int R(\lbrace a_i \rbrace, \xi) \delta(\xi - \xi_j) P(\xi) d\xi = 0$$
 (A. 4)

where δ is the Dirac delta function; and ξ_j is a sub-set of ξ whose entries are the collocation points. Equation (A.4) results in a set of independent equations whose solution yields the coefficients of the PCE.

Selection of the collocation points is the key issue of PCM. *Li and Zhang* (2007) suggest that the collocation points can be selected from the roots of the next higher order orthogonal polynomial for each uncertain parameter. Once the coefficients of the polynomial chaos expansions are obtained, (A.1) can be employed as a proxy for the original full system model. The statistics of Δ can be evaluated by the PCE coefficients (Li and Zhang, 2007; Li et al., 2009) or by sampling on the surrogate model (A.1) (Li et al., 2011 and Dai et al., 2014) with a group realizations of random variables.

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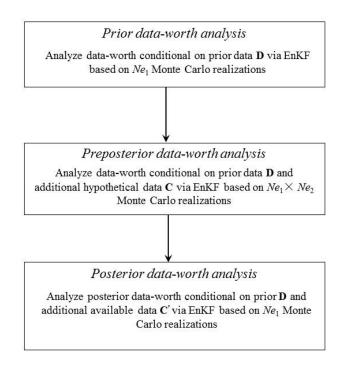


Figure 1. Workflow of data-worth analysis for a dynamic system.

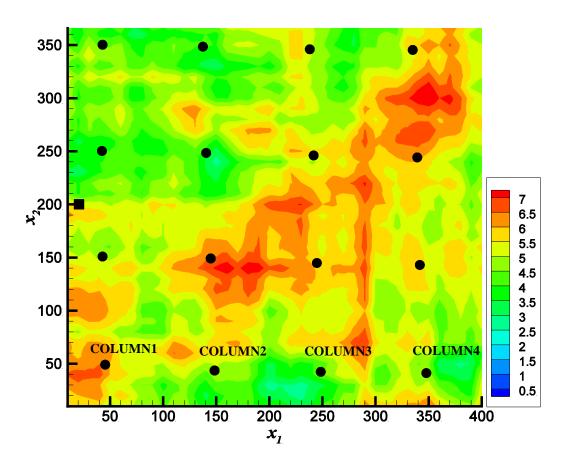


Figure 2. The true (or reference) log permeability field. Solid black square indicates the contaminant source; solid circles denote monitoring wells employed in the test cases examined.

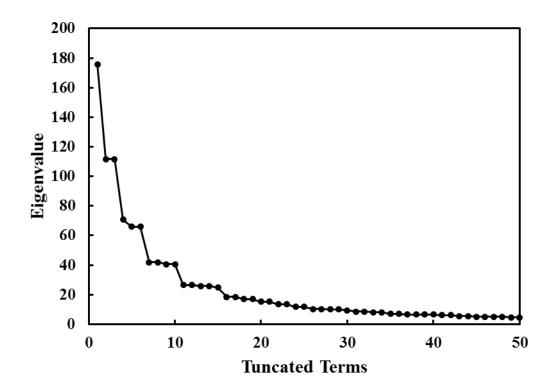
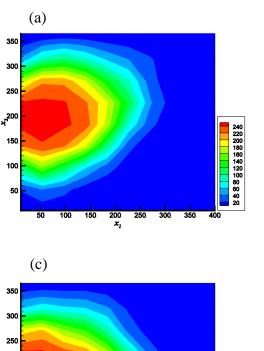
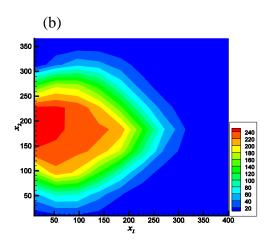


Figure 3. Decay of eigenvalues as function of the number of terms retained in the Karhunen-Loeve (KL) expansion (21).





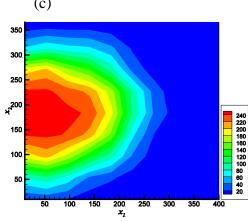
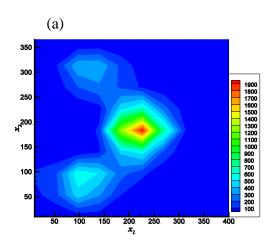


Figure 4. (a) Spatial distributions of true (reference) concentration field and (b) mean concentration field obtained via (b) the PCKF, and (c) EnKF at time T_{60} .



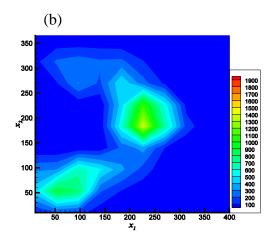


Figure 5. Spatial distribution of the predictive variance of solute concentrations obtained via (a) the PCKF and (b) the EnKF at time T_{60} .

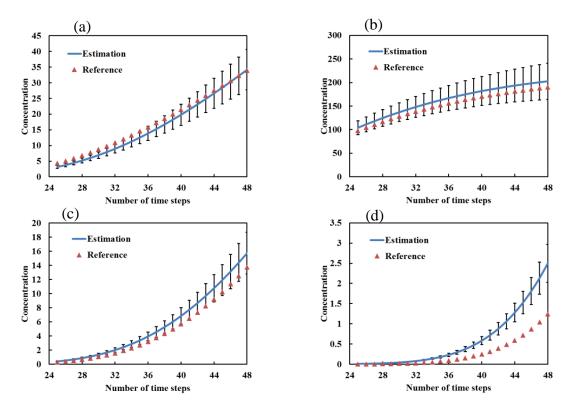


Figure 6. Solute concentration values observed at observation wells located at (a) (5, 5); (b) (15, 15); (c) (25, 15); and (d) (35, 25) at times $T_i = i \Delta T$ (i = 24, 25, 26, ..., 48) with corresponding estimates of mean concentrations and envelopes of width of ± 1 standard deviations about the mean obtained based only on prior **D** values.

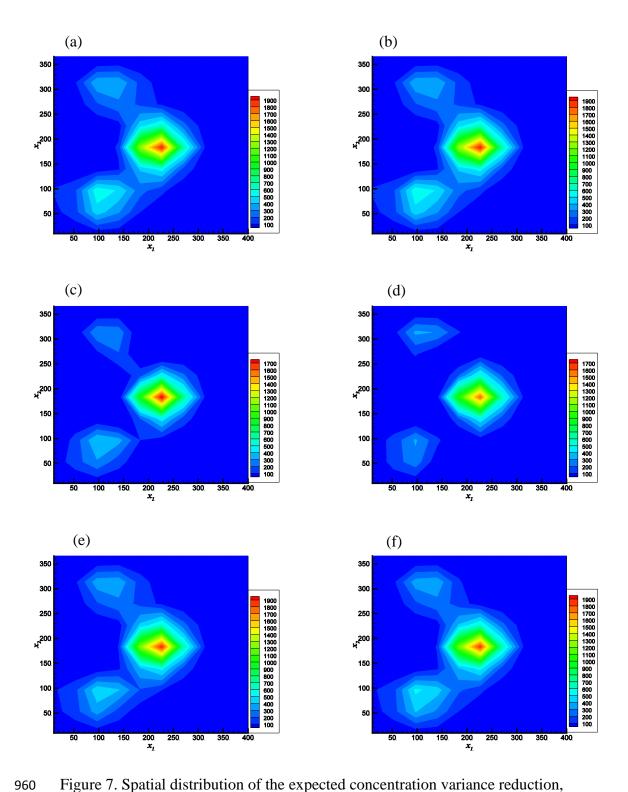


Figure 7. Spatial distribution of the expected concentration variance reduction, $Cov_{C|D}E(\Delta \mid \mathbf{D}, \mathbf{C})$, for (a) TC1; (c) TC2; and (e) TC3. Spatial distribution of concentration variance reduction, $Var(\Delta \mid \mathbf{D}) - Var(\Delta \mid \mathbf{D}, \mathbf{C}')$, obtained after \mathbf{C}' has been observed for (b) TC1; (d) TC2; and (f) TC3. Results are depicted for T_{60} .

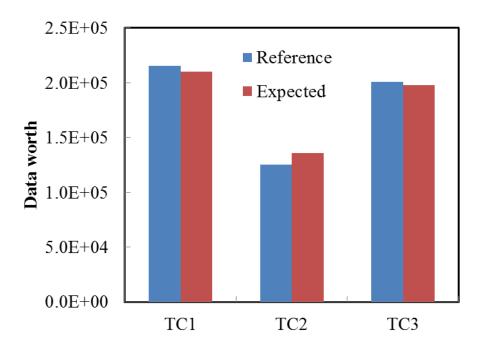


Figure 8. Scalar measures of the expected and reference data-worth for TC1, TC2, and TC3 at T_{60} .

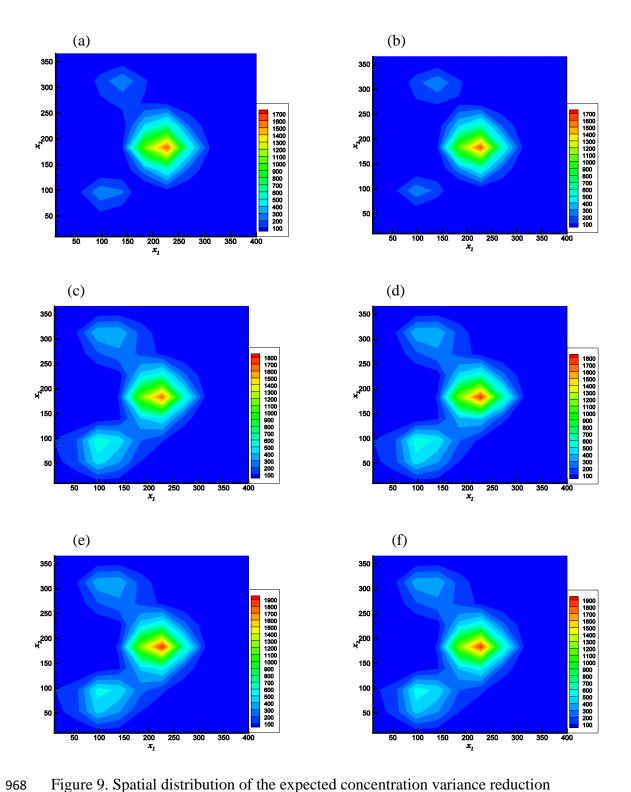


Figure 9. Spatial distribution of the expected concentration variance reduction $Cov_{C|D}E(\Delta \mid \mathbf{D}, \mathbf{C})$ for (a) TC4; (c) TC5; and (e) TC6. Spatial distribution of concentration variance reduction, $Var(\Delta \mid \mathbf{D}) - Var(\Delta \mid \mathbf{D}, \mathbf{C}')$ obtained after \mathbf{C}' has been sampled for (b) TC4; (d) TC5; and (f) TC6. Results are depicted for T_{60} .

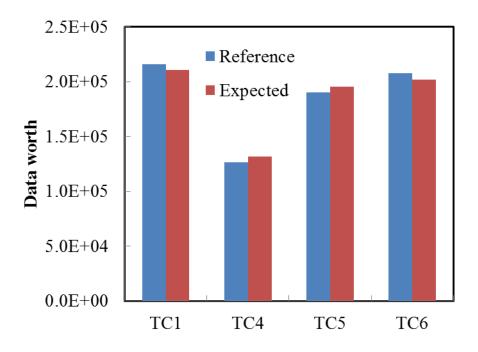


Figure 10. Scalar measures of the expected and reference data-worth for TC1, TC4, TC5, and TC6 at T_{60} .

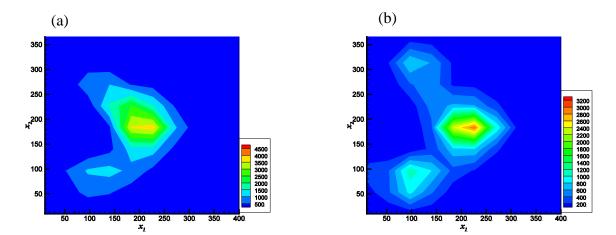


Figure 11. Spatial distribution of the predictive variance of contaminant concentration at time T_{60} conditional on \mathbf{D} , $Var(\Delta \mid \mathbf{D})$, for (a) TC7; and (b) TC8. Results are depicted for T_{60} .

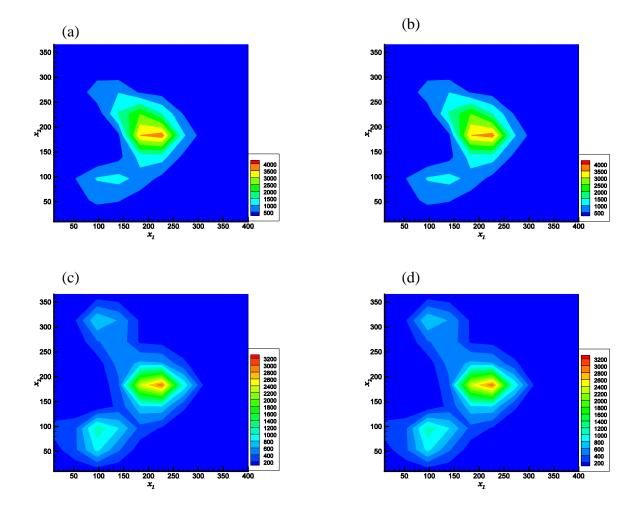


Figure 12. Spatial distribution of the expected concentration variance reduction, $Cov_{C|D}E(\Delta \mid \mathbf{D}, \mathbf{C})$, for (a) TC7; and (c) TC8. Spatial distribution of concentration variance reduction, $Var(\Delta \mid \mathbf{D}) - Var(\Delta \mid \mathbf{D}, \mathbf{C}')$, obtained after \mathbf{C}' has been observed for (b) TC7; and (d) TC8. Results are depicted for T_{60} .

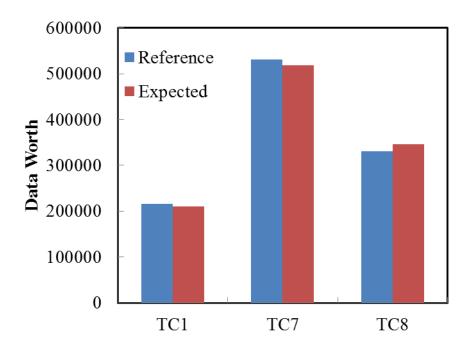


Figure 13. Scalar measures of the expected and reference data-worth for TC1, TC7 and TC8 at T_{60} .

Table 1. Summary of the system setup parameters employed in the two-dimensional synthetic examples.

Parameter	Value (in consistent units)			
Discretization and source ch	aracteristics			
No. of rows in the domain	40			
No. of columns in the domain	40			
Domain width	400			
Domain length	400			
Time step	30			
Solute source location $x_0 = (x_{01}, x_{02})$	(20, 200)			
Volumetric flux at source (q_s)	100			
Concentration at source (C_s)	250			
Flow and Transport Parameters				
Head along upstream boundary	12			
Head along downstream boundary	7			
Diffusion coefficient D_m	0.5			
Local-scale dispersivities	(1.5, 0.3)			
Porosity	0.15			
Log Permeability				
Mean value	5			
Variance	1			
Correlation Length ($\eta_1 = \eta_2$)	80			

Table 2. Main characteristics of test cases examined.

Test	No. of time		Sampling	
	steps in prior	Monitoring well	frequency of	Description
case	data D		additional data	
TC1	24	COLUMN1-4	ΔT	Base case
TC2	24	COLUMN 1-4	$6 \Delta T$	Effect of sampling frequency
TC3	24	COLUMN 1-4	$3 \Delta T$	Effect of sampling frequency
TC4	24	COLUMN 1,2	ΔT	Effect of monitoring network location
TC5	24	COLUMN 3,4	ΔT	Effect of monitoring network location
TC6	24	COLUMN 2,3	ΔT	Effect of monitoring network location
TC7	18	COLUMN 1-4	ΔT	Effect of prior data content
TC8	12	COLUMN 1-4	ΔT	Effect of prior data content

Table 3. Summary of the scalar measures of preposterior and posterior uncertainty reductions for all test cases (results are listed for T_{60}).

Test case	Preposterior predictive uncertainty reduction	Posterior predictive uncertainty reduction
TC1	215,742	210,452
TC2	125,292	135,982
TC3	200,672	197,962
TC4	126,698	131,980
TC5	190,141	195,533
TC6	207,585	201,750
TC7	530,745	518,494
TC8	330,028	345,770

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