A two-stage ensemble Kalman filter based on multiscale model reduction for inverse problems in time fractional diffusion-wave equations

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ABSTRACT

Ensemble Kalman filter (EnKF) has been widely used in state estimation and parameter estimation for the dynamic system where observational data is obtained sequentially in time. Very burdened simulations for the forward problem are needed to update a large number of EnKF ensemble samples. This will slow down the analysis efficiency of the EnKF for large-scale and high dimensional models. To reduce uncertainty and accelerate posterior inference, a two-stage ensemble Kalman filter is presented to improve the sequential analysis of EnKF.

It is known that the final posterior ensemble may be concentrated in a small portion of the entire support of the initial prior ensemble. It will be much more efficient if we first build a new prior by some partial observations, and construct a surrogate only over the significant region of the new prior. To this end, we construct a very coarse model using generalized multiscale finite element method (GMsFEM) and generate a new prior ensemble in the first stage. GMsFEM provides a set of hierarchical multiscale basis functions supported in coarse blocks. This gives flexibility and adaptivity to choosing degree of freedoms to construct a reduce model. In the second stage, we build an initial surrogate model based on the new prior by using GMsFEM and sparse generalized polynomial chaos (gPC)-based stochastic collocation methods. To improve the initial surrogate model, we dynamically update the surrogate model, which is adapted to the sequential availability of data and the updated analysis. The two-stage EnKF can achieve a better estimation than standard EnKF, and significantly improve the efficiency to update the ensemble analysis (posterior exploration).

To enhance the applicability and flexibility in Bayesian inverse problems, we extend the two-stage EnKF to non-Gaussian models and hierarchical models. In the paper, we focus on the time fractional diffusion-wave models in porous media and investigate their Bayesian inverse problems using the proposed two-stage EnKF. A few numerical examples are carried out to demonstrate the performance of the two-stage EnKF method by taking account of parameter and structure inversion in permeability fields and source functions.

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1 Introduction

The model inputs (parameters, source, domain geometry and system structure, et. al.) in many practical systems are often unknown. We need to identify or estimate these inputs by partial and noisy observations to construct predictive models and calibrate the models. This results in inverse problems. In the paper, we consider the inverse problems in anomalous diffusion models. The anomalous diffusion can be roughly classified into two categories: subdiffusion \((0 < \gamma < 1)\) and superdiffusion \((1 < \gamma < 2)\). Here \(\gamma\) denotes the fractional derivative with respect to time. The anomalous diffusion equations are also called fractional diffusion-wave equations. Theoretical results such as existence, uniqueness, stability and numerical error estimates are presented in [19] for some type of anomalous diffusion equations. The relationship between anomalous diffusion equations and regular diffusion equations is discussed in [28]. The fractional order diffusion-wave equation as a typical fractional partial differential equation [35], is a generalization of the classical diffusion and wave equation and can be used to better characterize anomalous diffusion phenomena in various fields. The fractional diffusion-wave equations can model porous media applications, viscoelastic mechanics, power-law phenomenon in fluid and complex network, allometric scaling laws in biology and ecology, quantum evolution of complex systems and fractional kinetics [27].

In practice, the inputs and parameters in the anomalous diffusion models are often unknown and need to be identified based on some observation data and prior information. The problem of identifying unknown inputs in mathematical models has been intensively studied in the framework of inverse problems and various numerical methods have been developed [1, 20, 26]. The mathematical model of inverse problem is featured with quantities which renders useful simulation prediction obtained by imperfect model equations and measurements. The inverse problem is usually ill-posed. Many methods [17, 11] such as regularization or penalty can be used to overcome the ill-posedness. The unknown inputs (e.g., permeability field) of the anomalous diffusion models in porous media may have multiscale structure, complex geometry patterns and uncertainty. This significantly increases the challenge of the inverse problems for these models.

Practical models usually involve uncertainty. Moreover, the prior information for unknown parameters and observations are often characterized by random variables. Thus, it is desirable to treat the computational model and its inverse problem in statistical perspective. Once of statistical approach for inverse problems is Bayesian inference. The Bayesian approach [17, 33] incorporates uncertainties in noisy observations and prior information, and can derive the posterior probability density of the parameters, which enables us to quantify the uncertainty in the parameters. The popular sampling methods in Bayesian inversion are Markov chain Monte Carlo (MCMC) [10] method and its variants [2, 22], which require costly computation to achieve convergence and explore the whole state space in high dimension sample spaces. MCMC simulation has to run a long enough chain to give an accurate estimate, and entails repeated solutions of the forward model. This leads to great challenge for solving the Bayesian inverse problem.
The ensemble Kalman filter (EnKF) is another Bayesian method. It can be seen as a reduced-order Kalman filter (KF) or a Monte Carlo implementation of KF \cite{Ev88,Mo82}. Since its introduction by Evenson in \cite{Ev88}, EnKF has been applied in many fields such as oceanography, numerical weather prediction, hydrology and petroleum reservoir history matching. EnKF can be used for both data assimilation, where the goal is to estimate model states by incorporating dynamical observation data, and inverse problem, where the objective is estimate unknown parameters appearing in models. Ensemble members of EnKF are forecasted in time by solving forward models and updated by an approximate Kalman filter scheme. EnKF has the significant advantage that its inherent recursive process is adapted to the sequential availability of observation data in dynamical systems. Unlike MCMC, the ensemble samples are updated independently each other in EnKF and it is not necessary to propose samplers in a very tricky manner. Thus, EnKF and other filter methods have attracted much more attention in community of Bayesian inversion \cite{Ev91,Ev03}. For nonlinear models, the linearization of model can be avoided in EnKF using ensemble covariance as an approximation of the posterior error covariance. The EnKF methods provide the first and second moments of random parameter, which are approximated by ensemble mean and ensemble covariance, respectively. Thus, the EnKF algorithm makes Gaussian approximation in a sequential manner. These approximation is accurate for Gaussian prior models. Some insight analysis of EnKF for inverse problems has been made in recent years \cite{Ev07,Al11}. For the non-Gaussian models, a normal-score ensemble Kalman filter is proposed in \cite{Va14}, where the normal-score transformation is applied to transform unknown non-Gaussian parameters to Gaussian and make the parameters follow marginal Gaussian distributions.

As a Bayesian sampling method, EnKF needs to compute the forward problem repeatedly. When the forward model is computationally intensive, such as multiscale models, a direct application of EnKF forecast with full order model would be computationally prohibitive. In order to significantly improve the simulation efficiency, seeking more efficient sampling from the previous posterior and building surrogates of the forward models \cite{Al11,Go15} are necessary to accelerate the EnKF analysis (posterior exploration). Multiscale models can be solved efficiently and accurately by the numerical multiscale methods in a coarse grid instead of resolving all scales in very fine grid. As a numerical multiscale method, Generalized Multiscale Finite Element Method (GMsFEM) \cite{Co13,Co15} can provide a reduce model with a good trade-off between accuracy and computation efficiency. The main idea of GMsFEM is to use a variational formulation to capture the impact of small-scale features on the coarse-scale by using multiscale basis functions. The small-scale information is integrated into multiscale basis functions, which can be used repeatedly for different source terms and boundary conditions of the model \cite{Co13}. GMsFEM has been developed to solve multiscale models with complex multiscale structures and its convergence is independent of the contrastness of the multiscales \cite{Co13}.

In the framework of EnKF, the output of model depends on random parameters. We use generalized polynomial chaos (gPC)-based stochastic collocation methods to propagate prior uncertainty through the forward model in a sequential manner. The gPC stochastic collocation methods require only a few number of uncoupled deterministic simulations with no reformulation of the governing equations of the forward model. We assume that the model’s output is a stochastic field and admits a gPC expansion. Then we select a set of collocation nodes and use least-squares methods to determine the coefficients of the gPC
basis functions. To taking account of the potential sparsity of the gPC expansion, we use $l_1$ regularization to the least-squares problem. This allows using much fewer samples to construct a gPC surrogate model. The idea has been employed in Bayesian inverse problems [15, 16, 21, 37]. To accelerate computation for the $l_1$ regularized least-squares problem, the lagged diffusivity fixed point method is used. The gPC surrogate is usually constructed based on a prior density [15, 21, 37]. However, the posterior is concentrated in a small portion of the entire prior support in many inference problems [16]. In EnKF, the prior is sequentially updated by incorporating new data information. Thus, it may be much more efficient to build surrogates only over the important region of the updated prior than the initial prior support.

In this paper, we propose a two-stage EnKF to take care of the challenges and concerns mentioned above. In the first stage, we construct a coarse GMsFEM model with very few multiscale basis functions, and build a new prior using standard EnKF based on the first partial measurement data information in time. The initial ensemble samples are drawn from the new prior for the second stage of EnKF. By integrating GMsFEM and sparse gPC stochastic collocation method based on the new prior, we build an initial surrogate model for the second stage. Because the ensemble samples are updated by the new analysis of EnKF, this also sequentially updates the prior based on the new ensemble samples. To improve the initial surrogate model, we dynamically update the surrogate model based on the updated prior in each assimilation step. We note that the surrogates are constructed efficiently in EnKF procedure by using GMsFEM and sparse gPC stochastic collocation method. By virtue of building new priors, we exclude the unimportant region of the posterior. We may use some other methods such as ensemble smoother (ES) [5] to build the new prior. In general, ES is used to estimate the parameters and states when simulation models are typically stable functions. To extend the two-stage EnKF to non-Gaussian models, we integrate the proposed EnKF with normal-score transformation to broaden the applicability. The two-stage EnKF is also explored in hierarchical Bayesian inverse problems. This increases flexibility in prior modeling for the Bayesian inference.

The structure of the paper is as follows. We begin with the general framework of EnKF for inverse problems. In Section 3, we focus on the time fractional diffusion-wave models and the surrogate model construction using GMsFEM and sparse gPC. Section 4 is devoted to presenting the two-stage EnKF based on the surrogate model. In Section 5, we present a few numerical examples to illustrate the performance of proposed EnKF with applications of inverse problems for time fractional diffusion-wave equations. Some conclusions and comments are made finally.

2 Ensemble Kalman filter for inverse problems

Let $\mathcal{U}$ be a Hilbert space and $\mathcal{N}$ a generic forward operator on $\mathcal{U}$ for some physical system. We assume that the forward operator describes the relation of parameter $\theta$, state $u$ and source term $f$, i.e.,

$$\mathcal{N}(u; \theta) = f,$$

(2.1)
where \( u \in \mathcal{U}, f \in \mathcal{U}^* \), the dual space of \( \mathcal{N} \). We assume that the solution of the problem has the form

\[
\mathbf{u = X(\theta; f)}.
\]

Let \( \mathbb{H} \) be the observation operator mapping the model state \( u \in \mathcal{U} \) to the observation space \( \mathcal{Y} \)

\[
y = \mathbb{H}(u(\theta)) := \mathbb{H}(\theta) \in \mathcal{Y}.
\]

Let \( \varepsilon \) be additive Gaussian noise for observation. Then the observation data can be expressed by

\[
d = \mathbb{H}(\theta) + \varepsilon.
\]

In the paper, we assume that \( \varepsilon \) is independent of \( \theta \). In practical setting, the observation data is in a finite dimensional space and can be expressed by

\[
d = \mathbb{H}(\theta) + \varepsilon \in \mathbb{R}^{n_d},
\]

where \( n_d \) is the dimension of observations.

## 2.1 Bayesian inference using EnKF

The EnKF was introduced by Evensen [8] as a powerful data assimilation method. Kalman filter is used for sequential update for states in linear dynamical systems and Gaussian distribution. It provides the mean and covariance information of the posterior distribution. When the prior is Gaussian, the filter gets the posterior Gaussian distribution from the joint Gaussian observation and the parameter. But for nonlinear dynamical system, EnKF has been widely used for data assimilation. In the paper, we use EnKF for Bayesian inverse problems. This is a particular application of EnKF in recent years [6, 32].

Given some observation data, we want to estimate the parameter \( \theta \). In Bayesian context, both \( \theta \) and \( d \) are random variables. Thus Bayes rule gives the posterior probability density for \( \theta \) by

\[
p(\theta|d) \propto p(d|\theta)p(\theta),
\]

where \( p(\theta) \) is the prior distribution before the data is observed. The data enter the Bayesian inference through the likelihood function \( p(d|\theta) \).

If the information of observations is incomplete, the covariance of the noise observation may be unknown. For this situation, we need to estimate the covariance of observation noise. Let \( \varepsilon \) be independent and identically distributed (i.i.d.) Gaussian random vector with zero mean and variance \( \sigma^2 \), i.e.,

\[
\varepsilon \sim N(0, \sigma^2 I),
\]

where \( I \) is the \( n_d \times n_d \) identity matrix. Thus the likelihood function \( p(d|\theta) \) obeys the Gaussian distribution. Let \( \| \cdot \| \) be the Euclidean norm and \( \| \cdot \|_P = \| P^{-\frac{1}{2}} \cdot \| \) the weighted norm, where \( P \) is the prior’s covariance matrix. If the prior \( p(\theta) \) is also Gaussian distribution, then

\[
p(\theta|d) \propto \exp \left( -\frac{\| d - \mathbb{H}(\theta) \|^2}{2\sigma^2} - \frac{\| \theta - \theta^b \|^2}{2P} \right), \quad (2.3)
\]
where \( \theta^b \) is the mean of prior (background information). When \( \sigma^2 \) is unknown, \( \sigma^2 \) is a hyperparameter in the hierarchical Bayesian model. Then the corresponding posterior
\[
p(\theta, \sigma^2 | d) \propto p(d | \theta, \sigma^2) p(\theta)p(\sigma^2).
\]
The marginal posterior of \( \sigma^2 \) is
\[
p(\sigma^2 | \theta, d) \propto p(d | \theta, \sigma^2) p(\sigma^2).
\]
Because the likelihood
\[
p(d, \theta | \sigma^2) = \frac{1}{(2\pi \sigma^2)^{n_d/2}} \exp \left( -\frac{\|d - H(\theta)\|^2}{2\sigma^2} \right)
\]
belongs to the inverse-gamma family, the conjugate prior \( p(\sigma^2) \) can be the inverse-gamma distribution
\[
p(\sigma^2) \propto (\sigma^2)^{-(\alpha+1)} e^{\beta/\sigma^2}.
\]
From (2.4) and (2.5), we get
\[
\sigma^2 | \theta, d \sim \text{Inv-gamma}\left(\alpha + \frac{n_d}{2}, \beta + \frac{\|d - H(\theta)\|^2}{2}\right).
\]
As in [9], we choose two numbers \( n_s \) (\( n_s \) often small and between 0.01 and 1) and
\[
\sigma^2_s = \frac{\|d - H(\theta)\|^2}{n_d - n_p},
\]
where \( \theta \in \mathbb{R}^{n_p} \), such that \( \alpha = n_s / 2 \) and \( \beta = \sigma^2_s \alpha \). Once the posterior distribution of \( \theta \) is inferred, we can extract the posterior mean or the maximum a posteriori (MAP) of the unknown parameter \( \theta \). We note that the MAP estimate is equivalent to the solution of a regularized optimization problem. In fact, maximizing the right hand side of (2.3) is equivalent to the minimization problem
\[
\min_{\theta \in \mathbb{R}^{n_p}} \left( \frac{1}{2} (d - H(\theta))^T R^{-1} (d - H(\theta)) + \frac{1}{2} (\theta - \theta^b)^T P^{-1} (\theta - \theta^b) \right),
\]
where \( R = \sigma^2 I \). The hyperparameter \( \sigma^2 \) can be drawn from Inv-gamma distribution of (2.6). When the observation operator \( H \) is linear, Kalman filter (KF) method can be derived from (2.7) by completing the squares on the variable \( \theta \) and gives the following analysis
\[
\begin{cases}
\theta^a = \theta^b + K(d - H(\theta^b)), \\
P^a = (I - KH)P,
\end{cases}
\]
where \( H = \mathbb{H} \) and the Kalman gain \( K \) is given by
\[
K = PH^T (HPH^T + R)^{-1}.
\]
Let \( k \) be an artificial time for the data assimilation in a dynamic system. We denote unknown parameter, hyperparameter and observation data as \( \theta_k \in \mathbb{R}^{n_p} \), \( \sigma_k^2 \in \mathbb{R} \) and \( \mathbf{d}_k \in \mathbb{R}^{n_d} \) at artificial time step \( k \), respectively. Then we define the artificial discrete dynamic system

\[
\begin{align*}
\theta_k &= \theta_{k-1}, \\
\mathbf{d}_k &= \mathbb{H}(\theta_k) + \varepsilon_k, \\
\sigma_k^2 &\sim \text{Inv-gamma}(\alpha + \frac{n_d}{2}, \beta + \frac{\|\mathbf{d}_k - \mathbb{H}(\theta_k)\|^2}{2}).
\end{align*}
\]

In the framework of EnKF, an estimate for \( \theta \) is updated in each data assimilation step. The sequential update needs forecast steps and analysis steps, which transport information of the current time to the next observation time in the forecast step. At the time step \( k \), we denote the forecast by \( \theta_f^k \), the analysis by \( \theta_a^k \), forecast error covariance matrix by \( P_f^k \) and analysis error covariance matrix by \( P_a^k \). Then we have

\[
\begin{align*}
\theta_f^k &= \theta_a^{k-1}, \\
\sigma_k^2 &\sim \text{Inv-gamma}(\alpha + \frac{n_d}{2}, \beta + \frac{\|\mathbf{d}_k - \mathbb{H}(\theta_f^k)\|^2}{2}), \\
P_f^k &= P_a^{k-1}.
\end{align*}
\]

As in (2.8), the posterior is the weighted sum of observations and forecast in the analysis step \( k \), i.e.,

\[
\begin{align*}
\theta_a^k &= \theta_f^k + K_k(\mathbf{d}_k - \mathbb{H}(\theta_f^k)), \\
P_a^k &= (I - K_k\mathbb{H})P_f^k,
\end{align*}
\]

where Kalman gain

\[
K_k = P_f^k\hat{H}^T(HP_f^k\hat{H}^T + R_k)^{-1}, \quad R_k = \sigma_k^2 I.
\]

Here \( \hat{H} \) is the Jacobian matrix of \( \mathbb{H} \) in Extended Kalman Filter (EKF) when \( \mathbb{H} \) is nonlinear.

It is well known that the KF and EKF are numerically scarcely affordable and the storage of a few state vectors is impossible in the high-dimensional systems. To overcome the difficulty, EnKF is desirable for nonlinear data assimilation problems in high-dimensional space. The advantage of EnKF is that we apply a useful approximation to the Kalman filter to avoid propagating the first and second order statistical moments. To this end, Monte Carlo method is used to propagate an ensemble of realizations from the prior distribution. In EnKF, we just update the propagating ensemble and the Kalman gain matrix is approximated by

\[
K_k = \text{Cov}(\Theta_f^k, Z_k)\text{Cov}(Z_k, Z_k)^{-1}, \quad Z_k \sim \mathcal{N}(0, \sigma_k^2 I).
\]

where \( \Theta_f^k \) is the forecast ensemble and \( Z_k \) is the ensemble of simulated observations. Thus, the forecast error covariance matrix and analysis error covariance matrix are not necessary to compute. The true mean and covariance are approximated by ensemble mean and ensemble covariance, respectively. In the paper, we make use of the stochastic analysis ensemble generation method, where the simulated observations are perturbed by simulated observation error \( \varepsilon_f \). The \( \varepsilon_f \) is independent of \( \varepsilon \).
Let \( \{d_1, d_2, \ldots, d_k, \ldots, d_I\} \) be a time series of observations and \( d_k \in \mathbb{R}^{n_d} \) \((k = 1, \ldots, I)\). We assume that the prior distribution of \( \theta \) is \( \mu_0 \) (Gaussian), and \( \varepsilon_k^f \sim N(0, \sigma_k^2 I) \), where \( \sigma_k^2 \) is unknown. We initially pick \( M \) ensemble members for EnKF, through which we obtain the analysis ensemble \( \Theta^a_I \). Furthermore, the mean and covariance of \( \Theta^a_I \) can be used to estimate the unknown parameter. The pseudo-code of EnKF algorithm is presented in Algorithm 1.

**Algorithm 1** Sequential EnKF algorithm with unknown \( \sigma^2 \)

**Input**: number of ensemble members \( M \), initial ensemble members \( \{\theta_1,0, \ldots, \theta_M,0\} \) drawn from prior \( \mu_0 \), the number of data assimilation steps \( I \), observations \( \{d_1, d_2, \ldots, d_k, \ldots, d_I\} \).

**Output**: \( \Theta^a_I \).

1. \( \Theta^a_0 = (\theta_1,0, \ldots, \theta_M,0) \)
2. for \( k = 1 : I \)
   1. **Forecast/predictor**: Generate ensemble of \( z_k \) by
      \[
      \Theta^f_k = \Theta^a_{k-1}
      \]
      for \( j = 1 : M \)
      \[
      \theta^f_{j,k} = \Theta^f_k(:,j), \quad Z(:,j) = H(\theta^f_{j,k}), \quad \sigma_s^2 = \frac{\|d-E(\theta^f_{j,k})\|^2}{n_d-n_p}, \quad \beta = \sigma_s^2 n_s, \quad S(j) \sim \text{Inv-gamma}\left(\alpha + \frac{n_d}{2}, \beta + \frac{\|d_k-E(\theta^f_{j,k})\|^2}{2}\right).
      \]
      end for
      \[
      \sigma_k^2 = \frac{\sum_{j=1}^M S(j)}{M}, \quad E(:,j) = \varepsilon^f_k \sim N(0, \sigma_k^2 I) \quad (j = 1, \ldots, M), \quad Z_k = Z + E.
      \]
   2. **Analysis/corrector**: Update the previous ensemble \( \Theta^a_{k-1} = (\theta^a_{1,k-1}, \ldots, \theta^a_{M,k-1}) \) by
      \[
      \Theta^a_k = \Theta^f_k + K_k(D_k - Z_k),
      \]
      where \( D_k = [d_1, \ldots, d_k] \in \mathbb{R}^{n_d \times M} \) and \( K_k = \text{Cov}(\theta^f_k, Z_k)\text{Cov}(Z_k, Z_k)^{-1} \).
   end for

**Remark 2.1**. We denote \( (d^I_1, d^I_2, \ldots, d^I_k, \ldots, d^I_I) \in \mathbb{R}^{(I-n_d) \times 1} \) by \( d^{1:I} \). We replace \( H \) by \( H^{1:I} \) in (2.2). We can use ensemble smoother (ES) to do a single global update. Then the analysis in ES is

\[
\Theta^a = \Theta^f + K(D - Z),
\]

where \( D = [d^{1:I}, \ldots, d^{1:I}] \in \mathbb{R}^{(I-n_d) \times M} \).

**2.2 EnKF for non-Gaussian model using normal-score transformation**

In general, EnKF is a Gaussian approximation for the estimated parameter because it reproduces the mean and covariance. If the target distribution is Gaussian and unimodal, EnKF inherently gives an accurate estimation. However, if the target distribution is non-Gaussian or multimodal, the approximation may not capture the properties of target distribution. In this situation, we can invoke the normal-score transformation, which maps
non-Gaussian into Gaussian and is invertible \cite{29}. We perform the normal-score transformation after each forecast step. Let $F_k$ be the normal-score operator at the assimilation step $k$ and satisfy

$$q_k = F_k(\theta_k) \sim N(0, I).$$

Because the support of cumulative distribution function (CDF) is $[0, 1]$, the transformation can be fulfilled by CDF. The normal-score transformation renders the Gaussian random variables one by one, and the multivariate properties of parameter vector are also changed but not necessary to be multi-Gaussian \cite{38}.

We want to incorporate the normal-score transformation into EnKF for non-Gaussian cases. Let $q^f_k$ and $q^a_k$ be the forecast and analysis after the normal-score transformation. Then the forecast step of EnKF implies that

$$\begin{align*}
\theta^f_k &= \theta^a_{k-1}, \\
q^a_{k-1} &= F_{k-1}(\theta^a_{k-1}), \\
q^f_k &= q^a_{k-1}.
\end{align*}$$

The analysis step is followed by

$$\begin{align*}
q^a_k &= q^f_{k-1} + K_k(d_k - H(\theta^f_k)), \\
\theta^a_k &= F^{-1}_k(q^a_k),
\end{align*}$$

where the Kalman gain matrix $K_k$ is approximated by

$$K_k = \text{Cov}(\Xi^f_k, Z_k) \text{Cov}(Z_k, Z_k)^{-1}.$$ 

Here $\Xi^f_k$ is the forecast ensemble after the transformation and $Z_k$ is the ensemble of simulated observations. Let $\mu_0$ be a non-Gaussian distribution and $\varepsilon^f_k \sim N(0, \sigma^2 I)$. We describe the normal-score EnKF (NS-EnKF) in Algorithm 2.

### 3 Surrogate model construction using GMsFEM and sparse gPC

For the EnKF methods presented in Algorithm 1 and Algorithm 2 we need to repeatedly compute the forward model for all ensemble members. This computation is very expensive when the forward model is a complex PDE model and the number of ensemble members is large. In order to significantly accelerate the forward model computation, we construct a surrogate model for the forward model using model reduction methods.

The goal is to approximate a large-scale problem in a low dimensional space. To this end, the key idea is to choose a set of appropriate basis functions, which can span a good approximation space for the solution. If equation (2.1) is linear with respect to $u$, we can derive an algebraic system for (2.1) as follows by applying suitable discretization method

$$K(\theta)u = f,$$  (3.9)
Algorithm 2 NS-EnKF algorithm

Input: number of ensemble members $M$, initial ensemble members $\{\theta_{1,0}, \cdots, \theta_{M,0}\}$ drawn from prior $\mu_0$, the number of data assimilation steps $I$, observations $\{d_1, d_2, \cdots, d_k, \cdots, d_I\}$.

Output: $\Theta^a_I$.

1. $\Theta^a_0 = (\theta_{1,0}, \cdots, \theta_{M,0})$, $\Xi^a_0 = \mathbb{F}_0(\Theta^a_0)$

2. for $k = 1 : I$

   (1). Forecast/predictor: Generate ensemble of $z_k$ by
   
   $\Theta^f_k = \Theta^a_{k-1}$, $\Xi^f_k = \mathbb{F}_{k-1}(\Theta^a_{k-1})$, $\Xi^f_k = \Xi^a_{k-1}$.

   for $j = 1 : M$
   
   $\theta_{j,k} = \Theta^f_k(:, j)$, $q_k = \Xi^f_k(:, j)$, $Z(:, j) = \mathbb{H}(\theta_{j,k})$.
   
   end for

   $E(:, j) = \epsilon^f_j \sim N(0, \sigma^2 I)$ (j = 1, $\cdots$, M), $Z_k = Z + E$.

   (2). Analysis/corrector: Update the previous ensemble $\Theta^a_{k-1}$ and $\Xi^a_{k-1}$ by

   $\Xi^a_k = \Xi^f_k + K_k(D_k - Z_k)$, $\Theta^a_k = \mathbb{F}^{-1}_k(\Xi^a_k)$,

   where $D_k = [d_k, \cdots, d_k] \in \mathbb{R}^{n_d \times M}$ and $K_k = \text{Cov}(\Xi^f_k, Z_k)\text{Cov}(Z_k, Z_k)^{-1}$.

end for

where $u \in \mathbb{R}^{N_h}$ is the numerical solution vector and $f \in \mathbb{R}^{N_h}$ the source vector. The $N_h$ is the number of spatial degree of freedoms and is usually very large if we straightforwardly solve the equation in fine grid. We can use a model reduction method and reduce the number of basis functions to improve the efficiency. Then we can get a reduced algebraic system for (2.1),

$$K_r(\theta)u_r = f_r.$$ 

Let $R \in \mathbb{R}^{N_h \times M_v}$ ($M_v \ll N_h$) be the matrix comprised of the $M_v$ reduced basis functions. Then a projection reduce method implies

$$K_r(\theta) = R^T K(\theta) R, \quad f_r = R^T f.$$ 

In order to accelerate evaluations of the posterior density for each updated parameter ensemble, we use stochastic response surface methods to construct surrogate. The solution $u_r(\theta)$ of the reduced model can be expressed by stochastic basis functions such as polynomial chaos [36], radial basis functions [30], and wavelet basis functions [24]. The surrogate model is constructed through the stochastic collocation method by solving a $l_1$ penalized least-square problem. We use the lagged diffusivity fixed point method for the $l_1$ optimization problem and get a sparse representation for $u_r(\theta)$ using fewer samples.

3.1 GMsFEM

In the paper, we consider the following time fractional PDE model

$$^cD^\alpha_t u - \nabla (k(x) \nabla u) = f(x,t), \quad x \in \Omega, \quad t \in (0, T] \quad (3.10)$$
where $\gamma \in (0, 1) \cup (1, 2)$ and is the fractional order of the derivative with respect to time. Here we consider the Caputo fractional derivative defined by
\[
c^D_t u = \frac{1}{\Gamma(m - \gamma)} \int_0^t (t - \tau)^{m-\gamma-1} \frac{\partial^m u(x, \tau)}{\partial \tau^m} d\tau, \quad m - 1 < \gamma < m,
\]
where $\Gamma(\cdot)$ is the Gamma function and $m$ is a positive integer. The equation (3.10) has a close form subject to a suitable boundary condition and initial condition.

In the model equation, $k(x)$ usually refers to a permeability field in porous media applications. The permeability field has heterogeneous and multiscale structure inherently and results in a multiscale model. We will use general multiscale finite element method (GMsFEM) to reduce the model and get a coarse GMsFEM model. This can achieve a good trade-off between efficiency and accuracy for simulating the forward model. We will apply GMsFEM presented in [4] to the time-fractional diffusion-wave equation (3.10). For GMsFEM, we need to pre-compute a set of multiscale basis functions. To this end, the first step is to construct a snapshot space $V_{\omega_{\text{snap}}}^\omega_i$ for multiscale basis by solving local eigenvalue problem on each coarse block $\omega_i$,
\[
\begin{align*}
-\text{div}(k(x, \theta_j)\nabla \varphi_{l,j}) &= \lambda_{l,j} k(x, \theta_j) \varphi_{l,j} \quad \text{in } \omega_i, \\
k(x, \theta_j) \nabla \varphi_{l,j} \cdot \vec{n} &= 0 \quad \text{on } \partial \omega_i,
\end{align*}
\]
where the samplers $\{\theta_j\}_{j}^{N_{\theta}}$ are drawn from the prior distribution of $\theta$. By a finite element method discretization on underlying fine grid, the local eigenvalue problem can be formulated as an algebraic system,
\[
A(\theta_j) \varphi_{l,j} = \lambda_{l,j} S(\theta_j) \varphi_{l,j},
\]
where
\[
[A(\theta_j)]_{mn} = \int_{\omega_i} k(x, \theta_j) \nabla v_n \nabla v_m, \quad [S(\theta_j)]_{mn} = \int_{\omega_i} k(x, \theta_j) v_n v_m,
\]
and $v_n$ are the basis functions in fine grid. We take the first $M_{\text{snap}}^i$ eigenfunctions corresponding to the dominant eigenvalues for each coarse neighborhood $\omega_i$ (see Figure 3.1), $i = 1, 2, \ldots, N_H$, where $N_H$ is the number of coarse nodes. Hence we construct the space of snapshots by
\[
V_{\text{snap}}^\omega_i = \text{span}\{\varphi_{l,j}, 1 \leq j \leq N_{\theta}, 1 \leq l \leq M_{\text{snap}}^i\}.
\]
The snapshot functions can be stacked into a matrix as
\[
R_{\text{snap}} = [\varphi_1, \cdots, \varphi_{M_{\text{snap}}}] ,
\]
where $M_{\text{snap}} = N_{\theta} \times M_{\text{snap}}^i$ denotes the total number of snapshots used in the construction. The second step is to solve the following local problems in the snapshot space
\[
\begin{align*}
-\text{div}(k(x, \bar{\theta})\nabla \psi_k^j) &= \lambda_k k(x, \bar{\theta}) \psi_k^j \quad \text{in } \omega_i, \\
k(x, \bar{\theta}) \nabla \psi_k^j \cdot \vec{n} &= 0 \quad \text{on } \partial \omega_i,
\end{align*}
\]
where \( \bar{\theta} = \frac{1}{N} \sum_{j=1}^{N} \theta_j \). We define

\[
\begin{aligned}
[A]_{mn} &= \int_{\omega_i} k(x, \bar{\theta}) \nabla \varphi_n \nabla \varphi_m = R_{\text{snap}}^T \bar{A} R_{\text{snap}}, \\
[S]_{mn} &= \int_{\omega_i} k(x, \bar{\theta}) \varphi_n \varphi_m = R_{\text{snap}}^T \bar{S} R_{\text{snap}},
\end{aligned}
\]

where \( \bar{A} \) and \( \bar{S} \) denote fine-scale matrices corresponding to the stiffness and mass matrices, respectively, with the permeability \( k(x, \bar{\theta}) \). We choose the smallest \( M_i \) eigenvalues of the equation

\[ A\psi^i_k = \lambda_k S\psi^i_k \]

and take the corresponding eigenvectors in the snapshot space by setting \( \psi^i_k = \sum_j \psi^i_{k,j} \varphi_j \), for \( k = 1, \cdots, M_i \), to form the reduced snapshot space, where \( \psi^i_{k,j} \) are the coordinates of the vector \( \psi^i_k \).

Let \( \{\chi_i\}_{i=1}^{N_H} \) be a set of partition of unity functions associated with the open cover \( \{\omega_i\}_{i=1}^{N_H} \) of \( \Omega \). Then we multiply the partition of unity functions by the eigenfunctions to construct

\[ V_H = \text{span}\{\Psi_{il} : \Psi_{il} = \chi_i \psi^i_l : 1 \leq i \leq N_H \text{ and } 1 \leq l \leq M_i\}. \]

We can use a single index for the multiscale basis function set \( \{\Psi_{il}\} \) and place them in the following matrix

\[ R = [\Psi_1, \Psi_2, \cdots, \Psi_{M_e}], \]

where \( M_e = \sum_{i=1}^{N_H} M_i \) denotes the total number of multiscale basis functions. We note that once the matrix \( R \) is constructed, it can be repeatedly used for simulation.

Next we present the temporal discretization for the equation (3.10). When \( 0 < \gamma < 1 \), the equation (3.10) is the subdiffusion equation. We use the method in [20] to discretize the fractional derivative and have

\[
\begin{aligned}
\int_0^t (t-\tau)^{-\gamma} \frac{\partial u(x, \tau)}{\partial \tau} d\tau &= \sum_{k=0}^{n-1} \frac{u^{k+1} - u^k}{\Delta t} \int_{t_k}^{t_{k+1}} \frac{d\tau}{(t_n - \tau)^\gamma} + O(\Delta t) \\
&= \frac{\Gamma(1-\gamma)}{s} \sum_{k=0}^{n-1} b_k (u^{k+1} - u^k) + O(\Delta t),
\end{aligned}
\]
where $u^k = u(x, t_k)$, $0 = t_0 < t_1 \cdots < t_n = t$, $t_k = k\Delta t$ and
\begin{equation}
\begin{aligned}
&b_k := (n - k)^{1-\gamma} - (n - k - 1)^{1-\gamma}, \quad k = 0, 1, \cdots, n - 1, \\
&s := \Delta t^\gamma \Gamma(2 - \gamma).
\end{aligned}
\end{equation}

When $1 < \gamma < 2$, the (3.10) is the superdiffusion equation. We can use the method in [18] to discretize the fractional derivative and have
\begin{equation}
\begin{aligned}
\int_0^t (t - \tau)^{1-\gamma} \frac{\partial^2 u(x, \tau)}{\partial \tau^2} d\tau &= \sum_{k=0}^{n-1} \frac{u^{k+2} - 2u^{k+1} + u^k}{\Delta t^2} \int_{t_k}^{t_{k+1}} \frac{d\tau}{(t_n - \tau)^{1-\gamma}} + O(\Delta t)^2 \\
&= \frac{\Gamma(2 - \gamma)}{\bar{s}} \sum_{k=0}^{n-1} \bar{b}_k (u^{k+2} - 2u^{k+1} + u^k) + O(\Delta t)^2;
\end{aligned}
\end{equation}
where
\begin{equation}
\begin{aligned}
&\bar{b}_k := (n - k)^{2-\gamma} - (n - k - 1)^{2-\gamma}, \quad k = 0, 1, \cdots, n - 1, \\
&\bar{s} := \Delta t^\gamma \Gamma(3 - \gamma).
\end{aligned}
\end{equation}

Let $U^n$ be the solution at the $n$–th time level. Then we have the weak formulation for the subdiffusion equation $(0 < \gamma < 1)$,
\begin{equation}
\begin{aligned}
\left\{ \begin{array}{l}
\frac{1}{\bar{a}} \left( \sum_{k=0}^{n-1} (U^{k+1} - U^k) b_k, v \right) + a(U^n, v) = (f(t_n), v), \quad \forall v \in V_H \\
(U^0, v) = (u(x, 0), v), \quad \forall v \in V_H,
\end{array} \right.
\end{aligned}
\end{equation}
where $s$ and $b_k$ are defined in (3.14) and
\begin{equation}
a(u, v) = \int k(x, \theta) \nabla u \nabla v dx, \quad \bar{a}(u, v) = \int q(x) uv dx.
\end{equation}
The weak formulation for the superdiffusion equation $(1 < \gamma < 2)$ reads
\begin{equation}
\begin{aligned}
\left\{ \begin{array}{l}
\frac{1}{\bar{a}} \left( \sum_{k=0}^{n-1} (U^{k+2} - 2U^{k+1} + U^k) \bar{b}_k, v \right) + a(U^{n+1}, v) = (f(t_{n+1}), v) \quad \forall v \in V_H, \\
(U^0, v) = (u(x, 0), v) \quad \forall v \in V_H.
\end{array} \right.
\end{aligned}
\end{equation}

For subdiffusion case $(0 < \gamma < 1)$, we define
\begin{equation}
c_k := \begin{cases}
  b_0 & k = 0, \\
  b_k - b_{k-1} & 1 \leq k \leq n - 1.
\end{cases}
\end{equation}

For superdiffusion case $(1 < \gamma < 2)$, we define
\begin{equation}
\bar{c}_k = \begin{cases}
n^{2-\gamma} - (n - 1)^{2-\gamma} & k = 0, \\
-2n^{2-\gamma} + 3(n - 1)^{2-\gamma} - (n - 2)^{2-\gamma} & k = 1, \\
(n + 2 - k)^{2-\gamma} - 3(n + 1 - k)^{2-\gamma} + 3(n - k)^{2-\gamma} - (n - k - 1)^{2-\gamma} & 2 \leq k \leq n - 1, \\
-3 + 2^{2-\gamma} & k = n, \\
1 & k = n + 1.
\end{cases}
\end{equation}
Then the weak formulation of (3.15) can be rewritten as
\[ \tilde{a}(U^n, v) + sa(U^n, v) = \tilde{a}\left(\sum_{k=0}^{n-1} U^k c_k, v\right) + s\left(f(t_n), v\right). \]

We assume that \( U^n \) has the expansion
\[ U^n = \sum_{j=1}^{M_v} c_j \Psi_j(x), \]
where \{\( \Psi_j(x) \)\} denote the GMsFEM basis functions. Let
\[ u^n_H = (u^n_{H1}, u^n_{H2}, \ldots, u^n_{HM_v})^T. \]
Then for \( k = 1, \ldots, M_v, \)
\[
\sum_{j=1}^{M_v} u^n_{Hj} \tilde{a}(\Psi_j, \Psi_k) + s \sum_{j=1}^{M_v} u^n_{Hj} a(\Psi_j, \Psi_k) = \sum_{i=1}^{n} \sum_{j=1}^{M_v} c_i u^n_{Hj} \tilde{a}(\Psi_j, \Psi_k) + s(f^n, \Psi_k).
\]

Let \( B, K \) and \( F \) be the weighted mass, stiffness matrices and load vector using FEM basis functions in fine grid, respectively. Then the equation gives the following algebraic system,
\[ R^T BR u^n_H + sR^T KR u^n_H = \sum_{i=1}^{n} c_i R^T BR u^n_i + sR^T F. \]

If we define
\[ \tilde{B} = R^T BR, \quad \tilde{K} = R^T KR, \]
then \( u^n_H \) can be computed by the iteration
\[ u^n_H = \left(\tilde{B} + s\tilde{K}\right)^{-1}\left(\sum_{i=0}^{n-1} c_i \tilde{B} u^n_i + sR^T F\right). \]

By using the multiscale basis functions, the solution of (3.15) in fine grid can be obtained by downscaling through the transformation \( Ru^n_H. \)

In a similar way to solving equation (3.15), the GMsFEM solution of (3.16) can be computed by the iteration
\[ u^{n+1}_H = \left(\tilde{B} + s\tilde{K}\right)^{-1}\left(\sum_{i=0}^{n} c_i \tilde{B} u^n_i + sR^T F\right). \]

We note that when GMsFEM is not applied, the full order model solution in fine grid is obtained by the iteration
\[ u^n_h = \left(B + sK\right)^{-1}\left(\sum_{i=0}^{n-1} c_i B u^n_i + sF\right) \quad \text{for} \quad 0 < \gamma < 1, \]
and
\[ u_{n+1}^h = \left( B + \tilde{s}K \right)^{-1} \left( \sum_{i=0}^{n} \tilde{c}_i Bu_i^h + \tilde{s}F \right) \text{ for } 1 < \gamma < 2. \]

By comparing the GMsFEM model with the full order model, we see that the size of \( \tilde{K} \) and \( \tilde{B} \) are \( M_v \times M_v \), but the size of \( K \) and \( B \) are \( N_h \times N_h \) (\( M_v \ll N_h \)). Thus a much smaller system is solved in GMsFEM. The matrix \( R \) for multiscale basis functions is computed overhead and it can be repeatedly used for all the time levels. This significantly improves the efficiency for forward model simulations.

**Remark 3.1.** In the sequential data assimilation process, ensemble members \( \{ \theta_j \} \) update and so \( \bar{\theta} \) of (3.13) updates as well. We can update the GMsFEM basis matrix \( R \) to improve the GMsFEM model.

### 3.2 Stochastic collation method using \( l_1 \) regularized least-squares

Stochastic collocation method is an efficient approach to approximate the solution of PDEs with random inputs. In this paper, we use stochastic collocation method (SCM) to obtain an expansion for observation operator \( \mathbb{H}(\theta) \) and efficiently evaluate the simulated observation ensemble.

We use generalized polynomial chaos (gPC) functions to represent \( \mathbb{H}(\theta) \) by \( l_1 \) penalized least-squares method. Let \( i \) be a multi-index with \( |i| = i_1 + \cdots + i_n \) and \( N_0 \) be a nonnegative integer. The \( N_0 \)th-degree gPC expansion of \( \mathbb{H}(\theta) \) is then approximated by a linear combination of gPC basis \( \{ \Phi_i(\theta) \}_{i=1}^P \), i.e.,

\[ \mathbb{H}(\theta) \approx \mathbb{H}^{N_0}(\theta) := \sum_{i} c_i \Phi_i(\theta), \quad P = \frac{(N_0 + n_z)!}{N_0!n_z!}. \]  

(3.17)

The coefficients of expansion are obtained by choosing some collocation points and least-squares method. We first take \( Q \) realizations \( \{ \theta^i \}_{i=1}^Q \) of \( \theta \) in the support of prior distribution \( p(\theta) \). Then for each \( i = 1, \cdots, Q \), we solve a deterministic problem at the node \( \theta^i \) to obtain \( \mathbb{H}(\theta^i) \). After we obtain all pairs \( \{ \theta^i, \mathbb{H}(\theta^i) \} \) \( (i = 1, \cdots, Q) \), we are able to construct a approximation of \( \mathbb{H}(\theta) \) such that \( \mathbb{H}^{N_0}(\theta^i) = \mathbb{H}(\theta^i) \) for all \( i = 1, \cdots, Q \). Thus, (3.17) can produce a system of linear equations

\[ Ac = b, \]  

(3.18)

where \( A \in \mathbb{R}^{Q \times P} \) is the matrix with the entries

\[ A_{ij} = \Phi_j(\theta^i), \quad i = 1, \cdots, Q, \quad j = 1, \cdots, P. \]

and the right term \( b \) satisfying

\[ b_i = \mathbb{H}(\theta^i), \quad i = 1, \cdots, Q. \]

If we solve the system in the ordinary least-squares method, the system (3.18) should be overdetermined, i.e., \( Q \) should be much larger than \( P \). This means that we need solve the
forward model for a large number of samples. To reduce the computation burden, we can take much fewer samples \((Q \leq P)\) and use \(l_1\) regularized least-squares, i.e.,

\[
\min_{\mathbf{c}} \|\mathbf{c}\|_1 \quad \text{s.t.} \quad \mathbf{A}\mathbf{c} = \mathbf{b},
\]

which is equivalent to the optimization problem

\[
\min_{\mathbf{c}} \|\mathbf{A}\mathbf{c} - \mathbf{b}\|_2 + \alpha \|\mathbf{c}\|_1. \quad (3.19)
\]

We use the lagged diffusivity fixed point method [34] for the \(l_1\) penalized least-squares problem. Due to the nondifferentiability of the \(l_1\) norm, we take an approximation to the penalty \(\|\mathbf{x}\|_1\) such as \(\sqrt{\|\mathbf{x}\|^2 + \beta^2}\), where \(\mathbf{x}\) is a scalar and \(\beta\) is a small positive parameter. We denote the approximated penalty by

\[
\mathcal{J}(\mathbf{c}) = \sum_{i=1}^{P} \psi(|c_i|^2),
\]

where

\[
\psi(t) = 2\sqrt{t + \beta^2}.
\]

For any \(\mathbf{v} \in \mathbb{R}^P\),

\[
\frac{d}{d\tau} \mathcal{J}(\mathbf{c} + \tau \mathbf{v}) = \sum_{i=1}^{P} \psi'(|c_i|^2)c_i v_i = \langle \text{diag}(\psi'(\mathbf{c}))\mathbf{c}, \mathbf{v} \rangle,
\]

where \(\text{diag}(\psi'(\mathbf{c}))\) denotes the \(n \times n\) diagonal matrix whose \(i\)th diagonal entry is \(\psi'(|c_i|^2)\), and \(\langle \cdot, \cdot \rangle\) denotes the Euclidean inner product on \(\mathbb{R}^P\). From this we obtain the gradient

\[
\text{grad} \mathcal{J}(\mathbf{c}) = \mathcal{L}(\mathbf{c})\mathbf{c},
\]

where \(\mathcal{L}(\mathbf{c}) = \text{diag}(\psi'(\mathbf{c}))\) and is positive semidefinite. For convenience, we present the lagged diffusivity fixed point method for the \(l_1\) regularized least-squares problem \((3.19)\) in Algorithm 3.

**Algorithm 3** Lagged diffusivity fixed point method for the \(l_1\) regularized least-squares problem \((3.19)\)

**Input:** \(\nu := 0, \mathbf{c}_0 := \text{initial guess}, \alpha\)

**Output:** \(\mathbf{c}\)

begin fixed point iterations

\[
\mathcal{L}_\nu := \mathcal{L}(\mathbf{c}_\nu);
\]

\[
\mathbf{g}_\nu := A^T(\mathbf{A}\mathbf{c}_\nu - \mathbf{d}) + \alpha\mathcal{L}_\nu\mathbf{c}_\nu;
\]

\[
H = A^T A + \alpha \mathcal{L}_\nu;
\]

\[
\mathbf{s}_{\nu+1} := -H^{-1}\mathbf{g}_\nu;
\]

\[
\mathbf{c}_{\nu+1} := \mathbf{c}_\nu + \mathbf{s}_\nu;
\]

\[
\nu := \nu + 1;
\]

end fixed point iterations.

We note that the forward model is solved \(Q\) times to obtain the sampling vector \(\mathbf{b}\). The accuracy of surrogate model can be ensured using much fewer samples (i.e., \(Q < P\)
when the $l_1$ regularized least-squares method is used. As the number of unknown parameters increases, the simulation times for the forward model will increase significantly. To treat the challenge, we use GMsFEM to solve the forward model on a coarse grid for each sample to improve the simulation efficiency.

## 4 Two-stage ensemble Kalman filter using GMsFEM coarse model

In this section, we present a two-stage EnKF using GMsFEM coarse model to accelerate posterior exploration and improve the sequential assimilation performance. The goal of inverse problems is to identify an appropriate solution which can minimize the misfit between the forward model and measurements. Here it is equivalent to solving the minimization problem (2.7). In this paper, we use EnKF methods to solve the minimization problem. EnKF is a sampling method and avoids expensive gradient computation for solving minimization problem.

Although EnKF method can avoid linearization and repeated sampling to explore posterior density, it requires to compute the forward model many times in each forecast step described in algorithm [1]. When the number of unknown parameters $\theta$ is large, we need a large number of ensemble members to estimate $\theta$. This implies that the computation of the forecast is very expensive. To improve the computation efficiency, we construct a surrogate model based on sparse gPC and GMsFEM to approximately represent the full order model. However, EnKF has an inherent constraint assumption for the prior distribution, which must be a Gaussian distribution. As we know, the support of Gaussian distribution is $\mathbb{R}$, but the posterior is often concentrated in a small portion of the entire prior support in many inference problems. Thus, it may be much more efficient to build a surrogate only over the important region of a posterior than the entire prior support. Inspired by the idea, we proposed a two-stage EnKF method. In the first stage, we build a new prior by a very coarse GMsFEM model, where we exclude the unimportant region of the posterior, and the initial ensemble members are drawn from the new prior to enter the EnKF assimilation process. The second stage is the surrogate model based EnKF. We update the surrogate model dynamically when a new analysis is obtained.

The objective of the surrogate model is to construct a representation that quantifies the primary features of the high-fidelity model while providing the computational efficiency required for uncertainty quantification. In stationary Bayesian inference, we may need to build surrogate model only once. However, EnKF method integrates new measurement data in each assimilation step. Thus, the surrogate model need to be updated sequentially. To construct the current surrogate model, we exclude the unimportant region by the previous analysis.

In the first stage, we can apply standard EnKF method based on the very first few levels of measurement data to construct the new prior. In the second stage, we use only a few ensemble members from the previous analysis and $l_1$ regularized least-squares to build the current surrogate model, which allows fast forward model evaluations to generate observation ensemble. The outline of two-stage EnKF is presented in Algorithm [4].
Algorithm 4 Two-stage EnKF with unknown measurement noise algorithm

Input: \( n_1 \) and \( n_2 (n_1 < n_2) \), \( M_1 \) and \( M_2 (M_1 < M_2) \), \( I_1 \) and \( I_2 (I_1 < I_2) \)

Output: final ensemble \( \Theta_{I_2} \)

1. As the GMsFEM described in section 3, we build the multiscale basis functions matrix \( R \).

First stage:

2. Take \( M_1 \) basis functions to construct a very coarse model

3. Set the initial ensemble \( \Theta_0 \) with \( n_1 \) samples

   for \( k = 1, \ldots, I_1 \)
   
   Run algorithm 1 where \( \mathbb{H} \) is the GMsFEM model based on \( M_1 \) basis functions

end for

Second stage:

4. The step 2 and 3 is to obtain \( \Theta_{I_1}^a \) based on the first \( I_1 \) levels of data information. The support of \( \Theta_{I_1}^a \) can be much smaller than the original prior support. We calculate the mean and covariance of \( \Theta_{I_1}^a \) obtained in step 3. Take new initial ensemble \( \Theta_0^{\text{new}} \) with \( n_2 \) samples from the new prior.

5. for \( k = I_1 + 1, \ldots, I_2 \)
   
   (a) Let \( \Theta_{I_1}^a = \Theta_0^{\text{new}} \). Construct the surrogate model using multiscale basis functions based on \( \Theta_{k-1}^a \), where the forward model is solved by GMsFEM with \( M_2 \) basis functions.
   
   (b) Generate the observation ensemble by substituting ensemble members into the surrogate model based on algorithm 3
   
   (c) Run algorithm 1 and obtain the new analysis \( \Theta_k^a \)

end for
The algorithm 4 can render an accurate posterior for Gaussian models. However, for non-Gaussian distributed parameter, we need to construct the surrogate model based on the support of non-Gaussian distribution, while the EnKF only works well for Gaussian prior. For this situation, we use normal-score EnKF to perform non-Gaussian models. We note that the Legendre orthogonal polynomials can be used in surrogate model construction if the support of unknown parameters is bounded.

Algorithm 5 Two-stage EnKF by normal-score transformation

**Input:** \( n_1 \) and \( n_2 (n_1 < n_2) \), \( M_1 \) and \( M_2 (M_1 < M_2) \), \( I_1 \) and \( I_2 (I_1 < I_2) \)

**Output:** final ensemble \( \Theta_{I_2} \)

The step 1 and 2 is the same as 1 and 2 in algorithm 4

**First stage:**

3. Set the initial ensemble \( \Theta_0 \) with \( n_1 \) samples, \( \Xi_0 = F_0(\Theta_0) \)
   
   for \( k = 1, \cdots, I_1 \)
   
   Run algorithm 2 where \( \mathbb{H} \) is the GMsFEM model based on \( M_1 \) basis functions
   
   end for

**Second stage:**

4. The step 2 and 3 is to obtain \( \Theta^a_{I_1} \) and \( \Xi^a_{I_1} \), and we calculate the mean and covariance of \( \Theta^a_{I_1} \) obtained in step 3. Take new initial ensemble \( \Theta^\text{new}_0 \) with \( n_2 \) samples from the new prior.

5. for \( k = I_1 + 1, \cdots, I_2 \)

   The step (a) and (b) are the same as in algorithm 4

   (c) Run algorithm 2 and obtain the new analysis \( \Theta^a_k \) and \( \Xi^a_k = F_k(\Theta^a_k) \)

end for

Remark 4.1. In algorithm 4 and 5, \( I_2 \) is the total number of data assimilation steps. In step 3, we can use ES method to build the new prior when some of the very first measurement data is uninformative.

5 Numerical examples

In this section, we consider the time fractional PDE model (3.10) and estimate the model’s unknown parameters and structures using the proposed two-stage EnKF. A few numerical results will be presented for the estimation for different unknown sources of the dynamic model. In Subsection 5.1, we recover a channel structure in permeability \( k(x) \) when the fractional derivative \( \gamma \) is known. In Subsection 5.2, we estimate the source locations when the diffusion type of equation (3.10) is unknown. In Subsection 5.3, we will recover a permeability pattern when measurement noise is unknown and is treated as a hyperparameter.

For the numerical examples, we consider a dimensionless square domain \( \Omega = [0, 1] \times [0, 1] \) for spatial variable and \( (0, T] \) for time, and we set the initial condition as \( u(x, 0) = 0 \) and in
addition, for the superdiffusion equation, we set
\[ \frac{\partial u(x; t)}{\partial t} \bigg|_{t=0} = 0. \]

Measurement data are generated synthetically by using FEM in a fine grid with time step \( \Delta t = 0.001 \), and the measurement noise is set to be \( \sigma = 0.01 \). For any given realizations of \( \theta \), we solve the time fractional diffusion-wave equation using GMsFEM with time step \( \Delta t = 0.002 \). The regularization parameter \( \alpha \) is set as 0.01 in the lagged diffusivity fixed point algorithm. For all numerical examples, the number of samplers in constructing the gPC surrogate is set to be the number of the gPC basis functions for approximation. We parameterize the Gaussian random fields by Karhunen-Loève expansion (KLE) with a given covariance function \( C(x, y) \) and truncate the KLE to approximately represent the random fields. For the random field \( h(x, \omega) \) by the first \( N \) terms can be represented by
\[
  h(x, \omega) = \mathbb{E}[h(x, \omega)] + \sum_{i=1}^{N} \sqrt{\lambda_i} \theta_i(\omega) \varphi_i(x),
\]
where \( \mathbb{E}[\cdot] \) is the expectation operator, \( \theta_i(\omega) \sim \mathcal{N}(0, 1) \) and \((\lambda_i, \varphi_i)\) are the eigenpairs of the eigenvalue problem
\[
  \int_{\Omega} C(x, y) \varphi_i(y) dy = \lambda_i \varphi_i(x). \tag{5.20}
\]

We sort the eigenvalues in ascending order, i.e., \( \lambda_1 \geq \lambda_2 \cdots \), and their corresponding eigenfunctions are also sorted accordingly and \( \{\theta_i\}_{i=1}^{N} \) are uncorrelated random variables. We will compare the estimation results obtained by using standard EnKF method with the proposed two-stage EnKF.

### 5.1 Recover a channel structure in permeability field

In this subsection, we consider the subdiffusion model (3.10) with mixed boundary condition, where Dirichlet boundary conditions is
\[
  u(0, y; t) = 1, \quad u(1, y; t) = 0, \tag{5.21}
\]
and there is no flow on the other boundaries. The source term is set as \( f = 10 \), the end time is \( T = 0.11 \), and the fractional derivative is given by \( \gamma = 0.5 \). The permeability field is unknown here, and we only have the prior information of the permeability field, which is structured with a channel that lies between \( y = 0 \) and \( y = 1 \). The spatial domain is divided into 3 parts by this channel and the permeability is a constant at each subregion. Thus, we can describe the boundaries of the channel by two curves \( \Gamma_1 \) and \( \Gamma_2 \), which can be expressed as
\[
  \Gamma_1(x) = \sum_{i=1}^{m_1} w_i^1 \phi_i(x), \quad \Gamma_2(x) = \sum_{i=1}^{m_2} w_i^2 \phi_i(x),
\]
where $\phi_i(x)$ are interpolation basis functions. In order to reduce dimension of unknown parameters, $\phi_i(x)$ are set to be Karhunen-Loève expansion (KLE) basis functions, i.e. $\phi_i(x) = \sqrt{\lambda_i} \varphi_i(x)$, and $E[\Gamma_i(x)] = 0$ $(i = 1, 2)$. The covariance function is set as

$$C(x_1, x_2) = \xi^2 \exp(-\frac{||x_1 - x_2||^2}{2l^2})$$

to solve the eigenvalue problem (5.20), where $\xi = 1$ and $l^2 = 0.1$. A criterion which is adopted for the choice of truncated term $N$ is

$$\frac{\sum_{j=1}^{N} \lambda_j}{\sum_{j=1}^{\infty} \lambda_j} > 99.99\%.$$ 

We truncate the KLE expansion for functions $\Gamma_i(x)$, $(i = 1, 2)$ by this criterion, and then we get $m_1 = 5$ and $m_2 = 5$. In order to constrain the curves in the unit square domain, we make a bijective transformation

$$\tilde{\Gamma}_i(x) = \frac{1}{2} + \frac{1}{\pi} \arctan(\Gamma_i(x)), \quad i = 1, 2.$$ 

Thus, $\tilde{\Gamma}_1(x) \in [0, 1]$ and $\tilde{\Gamma}_2(x) \in [0, 1]$ can be guaranteed, i.e., the curves lie in the physical domain. Then we can construct the level set functions corresponding to the two curves as

$$L_i(x, y) = \text{Heaviside}(y > \tilde{\Gamma}_i(x)), \quad i = 1, 2.$$ 

Hence, the random field can be parameterized as

$$\log k(x, y) = c_1 L_1 L_2 + c_2 (1 - L_1)L_2 + c_3 (1 - L_1)(1 - L_2),$$

where $\{c_i\}_{i=1}^{3}$ are unknowns. Thus, we have the unknown parameter vector

$$\theta = (c_1, c_2, c_3, w_1^1, \cdots, w_{m_1}^1, w_1^2, \cdots, w_{m_2}^2).$$

In this example, the reference permeability is generated by setting

$$\Gamma_1(x) = 0.7 + 0.1 \sin(3\pi x),$$

$$\Gamma_2(x) = 0.4 + 0.2 \sin(2\pi x + 0.1),$$

and the value of each subregion is set as $(c_1, c_2, c_3) = (0, 4, 1)$, which is shown in Figure 5.2 (left). In the example, the number of artificial time steps for data assimilation is $I_2 = 9$. Measurements are taken at time instances $0.012 + 0.01I : 0.002 : 0.018 + 0.01I$ in each data assimilation step, where $I \in \{1, 2, \cdots, I_2\}$ and the measurement locations are distributed on the uniform $5 \times 5$ grid of the domain $[0.1, 0.9] \times [0.1, 0.9]$ as shown in Figure 5.2 (right).

The forward model is defined on a uniform $80 \times 80$ fine grid, and we set the coarse grid $5 \times 5$ for GMsFEM simulation. We construct the local snapshot space with dimension $M_{\text{snap}}^i = 20$ and select $M_c = 10$ multiscale basis functions at each block to construct the coarsen reduced order model and solve the optimization problem (2.7).
To construct a new prior, we use $5 \times 10^3$ ensemble members by the standard EnKF method in the first stage. Then the new prior is constructed by incorporating data information from the first three data assimilation steps. Seven local multiscale basis functions ($M_i = 7$) are selected in constructing the basis matrix $R$ to construct coarse model. Then, we construct the surrogate model after obtaining the new prior ensemble in the second stage. Eight local multiscale basis functions ($M_i = 8$) are selected in constructing the matrix $R$ to construct the gPC surrogate model in the second stage. When the order of gPC is set as $N_0 = 3$, the number of random samplers is 560 for computing vectors $b$ and $A$ in Section 3.2. In this stage, the number of ensemble members is set as $10^4$. For a comparison, we also implement the standard EnKF, which directly uses the previous analysis as a prior to the current moment in data assimilation process. In the standard EnKF, we also use GMsFEM with 8 local GMsFE basis functions to solve forward model for $10^4$ ensemble members. But no new prior and no gPC surrogate model are build in the standard EnKF method. To complete the assimilation process, the standard EnKF method takes about 4.7 hours for CPU time, while the two-stage EnKF just take about 1 hour CPU time. Thus, the two-stage EnKF is much more efficient than the standard EnKF.

Figure 5.2: True profile of $\log k(x)$ (left) and observation locations (right).

Figure 5.3: $\log k(x, \omega)$ of initial prior for two different EnKF methods (left), $\log k(x, \omega)$ of new initial prior for the two-stage EnKF (right).

Figure 5.3 shows the initial prior ensemble (mean) and new prior ensemble (mean) for the two-stage EnKF method in logarithmic scale. We can see that the initial prior gives a very rough structure for the channel, and the new prior improves the channel pattern.
we compare the mean estimate by standard EnKF with the one by two-stage EnKF. The posterior mean and variance for the two methods are presented in Figure 5.4, from which we find that the two-stage EnKF offers more accurate estimation than standard EnKF. The large variance is concentrated around the two boundaries of the channel because two boundaries are unknown.

In order to measure the estimate accuracy in the update process, we define the relative errors $\varepsilon_{\Gamma_1}$ and $\varepsilon_{\Gamma_2}$ corresponding to the two curves after each update by

$$
\varepsilon_{\Gamma_1} := \frac{\| \tilde{\Gamma}_1(x) - \Gamma_1(x) \|}{\| \Gamma_1(x) \|}, \quad \varepsilon_{\Gamma_2} := \frac{\| \tilde{\Gamma}_2(x) - \Gamma_2(x) \|}{\| \Gamma_2(x) \|},
$$

where $\Gamma_1(x)$ and $\Gamma_2(x)$ are the reference boundary functions. We note that the first three data assimilation steps are used to produce the new prior for the two-stage EnKF. We list the relative errors in Table 1. As expected, the relative error gradually decreases when more measurement data is used in the inference. The table shows that the two-stage EnKF gives more accurate estimates than standard EnKF.

To access the prediction using the posterior model, we compute the 95% credible and predictive intervals for model response at $u((x, 0.5); t)$ and $u((0.5, y); t)$ for the two EnKF methods. We note that the realizations of model response of initial prior are constructed by GMsFE model. This coarsen model error is added to the estimated error variance to construct prediction intervals. As illustrated in Figure 5.5 and 5.6, both the credible interval and predictive interval become narrower as assimilation moves on. This means that as the uncertainty from input $\theta$ decreases with respect to assimilation step and the uncertainty associated with the model fit and predictions decreases. The observation/measurment data are almost contained in the predictive intervals when sufficient data is used in the posterior model.

---

Table 1: The relative error of $\tilde{\Gamma}_1$ and $\tilde{\Gamma}_2$

| Update     | Two-stage EnKF $\varepsilon_{\tilde{\Gamma}_1}$ | Two-stage EnKF $\varepsilon_{\tilde{\Gamma}_2}$ | Standard EnKF $\varepsilon_{\tilde{\Gamma}_1}$ | Standard EnKF $\varepsilon_{\tilde{\Gamma}_2}$ |
|------------|------------------|------------------|------------------|------------------|
| initial prior | 0.318            | 0.407            | 0.318            | 0.407            |
| I=4        | 0.120            | 0.129            | 0.183            | 0.181            |
| I=5        | 0.111            | 0.102            | 0.144            | 0.180            |
| I=6        | 0.109            | 0.095            | 0.118            | 0.176            |
| I=7        | 0.098            | 0.086            | 0.120            | 0.179            |
| I=8        | 0.099            | 0.088            | 0.120            | 0.179            |
| I=9        | 0.096            | 0.078            | 0.120            | 0.178            |
Figure 5.4: Mean and variance of log $k(x, \omega)$ by standard EnKF (the first row and the third row) and two-stage EnKF (the second row and the fourth row) at the assimilation step 4 and 9.
For this example, the measurement noise is the additional Gaussian type. Let $\theta^*$ denote the truth parameter, the discrepancy between the data and forward model is defined by

$$[d - \mathbb{H}(\theta^*)]_i \sim N(0, \sigma^2), \quad i = 1, \ldots, n_d,$$

where $n_d$ is the number of measurements and its expectation is given by

$$\mathbb{E}||d - \mathbb{H}(\theta^*)||^2 = n_d\sigma^2.$$

We plot $\mathbb{E}||d - \mathbb{H}(\theta^*)||^2$ against data assimilation step in Figure 5.7. Then we can see the expectation of the discrepancy tends to $n_d\sigma^2$ as assimilation time moves on. This implies that the final ensemble mean is an accurate estimate.

Figure 5.8 shows the marginal posterior density against the data assimilation step in the two-stage EnKF. Although there are no truth for these parameters $\{\omega^1\}_{i=1}^5$ and $\{\omega^2\}_{i=1}^5$, the data is sufficiently informative to identify a small range of values for the unknown parameters. The important region of marginal densities becomes narrower as data information gains.

### 5.2 Estimate the source locations and fractional derivative

In this section, we consider a diffusion-wave problem, i.e., the diffusion type is unknown, where the boundary conditions are the same as in Subsection 5.1. The end time is set as $T = 0.1$, and the source term is given by

$$f(x, t) = \frac{s_1}{2\pi t_1} \exp\{-\|x_1-x\|^2\} \left[1 - H(t - T_m)\right] + \sum_{i=2}^{m} \frac{s_i}{2\pi t_i} \exp\{-\|x_1-x\|^2\} H(t - T_{i-1}),$$

where $s_1$ and $s_i$ are the source terms at $x_1 = 0.5$ and $x_i$ for $i = 2, \ldots, m$. The weights $w_i$ are determined by the number of sources and the spatial distribution of the sources.
Discrepancy principle

Figure 5.7: \( \mathbb{E} [\| \mathbf{d} - \mathbb{H}(\boldsymbol{\theta}) \|^2] \) and \( n_d \sigma^2 \) via the assimilation time. We note \( I = 1 \) denotes the initial prior time and \( I = 3 \) the new prior time.

Figure 5.8: Marginal posterior density estimation for \( \omega_i^j \) \((i = 1, 2; \ j = 1, 3, 5)\) in different data assimilation steps.
where $H(t)$ is the heaviside function, $s_i$ is the strength, $\tau_i$ is the width and $\chi_i$ is the source location. In this example, both $\chi_i$ and $\gamma$ are unknown. We just have the prior $\chi_i = (\chi_i^1, \chi_i^2) \in [0, 1] \times [0, 1]$, and $\gamma \in (0, 2)$ and $\gamma \neq 1$. The prior is bounded, which can be seen as uniform distribution and $\gamma$ lies in the interval $(0, 2)$ for diffusion-wave equation. But its values are uncontrollable during the data assimilation steps, the samples may run out of the interval. Thus, the two-stage EnKF by normal-score method can be used to avoid the issue. To this end, we use a bijective map $F: E \rightarrow \mathbb{R}^5$, where 

$$E = (0, 2) \times (0, 1)^4.$$ 

Let $\theta = (\gamma, \chi_1, \chi_2)^T$ and $F(\theta) = q$, the map $F$ is set as

$$q_i = F_1(\theta_i), \quad i = 1, \ldots, 5.$$ 

The dynamic system correspondingly becomes

$$\begin{align*}
q_k &= \mathbb{H}(\theta_{k-1}) + \varepsilon_k, \\
\theta_k &= \theta_{k-1}, \\
q_k &= \mathbb{F}_k(\theta_{k-1}), \\
\theta_{k-1} &= \theta_{k-1}, \\
q_{k-1} &= F_{k-1}(\theta_{k-1}), \\
\end{align*}$$

where $q_k = (q_1^k, \ldots, q_5^k)^T$ and $k$ is the artificial time for data assimilation, and the Kalman gain is approximated by $K_k = \text{Cov}(q_k, y_k)\text{Cov}(y_k, y_k)^{-1}$.

For the source term, we take $m = 2$, $T_1 = 0.05$, $T_2 = T$, and $\tau_i = 0.1$ ($i = 1, 2$), $s_1 = 3$, $s_2 = 1$, and the truth source locations are set as $\chi_{1T}^{tr} = (0.2, 0.6)$, $\chi_{2T}^{tr} = (0.5, 0.3)$ and $\gamma^{tr} = 0.5$. Here, the total artificial time steps for data assimilation is set as $I_2 = 8$. Measurements are taken from the subdiffusion equation at time instances $0.012 + 0.01I : 0.006 : 0.018 + 0.01I$ in each data assimilation step, where $I \in \{1, 2, \ldots, 8\}$ and the locations are distributed on the uniform $5 \times 4$ grid of the domain $[0.1, 0.9] \times [0, 1]$ as shown in Figure 5.9 (right). The high-contrast permeability is known and has the spatial distribution shown in Figure 5.9 (left).

Figure 5.9: Spatial distribution of the permeability $k(x)$ (left) and observation locations (right).

The forward model is defined on $100 \times 100$ uniform fine grid, and GMsFEM is implemented on $5 \times 5$ coarse grid. We choose $M_{\text{snap}} = 10$ eigenfunctions for the local snapshot.
space and use GMsFEM with 9 local multiscale basis functions to approximate the forward model in solving the optimization problem (2.7).

In the subsection, we focus on the two-stage EnKF for the inverse problem. To this end, we use $3 \times 10^3$ ensemble members to construct a new prior by standard EnKF in the first stage. The new prior is constructed by incorporating data information from the first data assimilation step. 9 local multiscale basis functions ($M_i = 9$) are chosen in constructing the matrix $R$ to construct coarse model. Then, we construct the surrogate model after obtaining the new prior ensemble in the second stage. The data assimilation process begins from $I = 1$ in second stage. The same local multiscale basis functions are used in constructing the matrix $R$ to construct the gPC surrogate model. When the order of gPC is set as $N_0 = 7$, the number of random samplers is 792 when computing vectors $b$ and $A$ in Section 3.2. In the second stage of the proposed EnKF, the number of ensemble members is also set as $3 \times 10^3$.

![Number of ensemble members applied to different forward models against data assimilation steps](image)

Figure 5.10: Number of ensemble members applied to different forward models against data assimilation steps, where $I = 0$ is the process of generating a new prior.

We only know the problem is a diffusion-wave equation, which may be the subdiffusion or superdiffusion. Thus, it is important to identify the forward model before estimating the source locations. The result of identifying model is shown in Figure 5.10, where we count the number of ensemble members applied to the two diffusion models. By the figure, we can find that the new prior has actually identified the forward model, where $I = 0$ denotes the process of generating a new prior. In the second stage of two-stage EnKF, the superdiffusion model is almost not used. We define the relative error $\varepsilon$ corresponding to the posterior distribution by

$$
\varepsilon := \frac{\|E(\theta) - \theta^{tr}\|}{\|\theta^{tr}\|},
$$

where $\theta^{tr} = (\gamma^{tr}, \chi_1^{tr}, \chi_2^{tr})$. Then we compute the relative error of the final ensemble mean and get $\varepsilon = 0.08$. Figure 5.11 shows all of the one and two-dimensional posterior marginal of $\theta$, where there exist some correlation between the source locations and $\gamma$, such as $\chi_1^{2}$ and $\chi_2^{2}$, but the others appear uncorrelated and mutually independent based on the shape of their 2-D marginal.
To make prediction, we plot the 95% credible interval and predictive for the model response with the ensemble members and realizations by the surrogate model in Figure 5.12 and Figure 5.13. We note the realizations of model response for initial prior is constructed by GMsFE model. The prediction intervals are constructed by taking account of measurement errors. The credible interval and prediction interval, along with the truth, and observation data, are illustrated for $u((x,0);t)$ and $u((0.9,y);t)$ in Figure 5.12 and Figure 5.13 respectively. We note that the credible intervals are tight in the final data assimilation step. Then we can observe from these figures that the uncertainty associated with both the model fit and predictions decreases with respect to assimilation step. We also plot the marginal density for the initial prior, new prior and posteriors together in Figure 5.14 which clearly shows that the posterior is dynamically updated. From this figure, we note that the new prior identifies the model belonging to subdiffusion.

![Figure 5.11: 1-D and 2-D posterior marginals of $\theta$](image)

![Figure 5.12: 95% predictive interval, 95% credible interval, observation and true value for $u((x,0);t)$](image)

29
Figure 5.13: 95% predictive interval, 95% credible interval, observation and true value for $u((0.9, y); t)$.

Figure 5.14: Marginal posterior density estimation of $\theta$ at different data assimilation steps.
5.3 Estimate a permeability field in a hierarchical model

In this subsection, we consider the fractional superdiffusion equation with homogeneous Dirichlet boundary condition. The source term is set as $f = 20$, and the end time is set as $T = 0.11$. In the inversion model, both the permeability field and observation noise are unknown and we want to estimate them by the two-stage EnKF. To parameterize the permeability field, we set $l_x = 0.2$, $l_y = 0.3$, $\sigma^2 = 1$, and the covariance function is assumed to be

$$C(x, y) = \sigma^2 \exp \left( - \frac{\|x_1 - x_2\|^2}{2l_x^2} - \frac{\|y_1 - y_2\|^2}{2l_y^2} \right).$$

We assume that $\log k(x, \omega)$ can be represented by the following truncate the KLE

$$h(x, \omega) := \log k(x, \omega) = \mathbb{E}[h(x, \omega)] + \sum_{i=1}^{N} \sqrt{\lambda_i} \theta_i(\omega) \phi_i(x), \quad N = 20,$$

where the $\mathbb{E}[h(x, \omega)]$ is plotted in Figure 5.15.

The ground true parameter $\theta^*$ are randomly drawn from the standard multivariate normal distribution and the truth permeability map is depicted in Figure 5.16 (left). The total number of data assimilation steps is set as $I_2 = 9$. Measurements are taken at time instances $0.012 + 0.01i : 0.002 : 0.018 + 0.01i$ in each data assimilation step, where $I \in \{1, 2, \ldots, 9\}$ and the locations are distributed on the uniform $5 \times 5$ grid of the domain $[0.1, 0.9] \times [0.1, 0.9]$ as shown in Figure 5.16 (right).

The full order forward model is defined on $100 \times 100$ uniform fine grid, and GMsFEM is implemented on $5 \times 5$ coarse grid. We choose the number of eigenfunctions selected in calculating the snapshot space as $M_{\text{snap}} = 20$. GMsFEM with 10 offline local multiscale basis functions (i.e., $M_c = 10$) is used to solve optimization problem (2.7).

In the numerical simulation, we use $3 \times 10^3$ ensemble members to construct a new prior by standard EnKF in the first stage, where the first two levels of data information is used. Three online local multiscale basis functions ($M_i = 3$) are selected in constructing the matrix $R$ to construct coarse model to obtain the new prior. Then, we construct the surrogate model after obtaining the new prior ensemble in the second stage. Five online local multiscale basis functions ($M_i = 5$) are selected in constructing the multiscale basis matrix to construct the gPC surrogate model. When the order of gPC is set as $N_0 = 3$, we use 1771 samplers when
computing vectors $b$ and $A$ in Section 3.2. In this stage, the number of ensemble members is set as $10^4$. By Algorithm 4, we need to simultaneously update the hyperparameter $\sigma^2$. We take $n_s = 0.05$ in Algorithm 1. The $\sigma^2$ for true measurement noise is set as $10^{-4}$. The posterior mean and posterior variance via the data assimilation step are illustrated in Figure 5.17, which shows that the uncertainty mainly lies on high-contrast part and boundary.

Figure 5.18 depicts the discrepancy principle when $\sigma^2$ is unknown in the forecast step. As we have mentioned before in Subsection 5.1, the measurement noise is the additional Gaussian type. Then we can see that the expectation of the discrepancy tends to $n_d\sigma^2$ as assimilation time moves on. To measure the estimate accuracy, we define the relative errors corresponding to the posterior distribution by

$$
\varepsilon_k := \frac{\|k(x, \bar{\theta}) - k(x, \theta^*)\|}{\|k(x, \theta^*)\|}, \quad \varepsilon_{\sigma^2} := \frac{\|\sigma^2 - \sigma^2_t\|}{\|\sigma^2_t\|}, \quad \varepsilon_{\theta} := \frac{\|\bar{\theta} - \theta\|}{\|\theta\|},
$$

where $\bar{\theta}$ and $\sigma^2$ are the mean of final assimilation step by the two-stage EnKF and $\sigma^2_t$ is the truth observation noise. We just consider the error of the second stage in the two-stage
EnKF method. The relative error is plotted in the Figure 5.19. As expected, the relative error gradually decreases with the assimilation steps.

The credible interval and prediction interval, along with the true, and observation data, are illustrated for \( u((x, 0.5); t) \) and \( u((0, 5, y); t) \) in Figure 5.20 and Figure 5.21, respectively. We find that the final credible intervals are tight and that the uncertainty associated with both the model fit and predictions decreases with respect to data assimilation step. The corresponding uncertainty decreases as \( x \) and \( y \) get closer to boundary, which is due to the deterministic Dirichlet boundary condition. The marginal densities for the unknown parameters in different assimilation steps are plotted in Figure 5.22, which shows the the support of density distribution becomes narrower as more data information is used in the posterior exploration.
Figure 5.20: 95% predictive interval, 95% credible interval, observation and true value of $u((x, 0.5); t)$.

Figure 5.21: 95% predictive interval, 95% credible interval, observation and true value of $u((0.5, y); t)$.

Figure 5.22: Marginal posterior density estimation of $\theta_1$, $\theta_5$, $\theta_9$, $\theta_{13}$, $\theta_{16}$ and $\theta_{18}$ in different data assimilation steps.
6 Conclusion

We presented a two-stage EnKF using coarse GMsFEM models in the paper. In the first stage, we used a very coarse GMsFEM model to approximate the forward model, and the corresponding misfit-to-observed problem is solved by using GMsFEM. We then constructed the new prior based on the ensembles by traditional EnKF, and then used GMsFEM and gPC to obtain a compact representation for the model response based on the new prior. The two-stage EnKF was employed to explore the surrogate posterior density, which was incorporated by the surrogate likelihood and the updated prior. It showed that the proposed method leads to the approximate posterior with better efficiency and accuracy than traditional EnKF method.

The deterministic and statistical methods were combined together to solve inverse problems. We obtained not only the point estimate and confidence interval but also the statistical properties of the unknowns. A new prior was constructed for Bayesian inference using coarse GMsFEM models. The new prior contains the significant region or support of the posterior and is incorporated with the likelihood to be explored. For non-Gaussian models, we presented a two-stage EnKF using normal score transformation.

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