# Bayesian High-Rank Hankel Matrix Completion for Nonlinear Synchrophasor Data Recovery 

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#### Abstract

Phasor measurement units (PMUs) provide high temporal-resolution synchrophasor measurements for power system monitoring and control. The frequent data quality issues, such as missing and bad data, prevent the incorporation of synchrophasor data in real-time operations. Most existing data-driven data recovery methods assume the power system dynamics can be approximated by a linear dynamical system, and the recovery performance degrades significantly when the power system is experiencing nonlinear dynamics during significant events. This paper proposes a data-driven Bayesian nonlinear synchrophasor data recovery method (Ba-NSDR) that can recover a consecutive time period of simultaneous data losses or errors across all channels, even when the underlying system is highly nonlinear. The idea is to lift the Hankel matrix of the spatial-temporal synchrophasor data to a higher dimension such that the lifted Hankel matrix is low-rank in that space and can be processed with the kernel trick. Our proposed Bayesian method then infers the probabilistic distributions of synchrophasor from the partial observations. Some distinctive features of Ba-NSDR include an uncertainty index to measure the accuracy of the recovery result and the robustness to parameter selections. Our method is verified on both synthetic and recorded event datasets.


Index Terms-PMU data recovery, high-rank matrix completion, Bayesian robust matrix completion, kernel method, uncertainty modeling.

## I. Introduction

PHASOR Measurement Units (PMUs) provide synchronized voltage and current phasor measurements across different locations in the electric power system. With a high sampling rate of thirty or sixty samples per second per channel, synchrophasor data provide great visibility of power system dynamics, which is typically difficult to observe in the supervisory

[^0]control and data acquisition (SCADA) system. Synchrophasor data have been employed for event classification [1], [2], state estimation [3], [4], [5] and system identification [6], [7]. Synchrophasor data, however, suffer from quality issues such as missing and bad data, because of various reasons like PMU malfunctions, communication failure, and false data injections. Synchrophasor data usually have missing and bad data issues. The quality issues prevent synchrophasor data from being employed in real-time control operations.

Various approaches have been developed to handle missing and bad data. The model-based methods utilize a dynamic model [8] to fill the missing data or estimate the dynamic states based on the Kalman filter [9], [10]. The performance critically depends on accurate model estimation. Refs. [11], [12], [13] train deep neural networks to recover missing data. Refs. [14], [15], [16], [17] formulate the error correction as a hypothesis testing problem. Ref. [18] exploits spatial-temporal similarities in the synchrophasor measurements to correct bad data. Ref. [19] employs the independent component analysis to obtain the measurement structure and remove the errors. Refs. [20], [21], [22], [23] exploit the low-rank property of the spatial-temporal PMU data matrix to correct missing and bad data. These data-driven methods, however, cannot handle simultaneous and consecutive data issues across all channels.

When the power system dynamics can be approximated by a linear dynamical system, [24], [25], [26], [27] exploit the resulting low-rank property of the Hankel matrix of PMU data to recover simultaneous and consecutive data issues. The linear dynamical model, however, becomes inaccurate when the power system is experiencing nonlinear dynamics. To the best of our knowledge, only Ref. [28] considers missing data recovery in nonlinear dynamical systems and proposes a lifted low-rank Hankel property to characterize the data dynamics without explicitly modeling the dynamical system. This approach cannot handle bad data, and its performance is very sensitive to parameter selection. Moreover, the recovery performance drops significantly for long consecutive data loss. One major limitation of most methods mentioned above is that they only provide an estimation of the actual data without any evaluation of the accuracy of the estimation. Only Ref. [27] provides an uncertainty evaluation of the recovered data.

This paper proposes a Bayesian high-rank Hankel matrix recovery method (Ba-NSDR) to recover missing data and correct bad data when the power system exhibits significant nonlinear dynamics. The main idea is to lift the original high-rank

Hankel matrix into a higher-dimensional space so that the lifted matrix becomes low-rank. The nonlinear lifting function can be characterized implicitly by the kernel function [29], which has been exploited in high-rank matrix completion [30], [31], [32]. [33] employs the kernel function to incorporate the prior knowledge into the matrix completion, but it does not consider the nonlinear dynamics. [34] uses the Gaussian process with kernel functions to model the linear time-invariant (LTI) systems and approximate the nonlinear dynamical systems by LTI systems. [34] is not modeless and requires detailed system information. A prior probabilistic distribution is imposed over the Hankel matrix, and Ba-NSDR computes the approximate posterior distributions using variational inference based on the observed data. Ba-NSDR has multiple distinctive features. First, it can handle simultaneous and consecutive missing/bad data across all PMU channels. When the system is experiencing nonlinear dynamics, the recovery accuracy by Ba-NSDR is much higher than the existing methods. Second, Ba-NSDR returns an uncertainty index that reflects the accuracy of recovered data, while the recovery accuracy of most existing methods cannot be measured without the ground-truth value. Third, Ba-NSDR does not require any prior knowledge of the unknown ground-truth matrix rank and is robust to the initial rank selection. It can effectively estimate the rank from the observed data through pruning from a large rank.

The rest of the paper is organized as follows. The problem formulation, low-rank Hankel property, and low-rank lifted Hankel property of synchrophasor data are described in Section II. The methodology is introduced in Section III. Section IV reports the numerical results. Section V concludes the paper. The derivation details of our method are shown in the supplementary materials.

## II. Problem Formulation

Let a matrix $Y$ denote the ground truth of PMU measurements of $m$ channels at different locations during $n$ time instants,

$$
\begin{equation*}
\boldsymbol{Y}=\left[\boldsymbol{y}_{1}, \boldsymbol{y}_{2}, \ldots, \boldsymbol{y}_{n}\right] \in \mathbb{R}^{m \times n} \tag{1}
\end{equation*}
$$

where $\boldsymbol{y}_{i} \in \mathbb{R}^{m}$ denotes the measurement of $m$ channels at time instant $i$. Let $N \in \mathbb{R}^{m \times n}$ denote the measurement noise. Let $\boldsymbol{E} \in \mathbb{R}^{m \times n}$ denote the additive bad data. The entries in $\boldsymbol{E}$ can be arbitrarily large, modeling significant bad data. We assume such bad data only happen at a small fraction of measurements, i.e., $\boldsymbol{E}$ is sparse.

Let a matrix $\boldsymbol{Y}^{o} \in \mathbb{R}^{m \times n}$ denote the observed measurements. Each entry $Y_{i, j}^{o}$ in the set $\boldsymbol{\Omega}$ of observed entries is given by

$$
\begin{equation*}
Y_{i, j}^{o}=Y_{i, j}+E_{i, j}+N_{i, j} \quad(i, j) \in \Omega \tag{2}
\end{equation*}
$$

where $\Omega$ denotes the set of observed entries. The unobserved entries in $\boldsymbol{Y}^{o}$ are irrelevant and set as zeroes for the completeness of the definition.

The objective of this paper is to recover data $\boldsymbol{Y}$ with measurable accuracy from measurements $\boldsymbol{Y}^{o}$ that are corrupted by missing data, bad data, and noise. This is particularly challenging when the power system is under nonlinear dynamics.

Our proposed Ba-NSDR method exploits the low-rank property of the lifted Hankel matrix of the PMU data in nonlinear
dynamical systems. We first introduce the low-rank Hankel property for linear dynamical systems in Section II-A and then generalize to the lifted Hankel matrix for nonlinear dynamical systems in Section II-B. Detailed analyses of low-rank Hankel property can be found in Refs. [24] and [28], respectively.

## A. Low-Rank Hankel Property of PMU Data

Let $\mathcal{H}_{n_{2}}(\boldsymbol{Y}) \in \mathbb{R}^{m n_{2} \times n_{1}}\left(n_{1}+n_{2}=n+1\right)$ denote the Hankel matrix of $\boldsymbol{Y}$, where the $j$ th column of $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ includes all the measurements in $m$ channels from time $j$ to $j+n_{2}-1$, i.e.,

$$
\mathcal{H}_{n_{2}}(\boldsymbol{Y})=\left[\begin{array}{cccc}
\boldsymbol{y}_{1} & \boldsymbol{y}_{2} & \ldots & \boldsymbol{y}_{n_{1}}  \tag{3}\\
\boldsymbol{y}_{2} & \boldsymbol{y}_{3} & \ldots & \boldsymbol{y}_{n_{1}+1} \\
\vdots & \vdots & \ldots & \vdots \\
\boldsymbol{y}_{n_{2}} & \boldsymbol{y}_{n_{2}+1} & \ldots & \boldsymbol{y}_{n}
\end{array}\right] \in \mathbb{R}^{m n_{2} \times n_{1}}
$$

As shown in [24], if the underlying system that produces output $\boldsymbol{y}_{1}$ to $\boldsymbol{y}_{n}$ can be approximated by an order- $r$ (integer $r \geq 1$ ) linear dynamical system, then $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ can be approximated by a rank- $r$ matrix. $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ is low-rank because $r$ can be much smaller than $m$ and $n_{1}$. The rank- $r$ approximation $\mathcal{Q}^{r}\left(\mathcal{H}_{n_{2}}(\boldsymbol{Y})\right)$ to $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ can be computed by

$$
\begin{equation*}
\mathcal{Q}^{r}\left(\mathcal{H}_{n_{2}}(\boldsymbol{Y})\right)=\boldsymbol{A}_{1} \boldsymbol{S}_{1}^{r} \boldsymbol{B}_{1}^{T}, \tag{4}
\end{equation*}
$$

where $\mathcal{H}_{n_{2}}(\boldsymbol{Y})=\boldsymbol{A}_{1} \boldsymbol{S}_{1} \boldsymbol{B}_{1}{ }^{T}$ is the singular value decomposition of $\mathcal{H}_{n_{2}}(\boldsymbol{Y}) . \boldsymbol{A}_{1}, \boldsymbol{B}_{1}$, and $\boldsymbol{S}_{1}$ represent the left singular vectors, right singular vectors, and singular values, respectively. $\boldsymbol{S}_{1}^{r}$ keeps the largest $r$ singular values in $\boldsymbol{S}_{1}$ and sets all the others to zero. The corresponding normalized approximation error is computed by

$$
\begin{equation*}
\frac{\left\|\mathcal{Q}^{r}\left(\mathcal{H}_{n_{2}}(\boldsymbol{Y})\right)-\mathcal{H}_{n_{2}}(\boldsymbol{Y})\right\|_{F}}{\left\|\mathcal{H}_{n_{2}}(\boldsymbol{Y})\right\|_{F}}=\frac{\left\|\boldsymbol{S}_{1}^{r}-\boldsymbol{S}_{1}\right\|_{F}}{\left\|\boldsymbol{S}_{1}\right\|_{F}} \tag{5}
\end{equation*}
$$

where $\|.\|_{F}$ represents the Frobenious norm.

## B. Low-Rank Lifted Hankel Property in Nonlinear Dynamical System

When the underlying system is highly nonlinear such as immediately after a significant event, approximating a nonlinear system using a linear dynamical model usually requires a large order $r$. Thus, the corresponding $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ is no longer low-rank. The idea is to lift the measurements $\boldsymbol{y}_{i}$ to a higher dimensional space using a mapping function $\phi(\cdot): \mathbb{R}^{m} \rightarrow \mathbb{R}^{M}$, where $M$ is much larger than $m$ and can be infinite. As described in [28], there exists a mapping $\phi(\cdot)$ such that the nonlinear dynamical system can be a linear dynamical system in the lifted space. Let $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ be

$$
\mathcal{H}_{n_{2}}(\boldsymbol{Z})=\left[\begin{array}{cccc}
\boldsymbol{z}_{1} & \boldsymbol{z}_{2} & \ldots & \boldsymbol{z}_{n_{1}}  \tag{6}\\
\boldsymbol{z}_{2} & \boldsymbol{z}_{3} & \ldots & \boldsymbol{z}_{n_{1}+1} \\
\vdots & \vdots & \ldots & \vdots \\
\boldsymbol{z}_{n_{2}} & \boldsymbol{z}_{n_{2}+1} & \ldots & \boldsymbol{z}_{n}
\end{array}\right] \in \mathbb{R}^{M n_{2} \times n_{1}}
$$

where $\boldsymbol{z}_{i}=\phi\left(\boldsymbol{y}_{i}\right)$. The rank of $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ can be smaller than that of $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ for a proper $\phi$.

The rank-r approximation of $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ can be written as

$$
\begin{equation*}
\mathcal{Q}^{r}\left(\mathcal{H}_{n_{2}}(\boldsymbol{Z})\right)=\boldsymbol{A}_{2} \boldsymbol{S}_{2}^{r} \boldsymbol{B}_{2}^{T} \tag{7}
\end{equation*}
$$

where $\boldsymbol{S}_{2}^{r}$ contains the largest $r$ singular values of $\mathcal{H}_{n_{2}}(\boldsymbol{Z}), \boldsymbol{A}_{2}$ and $\boldsymbol{B}_{2}$ contain left and right singular vectors, respectively. The normalized approximation error of $\mathcal{Q}^{r}\left(\mathcal{H}_{n_{2}}(\boldsymbol{Z})\right)$ to $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ can be computed by

$$
\begin{equation*}
\frac{\left\|\mathcal{Q}^{r}\left(\mathcal{H}_{n_{2}}(\boldsymbol{Z})\right)-\mathcal{H}_{n_{2}}(\boldsymbol{Z})\right\|_{F}}{\left\|\mathcal{H}_{n_{2}}(\boldsymbol{Z})\right\|_{F}}=\sqrt{\frac{\sum_{i=r+1}^{n_{1}} \sigma_{i}^{2}}{\sum_{i=1}^{n_{1}} \sigma_{i}^{2}}} \tag{8}
\end{equation*}
$$

where $\sigma_{i}$ denotes the $i$ th largest singular value of $\boldsymbol{S}_{2}, \sigma_{i}$ cannot be computed directly from the SVD of $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ because $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ is unknown. Instead, one can compute $\mathcal{H}_{n_{2}}(\boldsymbol{Z})^{T} \mathcal{H}_{n_{2}}(\boldsymbol{Z})$ explicitly using the kernel trick [28] without knowing $\phi(\cdot)$. The $(i, j)$ th entry in $\mathcal{H}_{n_{2}}(\boldsymbol{Z})^{T} \mathcal{H}_{n_{2}}(\boldsymbol{Z})$ is computed by

$$
\left(\mathcal{H}_{n_{2}}(\boldsymbol{Z})^{T} \mathcal{H}_{n_{2}}(\boldsymbol{Z})\right)_{i, j}=\left[\begin{array}{lll}
\boldsymbol{z}_{i} & \ldots & \boldsymbol{z}_{i+n_{2}-1}
\end{array}\right]\left[\begin{array}{c}
\boldsymbol{z}_{j}  \tag{9}\\
\vdots \\
\boldsymbol{z}_{j+n_{2}-1}
\end{array}\right]
$$

$$
\begin{equation*}
=\sum_{p=0}^{p=n_{2}-1} \phi\left(\boldsymbol{y}_{i}\right)^{T} \phi\left(\boldsymbol{y}_{j}\right)=\sum_{p=0}^{p=n_{2}-1} \mathcal{K}_{Y Y}(i+p, j+p) \tag{10}
\end{equation*}
$$

where $\mathcal{K}_{Y Y}$ is the kernel function. The most popular kernel functions are the Gaussian kernel and the polynomial kernel. Reference [32] reports that the matrix completion methods with the Gaussian kernel perform better than the polynomial kernel. The Gaussian kernel corresponds to an infinite dimensional $\phi$. We employ the Gaussian kernel as follows,

$$
\begin{equation*}
\mathcal{K}_{Y Y}(i, j)=\phi\left(\boldsymbol{y}_{i}\right)^{T} \phi\left(\boldsymbol{y}_{j}\right)=\exp \left(-\frac{1}{2 c}\left\|\boldsymbol{y}_{i}-\boldsymbol{y}_{j}\right\|_{2}^{2}\right), \tag{11}
\end{equation*}
$$

where $c$ is a pre-defined scalar. One then solves the eigendecomposition of $\mathcal{H}_{n_{2}}(\boldsymbol{Z})^{T} \mathcal{H}_{n_{2}}(\boldsymbol{Z})$. The eigenvalues of $\mathcal{H}_{n_{2}}(\boldsymbol{Z})^{T} \mathcal{H}_{n_{2}}(\boldsymbol{Z})$ are $\sigma_{i}^{2}$, i.e.,

$$
\begin{equation*}
\mathcal{H}_{n_{2}}(\boldsymbol{Z})^{T} \mathcal{H}_{n_{2}}(\boldsymbol{Z})=\boldsymbol{B}_{2} \boldsymbol{S}_{2}^{2} \boldsymbol{B}_{2}^{T} \tag{12}
\end{equation*}
$$

Remark: When $\boldsymbol{Y}$ is obtained from a nonlinear dynamical system, to achieve the same normalized low-rank approximation error, it often requires a smaller rank to approximate the lifted Hankel matrix $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ (with a properly selected kernel function) than to approximate $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ with the same $n_{2}$. Therefore, the low-rank lifted Hankel property is more desirable in recovering PMU data in nonlinear dynamics.

To illustrate the low-rank lifted Hankel property, we consider a recorded generator trip event in New York State [24]. Fig. 1 shows the 10 seconds of voltage magnitude measurements in 11 channels at different locations. The data rate is 30 samples per second per channel. Let $\boldsymbol{Y} \in \mathbb{R}^{11 \times 300}$ contain all the measurements. Fig. 2(a) shows the normalized approximation errors of rank-r matrices to $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ and $\mathcal{H}_{n_{2}}(\boldsymbol{Y}) . c=200$ in (11). For example, the normalized error of rank-5 approximation to $\mathcal{H}_{10}(\boldsymbol{Z})$ is 0.0015 . In comparison, the matrix rank needs to be as least 10 to achieve a similar approximation error to $\mathcal{H}_{10}(\boldsymbol{Y})$. Moreover, with a large $n_{2}$, the dimension of $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ is very large but could be approximated by a matrix with a small rank. For


Fig. 1. The measurements of voltage magnitude [24].

(b)

Fig. 2. (a) The normalized approximation errors of the original Hankel matrices $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ and the corresponding lifted Hankel matrices $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$. (b) The normalized approximation errors of column-wise permuted Hankel matrices $\mathcal{H}_{n_{2}}(\overline{\boldsymbol{Y}})$ and the corresponding lifted Hankel matrices $\mathcal{H}_{n_{2}}(\overline{\boldsymbol{Z}})$.
instance, $\mathcal{H}_{20}(\boldsymbol{Y})$ is in $\mathbb{R}^{220 \times 281}$, and $\mathcal{H}_{20}(\boldsymbol{Z})$ is even higherdimensional due to the lifting. Still, $\mathcal{H}_{20}(\boldsymbol{Z})$ can be approximated by a rank-15 matrix with a normalized error of 0.00083 .

To illustrate that the low-rank (lifted) Hankel property is special for data from dynamical systems rather than an arbitrary matrix, we permute the columns in $\boldsymbol{Y}$ randomly and let $\overline{\boldsymbol{Y}}$ be the resulting matrix. Then $\boldsymbol{Y}$ and $\overline{\boldsymbol{Y}}$ have the same rank, but each row of $\overline{\boldsymbol{Y}}$ is no longer a time series. Fig. 2(b) shows the normalized approximation errors of $\mathcal{H}_{n_{2}}(\overline{\boldsymbol{Y}})$ and $\mathcal{H}_{n_{2}}(\overline{\boldsymbol{Z}})$, which are Hankel and lifted Hankel matrices constructed from $\overline{\boldsymbol{Y}}$. In contrast to Fig. 2(a), the approximation errors in Fig. 2(b) remain significant even when the rank is very large, because the low-rank (lifted) Hankel property does not hold for $\overline{\boldsymbol{Y}}$, which is not obtained from a dynamical system.

## III. Bayesian High-Rank Hankel Matrix Recovery (BA-NSDR) METHOD

The main idea of our proposed Ba-NSDR method is to estimate a matrix $\boldsymbol{Y}$ from partial observations $\boldsymbol{Y}^{o}$ such that the lifted


Fig. 3. An overall framework of the proposed method. The method maps the estimated data $\boldsymbol{Y}$ into a Hankel matrix $\boldsymbol{X}$ and then lifts $\boldsymbol{X}$ into higher dimensional space $\Phi(\boldsymbol{X}) . \Phi(\boldsymbol{X})$ is decomposed with a lifted factor $\Phi(\boldsymbol{U})$, and the coefficient matrix $\boldsymbol{V}$.

Hankel matrix of $\boldsymbol{Y}$ is low-rank. To simplify representation, given $n_{2}$, we use $\boldsymbol{X}$ and $\Phi(\boldsymbol{X})$ to denote $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ and $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$, respectively. With a bit of abuse of notation, $\Phi(\boldsymbol{A})$ means dividing each column of the matrix $\boldsymbol{A}$ into multiple vectors in $\mathbb{R}^{m}$ and lifting each vector to $\mathbb{R}^{M}$ by the lifting function $\phi$. Assuming $\Phi(\boldsymbol{X})$ is rank $K$, we view $\Phi(\boldsymbol{X})$ as the product of two matrix factors, $\Phi(\boldsymbol{U})$ in $\mathbb{R}^{M n_{2} \times K}$ and $\boldsymbol{V}$ in $\mathbb{R}^{K \times n_{1}}$, where $\Phi(\boldsymbol{U})$ is a lifted matrix to $\mathbb{R}^{M n_{2} \times K}$ from a matrix $\boldsymbol{U}$ in $\mathbb{R}^{m n_{2} \times K}$. Because the rank $K$ is small, the degree of freedom $K\left(m n_{2}+n_{1}\right)$ is much less than $m n$, the ambient dimension of $\boldsymbol{Y}$. Therefore, we could accurately recover $\boldsymbol{U}, \boldsymbol{V}$, and thus $\boldsymbol{Y}$ from partial observations that contain bad data. Note that every column of $\Phi(\boldsymbol{X})$ includes all data from $m$ channels in $n_{2}$ consecutive steps. Then as long as there exist $K$ reliable measurements in all channels in a length $-n_{2}$ window, all the remaining measurements in that window can be accurately recovered. Thus, by exploiting the low-rank lifted Hankel property, one can recover data losses/errors in all $m$ channels consecutively.

As a Bayesian approach, Ba-NSDR first imposes a prior distribution on $\boldsymbol{Y}$ and $\Phi(\boldsymbol{X})$ (Section III-A) and then computes the posterior distribution based on partial observations $\boldsymbol{Y}^{o}$ (Section III-B). Ba-NSDR then uses the posterior distribution of $\boldsymbol{Y}$ to estimate the data and compute the uncertainty index that reflects the estimation accuracy (Section III-C). Section III-D discusses the parameter selection.

## A. Proposed Probabilistic Model

Equations (13) to (20) show our hierarchical probabilistic model of the prior distributions (Fig. 3 shows the visualization). Readers can refer to [35] for prerequisites of the proposed Bayesian model. The latent variables are inferred using observations based on this probabilistic model. Equation (13) is a probabilistic version of equation (2), where $Y_{i, j}$ can be written as the Hankel inverse $\left(\mathcal{H}^{\dagger} \boldsymbol{X}\right)_{i, j}$, where the Hankel inverse operator $\mathcal{H}^{\dagger}$ is defined in (35) in the supplementary materials. $\Phi(\boldsymbol{X})_{. q} \in \mathbb{R}^{M n_{2}}$ is the $q$ th column of $\Phi(\boldsymbol{X})$. The


Fig. 4. The Graphical model of the proposed Bayesian high-rank Hankel matrix completion method.
prior knowledge of rank $K$ might be unavailable. Ba-NSDR sets the initial $K$ as a relatively large number and gradually prunes the basis based on learned coefficients $\boldsymbol{V}$.

The prior distributions of $\boldsymbol{U}_{. k}, \boldsymbol{X}_{. q}$, and $\boldsymbol{V}_{. q}$ are drawn from multivariate Gaussian distributions $\mathcal{N}\left(\mathbf{0}, \gamma_{u}^{-1} \boldsymbol{I}_{K}\right)$, $\mathcal{N}\left(\mathbf{0}, \gamma_{x}^{-1} \boldsymbol{I}_{K}\right)$, and $\mathcal{N}\left(\mathbf{0}, \gamma_{v}^{-1} \boldsymbol{I}_{K}\right)$, respectively. $\boldsymbol{I}_{m n_{2}}$ is an $m n_{2}$ by $m n_{2}$ identity matrix. $\gamma_{u}, \gamma_{x}$, and $\gamma_{v}$ are three pre-defined scalars. Each element in the error matrix $\boldsymbol{E}$ is drawn from a Gaussian distribution $\mathcal{N}\left(0, \beta_{i, j}^{-1}\right)$. Each element in the noise matrix $\boldsymbol{N}$ is drawn from a Gaussian distribution $\mathcal{N}\left(0, \gamma_{y}^{-1}\right)$. The Gamma prior distribution is placed on $\gamma_{y}$ and $\beta_{i, j}$, following parameters $\left(e_{0}, f_{0}\right)$ and $\left(g_{0}, h_{0}\right)$, respectively. The mathematical definition of the Gamma distribution is shown in the supplementary material. The conjugate priors are placed on $\boldsymbol{V}_{. q}, \gamma_{y}, E_{i, j}$, and $\beta_{i, j}$ to derive analytical solutions of posterior distributions. The graphical representation of the proposed probabilistic model is shown in Fig. 4.

For all $q=1,2,3, \ldots, n_{1}$, and $k=1,2,3, \ldots, K$,

$$
\begin{equation*}
Y_{i, j}^{o} \sim \mathcal{N}\left(\left(\mathcal{H}^{\dagger} \boldsymbol{X}\right)_{i, j}+E_{i, j}, \frac{1}{\gamma_{y}}\right) \quad(i, j) \in \Omega \tag{13}
\end{equation*}
$$

$$
\begin{align*}
\Phi(\boldsymbol{X})_{. q} & \sim \mathcal{N}\left(\Phi(\boldsymbol{U}) \boldsymbol{V}_{. q}, \frac{1}{\gamma_{\epsilon}} \boldsymbol{I}_{m n_{2}}\right)  \tag{14}\\
\boldsymbol{U}_{. k} & \sim \mathcal{N}\left(0, \frac{1}{\gamma_{u}} \boldsymbol{I}_{m n_{2}}\right)  \tag{15}\\
\boldsymbol{X}_{. q} & \sim \mathcal{N}\left(0, \frac{1}{\gamma_{x}} \boldsymbol{I}_{m n_{2}}\right)  \tag{16}\\
\boldsymbol{V}_{. q} & \sim \mathcal{N}\left(0, \frac{1}{\gamma_{v}} \boldsymbol{I}_{K}\right)  \tag{17}\\
\gamma_{y} & \sim \Gamma\left(e_{0}, f_{0}\right)  \tag{18}\\
E_{i, j} & \sim \mathcal{N}\left(0, \frac{1}{\beta_{i, j}}\right) \quad(i, j) \in \Omega  \tag{19}\\
\beta_{i, j} & \sim \Gamma\left(g_{0}, h_{0}\right) \tag{20}
\end{align*}
$$

## B. Variational Inference for Approximating the Posterior Distributions

To simplify representation, we denote $\boldsymbol{\Theta}=\left\{\boldsymbol{U}_{. k}, \boldsymbol{V}_{. q}\right.$, $\boldsymbol{X}_{. q}, \gamma_{y}, E_{i, j}, \beta_{i, j}, q=1,2,3, \ldots, n_{1}, k=1,2,3, \ldots, K,(i, j)$ $\in \Omega\}$ as the set of all the latent variables. Let $\boldsymbol{\Theta}_{i}$ denote one arbitrary variable in $\Theta$. Given partial observation $\boldsymbol{Y}_{\Omega}^{o}$, the goal is to compute the posterior distribution $p\left(\boldsymbol{\Theta}, \boldsymbol{Y} \mid \boldsymbol{Y}_{\Omega}^{o}\right)$. Based on Bayes' theorem,

$$
\begin{equation*}
p\left(\boldsymbol{\Theta}, \boldsymbol{Y} \mid \boldsymbol{Y}_{\Omega}^{o}\right)=\frac{p\left(\boldsymbol{\Theta}, \boldsymbol{Y}, \boldsymbol{Y}_{\Omega}^{o}\right)}{p\left(\boldsymbol{Y}_{\Omega}^{o}\right)} \tag{21}
\end{equation*}
$$

Computing (21) requires marginalizing out all the latent variables, which is usually intractable.

As a popular approach to approximate the complicated posterior distribution, the mean field variational inference [35] employs a simple distribution $q(\boldsymbol{\Theta})$ to approximate $p\left(\boldsymbol{\Theta}, \boldsymbol{Y} \mid \boldsymbol{Y}_{\Omega}^{o}\right)$. The mean field assumption assumes that each element in $\Theta$ is mutually independent. Then $q(\boldsymbol{\Theta})$ can be factorized as the product of each element, i.e.,

$$
\begin{align*}
q(\boldsymbol{\Theta})= & \prod_{k=1}^{K} q\left(\boldsymbol{U}_{. k}\right) \prod_{q=1}^{n_{1}} q\left(\boldsymbol{V}_{. q}\right) q\left(\boldsymbol{X}_{. q}\right) \\
& \times \prod_{(i, j) \in \boldsymbol{\Omega}} q\left(E_{i, j}\right) q\left(\beta_{i, j}\right) q\left(\gamma_{y}\right) \tag{22}
\end{align*}
$$

The best $q(\boldsymbol{\Theta})$ to approximate $p\left(\boldsymbol{\Theta}, \boldsymbol{Y}_{\Omega} \mid \boldsymbol{Y}_{\Omega}^{o}\right)$ is found by minimizing the Kullback-Leibler (KL) divergence, which measures the similarity of two probabilistic distributions. Specifically,

$$
\begin{align*}
q(\boldsymbol{\Theta}) & =\underset{q(\boldsymbol{\Theta})}{\arg \min } \mathbb{K} \mathbb{L}\left(q(\boldsymbol{\Theta}) \| p\left(\boldsymbol{\Theta}, \boldsymbol{Y} \mid \boldsymbol{Y}_{\Omega}^{o}\right)\right) \\
& =\underset{q(\boldsymbol{\Theta})}{\operatorname{argmax}} \mathbb{E}\left[\ln p\left(\boldsymbol{\Theta}, \boldsymbol{Y}, \boldsymbol{Y}_{\Omega}^{o}\right)\right]-\mathbb{E}[\ln q(\boldsymbol{\Theta})] . \tag{23}
\end{align*}
$$

where $\mathbb{K} \mathbb{L}(x \| y)$ denotes the $\operatorname{KL}$ divergence of distribution $x$ and $y$, and $\mathbb{E}$ is the expectation over $q(\boldsymbol{\Theta})$. The second equality follows from the definition of KL divergence and removes the term unrelated to $q(\boldsymbol{\Theta})$.

Because it is intractable to solve (23), a typical approach is to optimize each variable $\Theta_{i}$ in $\Theta$ via solving (23) while keeping
all other variables fixed using the most recent distributions.

$$
\begin{align*}
& q\left(\boldsymbol{\Theta}_{i}\right) \\
& =\underset{q\left(\boldsymbol{\Theta}_{i}\right)}{\arg \max \left(\int q\left(\boldsymbol{\Theta}_{i}\right) \mathbb{E}_{q\left(\boldsymbol{\Theta} \backslash \boldsymbol{\Theta}_{i}\right)}\left[\ln p\left(\boldsymbol{\Theta}, \boldsymbol{Y}, \boldsymbol{Y}_{\Omega}^{o}\right)\right] d\left(\boldsymbol{\Theta}_{i}\right)\right.} \\
& \left.-\int q\left(\boldsymbol{\Theta}_{i}\right) \ln q\left(\boldsymbol{\Theta}_{i}\right) d \boldsymbol{\Theta}_{i}\right) \tag{24}
\end{align*}
$$

where $\mathbb{E}_{q\left(\boldsymbol{\Theta} \backslash \Theta_{i}\right)}$ represents that the expectation is taken with respect to all the latent variables excluding $\boldsymbol{\Theta}_{i}$. These approximate distributions of variational inference finally converge to a local optimum of (23) [35], [36].

Because the conjugate priors are placed on latent variables $\boldsymbol{V}_{. q}, E_{i, j}, \beta_{i, j}$ and $\gamma_{y}$, (24) has analytical solutions for these variables. Please refer to steps (I), (IV), (V), and (VI) in supplementary materials for the respective updating equations. Because $\boldsymbol{U}_{. k}$ and $\boldsymbol{X}_{. q}$ are lifted to a higher dimensional space via the kernel method, (24) does not have analytical forms for these variables. To solve (24), we assume $\boldsymbol{U}_{. k}$ and $\boldsymbol{X}_{. q}$ are drawn from Gaussian distributions, and then the problem is simplified to find the corresponding mean and the variance of each variable. Then the reparameterization trick [37] is employed to differentiate and optimize the objective in (24) with respect to the mean and variance, respectively. Please refer to steps (II) and (III) in supplementary materials for the updating equations of $\boldsymbol{U}_{. k}$ and $\boldsymbol{X}_{. q}$.

Computing the objective function in (24) for $\boldsymbol{V}_{. q}, \boldsymbol{U}_{. k}$ and $\boldsymbol{X}_{. q}$ requires computing the inner product of the lifting function. We employ three Gaussian kernels $\mathcal{K}_{X X}, \mathcal{K}_{X U}$ and $\mathcal{K}_{U U}$ in (25)-(27) when updating $\boldsymbol{V}_{. q}, \boldsymbol{X}_{. q}$, and $\boldsymbol{U}_{. k}$, respectively.

$$
\begin{equation*}
\mathcal{K}_{X X}(p, q)=\Phi(\boldsymbol{X})_{. p}^{T} \Phi(\boldsymbol{X})_{. q}=\exp \left(-\frac{1}{2 c_{1}}\left\|\boldsymbol{X}_{. p}-\boldsymbol{X}_{. q}\right\|_{2}^{2}\right) \tag{25}
\end{equation*}
$$

$$
\begin{equation*}
\mathcal{K}_{X U}(q, k)=\Phi(\boldsymbol{X})_{. q}^{T} \Phi(\boldsymbol{U})_{. k}=\exp \left(-\frac{1}{2 c_{2}}\left\|\boldsymbol{X}_{. q}-\boldsymbol{U}_{. k}\right\|_{2}^{2}\right) \tag{26}
\end{equation*}
$$

$$
\begin{equation*}
\mathcal{K}_{U U}(i, j)=\Phi(\boldsymbol{U})_{. i}^{T} \Phi(\boldsymbol{U})_{. j}=\exp \left(-\frac{1}{2 c_{3}}\left\|\boldsymbol{U}_{. i}-\boldsymbol{U}_{. j}\right\|_{2}^{2}\right) \tag{27}
\end{equation*}
$$

where $c_{1}, c_{2}$ and $c_{3}$ are pre-defined scalars.
Initialization: Each entry in $\boldsymbol{U}$ is initialized from a Gaussian distribution $\mathcal{N}(0,1) . \boldsymbol{V}$ is initialized as an all-zero matrix. All the elements in initial variances for $\boldsymbol{U}_{. k}$ and $\boldsymbol{X}_{. q}$ are set as $\exp (-2)$. The initialization $\overline{\boldsymbol{X}}^{0}$ of $\boldsymbol{X}$ is initialized as the rank- $r$ approximation to $\mathcal{H}_{n_{2}}\left(\boldsymbol{Y}^{o}\right)$, where the missing entries are set as zero. The initial $\boldsymbol{E}$ is set as $\boldsymbol{Y}^{o}-\mathcal{P}_{\boldsymbol{\Omega}}\left(\mathcal{H}^{\dagger} \overline{\boldsymbol{X}}^{0}\right)$. The $\gamma_{y}$ is initialized as $10^{6}$.

Estimating the Rank of the Lifted Hankel Matrix: Because the actual rank of $\Phi(\boldsymbol{X})$ is unknown, one selects $K$ that is guaranteed to be larger than the actual rank. The deterministic methods such as [32] require $K$ to be an accurate estimation of the rank and often overfit when $K$ is larger than the actual rank. Here we propose to estimate the rank and remove the redundant factor by thresholding the entries in $\mathbb{E}[\boldsymbol{V}]$. If the sum of absolute
values of $\mathbb{E}\left[V_{k q}\right]$ for all $q$ is less than a threshold (e.g., $10^{-2}$ ), the algorithm removes the $k$ th column in $\mathbb{E}[\boldsymbol{U}]$, the $k$ th row in $\mathbb{E}[\boldsymbol{V}]$, and reduces the rank $K$ by one. That is because the $k$ th column in $\mathbb{E}[\boldsymbol{U}]$ is not selected to represent $\Phi(\boldsymbol{X})$ and is no longer needed. Therefore, our method is robust to the initial rank and can effectively infer the actual rank.

Estimating the Sparsity of the Error Matrix $\boldsymbol{E}$ : Reference [38] shows that the Gaussian distribution with Gamma priors promotes the sparsity of $\boldsymbol{E}$. We can make $\mathbb{E}[\boldsymbol{E}]$ sparser through thresholding, because significant errors do not happen frequently. When entries in $\mathbb{E}[\boldsymbol{E}]$ are very small (e.g., $10^{-1}$ ), the corresponding entries are set as zeroes.

Convergence Criteria: Let $\overline{\boldsymbol{X}}^{t}$ and $\overline{\boldsymbol{X}}^{t-1}$ denote the estimation of $\boldsymbol{X}$ at the $t$ th and $t-1$ th iteration, respectively. The algorithm terminates if $\frac{\left\|\overline{\boldsymbol{X}}^{t}-\overline{\boldsymbol{X}}^{t-1}\right\|_{F}}{\left\|\overline{\boldsymbol{X}}^{t-1}\right\|_{F}}<\xi$ where $\xi$ is a pre-determined threshold, or if the maximum iterations $T_{\max }$ is reached.

Missing Data Recovery Only: The algorithm can be simplified when the objective is to recover missing data only, assuming the observations do not contain bad data. Equations (19) and (20) in the prior model characterize the bad data distribution and can be removed. One can also skip steps (IV) and (V) (in the supplementary materials) that update $E_{i, j}$ and $\beta_{i, j}$.

Computational Complexity: The computational complexity per iteration is $\mathcal{O}\left(L m n_{2} n_{1} K t^{\max }\right)$, where $L$ and $t^{\max }$ are the Monte-Carlo samples and maximum iterations of inner loops, respectively, when computing $\boldsymbol{U}$ and $\boldsymbol{X}$. Thus, the computational complexity scales at most linearly in the size of the Hankel matrix. The details of derivation are provided in Section F in the supplementary materials. Our algorithm is a block processing method and is most suitable for offline data recovery. It could possibly be used for online processing with sufficient computational power.

## C. Data Recovery and Uncertainty Index

With the computed posterior distributions, we use the mean of the distribution of $Y_{i, j}$ as an estimate of the corresponding entry in $\boldsymbol{Y}$ for every $i=1, \ldots, m$, and $j=1, \ldots, n$. The variance of $Y_{i, j}$ is employed to estimate the accuracy of data recovery. Because the mean and variance do not have closed-form solutions, the Monte Carlo integration [39] is employed to compute them approximately. The predictive mean is derived as follows:

$$
\begin{equation*}
\mathbb{E}\left[Y_{i, j}\right] \approx \frac{1}{J} \sum_{l=1}^{J}\left(\mathcal{H}^{\dagger} \boldsymbol{X}^{(l)}\right)_{i, j} \quad \boldsymbol{X}^{(l)} \sim q\left(\boldsymbol{X} \mid \boldsymbol{Y}_{\Omega}^{o}\right) \tag{28}
\end{equation*}
$$

where $J$ is the number of Monte-Carlo samples. Each $\boldsymbol{X}^{(l)}$ is sampled from learned posterior distributions. The predictive variance is computed by:
$\operatorname{Var}\left[Y_{i, j}\right]=\mathbb{E}\left[Y_{i, j}^{2}\right]-\mathbb{E}\left[Y_{i, j}\right]^{2}$
$\approx \frac{1}{J} \sum_{l=1}^{J} \frac{1}{\gamma_{y}^{(l)}}+\frac{1}{J} \sum_{l=1}^{J}\left(\mathcal{H}^{\dagger} \boldsymbol{X}^{(l)}\right)_{i, j}^{2}-\left(\frac{1}{J} \sum_{l=1}^{J}\left(\mathcal{H}^{\dagger} \boldsymbol{X}^{(l)}\right)_{i, j}\right)^{2}$,
where each $\gamma_{y}^{(l)}$ is sampled from learned posterior distribution $q\left(\gamma_{y} \mid \boldsymbol{Y}_{\Omega}^{o}\right)$. We use the average variance as an uncertainty index of the data estimation, i.e.,

$$
\begin{equation*}
U_{\mathrm{index}}=\left(\sum_{i=1}^{m} \sum_{j=1}^{n} \operatorname{Var}\left[Y_{i, j}\right]\right) /(m n) \tag{30}
\end{equation*}
$$

A higher average variance leads to a larger uncertainty index. That means the algorithm is less confident about the recovery results.

## D. Parameter Selection

The prior distributions (18) and (20) require setting parameters $\left(e_{0}, f_{0}\right)$ and $\left(g_{0}, h_{0}\right)$. When $e_{0}$ is fixed, a larger $f_{0}$ corresponds to a smaller $\gamma_{y}$, which in turn increases the variance $1 / \gamma_{y}$ of the noise $\boldsymbol{N}$. When $h_{0}$ is fixed, a larger $g_{0}$ corresponds to a larger $\beta_{i, j}$, which in turn decreases the value of $E_{i, j}$. Note that $\left(e_{0}, f_{0}\right)$ and $\left(g_{0}, h_{0}\right)$ have a minor impact on the recovery results. Another important parameter is the Hankel size $n_{2}$. With a larger $n_{2}$, the method can recover consecutive data losses and errors for all channels for a longer time window (close to $n_{2}$ time steps). On the other hand, increasing $n_{2}$ leads to a higher computational cost. In our experiments, setting $n_{2}$ as at most 80 is sufficient to obtain accurate recovery performance. In Section IV-C3, we show that Ba-NSDR is not sensitive to these parameter selections.

## IV. Numerical Experiments

## A. Experimental Setup

We compare our proposed Ba-NSDR approach with the following nine methods: the Bayesian robust Hankel matrix completion method (BRHMC) in [27], the Bayesian robust Hankel matrix completion method employing the sliding window (BRHMC-S), the Bayesian Hankel matrix completion method (BHMC) in [27], the Bayesian Hankel matrix completion method employing the sliding window (BHMC-S), the deterministic kernel-based matrix completion method (KMC) in [32], the deterministic Hankel matrix completion method (AM-FIHT) in [25], the deterministic robust Hankel matrix completion method (SAP) in [26], the deterministic streaming data recovery method (SDR) in [24], the deterministic streaming data recovery method considering the nonlinear dynamics (SDR-K) in [28], The streaming methods "SDR" and "SDR-K" require that the observations in the first time window contain no missing and bad data, which is one disadvantage compared with offline methods. In the following experiments, we do not include missing and bad data in the first time window of these two methods to make a fair comparison.

Some parameters of Ba-NSDR are set as follows for all experiments if not otherwise stated: $\gamma_{\epsilon}=10^{5}, \gamma_{v}=10^{2}$, $J=$ $50, L=1, \gamma_{x}=\gamma_{u}=1, e_{0}=10^{-6}, f_{0}=10^{-4}, g_{0}=1, h_{0}=$ $10^{-6}, \xi=10^{-4} \cdot \lambda_{1}=10, \lambda_{2}=\lambda_{4}=0.1, \lambda_{3}=1 . T_{\max }=100$. $t_{1}^{\max }=t_{2}^{\max }=t_{3}^{\max }=t_{4}^{\max }=100$. The experiments are conducted on Matlab 2019 with a desktop with 3.1 GHz Intel i9-9900 and 32 GB memory. Fig. 5 shows three modes of


Fig. 5. Three modes of missing/bad data generation. "M" stands for missing data. "B" stands for bad data.
missing/bad data considered in this experiment. For example, M3 represents Mode 3 of missing data. B2 represents Mode 2 of bad data.

- Mode 1: Missing/bad entries independently and randomly distribute across all channels and time instants.
- Mode 2: Missing/bad entries distribute across all channels, and the time instants are randomly selected.
- Mode 3: Missing/bad entries distribute across all channels. The time instants are consecutive instants and the starting instant is randomly selected.
Evaluation Metrics: The Normalized Estimation Error (NEE) is employed to evaluate the data recovery performance. The NEE is defined as

$$
\begin{equation*}
\mathrm{NEE}=\|\hat{\boldsymbol{Y}}-\boldsymbol{Y}\|_{F} /\|\boldsymbol{Y}\|_{F} \tag{31}
\end{equation*}
$$

where $\hat{\boldsymbol{Y}}$ in $\mathbb{R}^{m \times n}$ is the estimate and $\boldsymbol{Y}$ in $\mathbb{R}^{m \times n}$ is the groundtruth data. Note that the computation of NEE requires groundtruth data and can only be used for evaluation. When there is no ground-truth data provided, the uncertainty index reflects the estimation accuracy. We will report both the uncertainty index and NEE in the following experiments.

## B. Performance on Synthetic Datasets

1) Dataset Generation: We first evaluated the data recovery performance on synthetic data where each row of $\boldsymbol{Y}$ is a weighted sum of $r$ time-varying damping noisy sinusoids. Each entry $Y_{i, j}$ in $\boldsymbol{Y}$ is generated by
$Y_{i, j}=\sum_{k=1}^{r} b_{k, j} e^{-a_{i} t_{j}} \sin \left(2 \pi f_{k, j} t_{j}\right) \quad i=1, \ldots, m, j=1, \ldots, n$,
where $f_{k, j}$ is the time-varying frequency, $b_{k, j}$ is the time-varying amplitude of the $k$ th sinusoid. The general form of time-varying frequency and amplitude can characterize the dynamic transitions during a significant disturbance in power systems. The frequency $f_{k, j}$ is randomly selected from $(100,102)$. The amplitude $b_{j, k}$ is randomly selected from (1,1.3). $r=2, a_{1}=$ $30, a_{2}=40, a_{3}=35$. The generated matrix $\boldsymbol{Y}$ has three rows

TABLE I
The Recovery Error and the Uncertainty Index by BA-NSDR on M2 Missing Data of Synthetic Data

| Missing rate \% | 5 | 15 | 25 | 35 |
| :---: | :---: | :---: | :---: | :---: |
| NEE | 0.028 | 0.044 | 0.057 | 0.071 |
| $\mathrm{U}_{\text {index }}$ | $1.2 \times 10^{-3}$ | $1.6 \times 10^{-3}$ | $2.0 \times 10^{-3}$ | $2.9 \times 10^{-3}$ |
| Missing rate \% | 45 | 55 | 65 |  |
| NEE | 0.10 | 0.28 | 0.54 |  |
| $\mathrm{U}_{\text {index }}$ | $3.8 \times 10^{-3}$ | $7.2 \times 10^{-3}$ | $8.3 \times 10^{-2}$ |  |

and 300 columns. Fig. 7 shows the normalized approximation errors of rank- $r$ matrices to the Hankel matrix $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ and the lifted Hankel matrix $\mathcal{H}_{n_{2}}(\boldsymbol{Z}) . c=200$ in (11). One can see that it requires a much smaller rank to approximate $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ than $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ with the same normalized approximation error. For example, a rank-2 approximation to $\mathcal{H}_{10}(\boldsymbol{Z})$ is 0.019 , while it requires at least rank-27 to achieve a similar error to approximate $\mathcal{H}_{10}(\boldsymbol{Y})$.

We used a simple signal with nonlinear dynamics in (32) to verify the performance of our algorithm. The signals in (32) simulate the nonlinear dynamics from a nonlinear dynamical system. As stated in reference [40], a linear dynamical system should hold homogeneity property and additive property at the same time. Therefore, if an input is a sinusoidal signal $x(t)=\sin (2 \pi f t)$, where $f$ is the frequency and $t$ is the time instant, the output of a linear dynamical system should be $y(t)=A \sin (2 \pi f t+\alpha)$, where $A$ is a scaling amplitude and $A$ is a scalar, and $\alpha$ is the time-shifting phase. Because the amplitude in (32) is time-varying, the resulting signals are not generated from a linear dynamical system but from a nonlinear system.
2) Recovery Performance: Some parameters of Ba-NSDR are: $c_{2}=c_{3}=200, \xi=10^{-4}, K=50, T_{\max }=150 . n_{2}=$ 20 for all cases except that $n_{2}=30$ for M3 missing mode (Fig. 6(c)-(f)). The results are averaged over 10 trails. Fig. 6(a)(c) compare the missing data recovery performance of Ba-NSDR with KMC, SDR-K, AM-FIHT, BHMC-S, and BHMC on three missing data modes. Ba-NSDR achieves the lowest recovery error among all the methods. Specifically, the conventional kernelbased method KMC does not consider the Hankel structure and, thus, performs poorly on M2 and M3 modes. Deterministic Hankel-based method AM-FIHT and Bayesian Hankel-based methods BHMC, BHMC-S, approximate the data generated from nonlinear dynamical systems using linear dynamical systems and, thus, cannot accurately recover the highly nonlinear components. SDR-K employs the low-rank lifted Hankel property to characterize nonlinear dynamics and performs better than all other methods except our method Ba-NSDR. SDR-K does not provide any uncertainty index and cannot handle bad data. Moreover, SDR-K is sensitive to parameter selections, especially the selection of rank. Table I shows the NEE and the corresponding uncertainty indices when the missing data follow M2 mode. The uncertain index increases when the recovery error increases. This indicates that the uncertainty index is able to differentiate reliable estimations from unreliable estimations.

Fig. 6(d)-(f) compare the data recovery performance of BaNSDR with SAP, BRHMC-S and BRHMC when data contain both missing and bad data. Except for Ba-NSDR, all other


Fig. 6. Comparison of Ba-NSDR with other methods. (a)-(c) show the missing data recovery results with three missing modes. (d)-(f) show the recovery results with both missing and bad data.


Fig. 7. Low-rank approximations to Hankel matrices $\mathcal{H}_{n_{2}}(\boldsymbol{Y})$ and the corresponding lifted Hankel matrices $\mathcal{H}_{n_{2}}(\boldsymbol{Z})$ for synthetic data.
methods do not characterize nonlinear dynamics. One can see from Fig. 6(d)-(f) that Ba-NSDR performs the best among all the methods. Note that the signal in (32) does not include the phase for simplicity. We also tested the performance of our algorithm on a sinusoid with a time-varying phase, and the recovery results are shown in Fig. 11 in supplementary materials. Our method achieves similar performance as the results in Fig. 6.

## C. Performance on Practical PMU Dataset

We then conducted the experiments on the recorded dataset as shown in Fig. 1 in Central New York Power System. ${ }^{1}$ The PMU data type is voltage in rectangular coordinates. The proposed method can also be easily extended to other data types such as current and frequency. Observations in all channels are available in this 10 -second window and are treated as ground-truth data. We remove some data points and add bad data following different

[^1]TABLE II
the Recovery Performance of Recorded PMU Data on 6.7\% M3 Mode

| Method | Ba-NSDR | BHMC | BHMC-S | AM-FIHT | SDR-K |
| :---: | :---: | :---: | :---: | :---: | :---: |
| NEE | $8.3 \times 10^{-4}$ | $5.6 \times 10^{-3}$ | $3.0 \times 10^{-3}$ | $6.0 \times 10^{-3}$ | $2.1 \times 10^{-3}$ |
| NEE $_{2-4}$ | $1.9 \times 10^{-3}$ | $1.2 \times 10^{-2}$ | $6.6 \times 10^{-3}$ | $1.3 \times 10^{-2}$ | $4.7 \times 10^{-3}$ |
| Time(sec.) | 28.5 | 13.1 | 281.5 | 0.30 | 0.45 |

TABLE III
the Recovery Performance of Recorded PMU Data on 5\% M1 and 3.7\% B3 MODE

| Method | Ba-NSDR | BRHMC | BRHMC-S | SAP | SDR |
| :---: | :---: | :---: | :---: | :---: | :---: |
| NEE | $9.8 \times 10^{-4}$ | $7.1 \times 10^{-3}$ | $3.6 \times 10^{-3}$ | $6.0 \times 10^{-3}$ | $5.6 \times 10^{-3}$ |
| NEE $_{2-4}$ | $1.9 \times 10^{-3}$ | $1.5 \times 10^{-2}$ | $7.9 \times 10^{-3}$ | $1.3 \times 10^{-2}$ | $1.2 \times 10^{-2}$ |
| Time(sec.) | 19.5 | 2.2 | 276.8 | 0.054 | 0.13 |

patterns. The recovered data are evaluated by comparing them with the ground-truth data.

1) Recovery Performance: We first evaluated our method on two case studies.

- Case 1: 6.7\% data are removed following Mode M3. The length of M3 missing data is 20 consecutive time instants, which correspond to 0.67 seconds.
- Case 2: 5\% data are removed following Mode M1 and 3.7\% bad data following Mode B3 are added. The length of B3 bad data is 10 consecutive time instants, which correspond to 0.33 seconds. The bad data is randomly sampled from (0.1, 0.4).

The parameter setting of Ba-NSDR is as follows. The initial rank is set as 10. $n_{2}=30, c_{2}=c_{3}=40, f_{0}=10^{-6}$ in Case 1. $n_{2}=80, n_{2}=5, c_{2}=c_{3}=7, f_{0}=10^{-4}$ in Case 2.

Figs. 8 and 9 compare the recovery performance of Ba-NSDR with other methods on Case 1 and Case 2, respectively. BaNSDR can accurately recover the nonlinear dynamics during the event and clearly outperform all the existing methods. Tables II and III report the NEE over the whole ten-second window, the


Fig. 8. The recovery performance on $6.7 \% \mathrm{M} 3$ missing data. (a) the observed data, (b) the estimated data by the proposed Ba-NSDR method, (c) the estimated data by the BHMC method, (d) the estimated data by the BHMC-S method, (e) the estimated data by the AM-FIHT method, (f) the estimated data by the SDR-K method.


Fig. 9. The recovery performance on $5 \% \mathrm{M} 1$ missing data and $3.7 \% \mathrm{~B} 3$ bad data. (a) the observed data, (b) the estimated data by the proposed Ba-NSDR method, (c) the estimated data by the BRHMC method, (d) the estimated data by the BRHMC-S method, (e) the estimated data by the SAP method, (f) the estimated data by the SDR method.

NEE of a window between 2-4 seconds where missing data occur, denoted by $\mathrm{NEE}_{2-4}$, and the computational time of these methods over the whole ten-second window. Ba-NSDR achieves a great balance of recovery accuracy and computational cost. AM-FIHT, SAP, SDR, and SDR-K are computationally efficient, but their recovery performances are worse than our method. BHMC-S and BRHMC-S truncate the data into small windows and approximate each window using low-rank Hankel matrices and thus are much more computationally expensive than other methods.

The major disadvantage of our method is that it is more computationally expensive than the deterministic low-rank Hankel methods. However, we can see from Tables II and III that the
proposed Ba-NSDR method achieves a great balance of recovery accuracy and computational cost. Moreover, the proposed probabilistic framework is able to model the uncertainty of the recovery results, while other works cannot provide such an uncertainty index.
2) Uncertainty Modeling: One major advantage of BaNSDR over existing PMU data recovery methods is that it provides an uncertainty index, which can be employed to evaluate the reliability of the recovery results. Table IV shows the recovery performance and the corresponding uncertainty index on $5 \% \mathrm{~B} 1$ with varying missing data percentages of mode M2. Table V shows the recovery performance and the corresponding uncertainty index on 5\% M2 with varying bad data percentages

TABLE IV
The Recovery Error and the Uncertainty Index on 5\% B1 With Varying Missing Data Percentage of M2

| Missing rate | 5 | 15 | 25 | 35 | 45 |
| :---: | :---: | :---: | :---: | :---: | :---: |
| NEE | 0.0019 | 0.0037 | 0.0057 | 0.0060 | 0.18 |
| $\mathrm{U}_{\text {index }}$ | $2.6 \times 10^{-5}$ | $4.8 \times 10^{-5}$ | $1.5 \times 10^{-4}$ | $4.5 \times 10^{-4}$ | $1.1 \times 10^{-2}$ |

TABLE V
The Recovery Error and the Uncertainty Index on 5\% M2 With Varying Bad Data Percentage of B1

| Bad rate | 5 | 15 | 25 | 35 | 45 |
| :---: | :---: | :---: | :---: | :---: | :---: |
| NEE | 0.0019 | 0.0091 | 0.016 | 0.017 | 0.032 |
| $\mathrm{U}_{\text {index }}$ | $2.6 \times 10^{-5}$ | $5.8 \times 10^{-5}$ | $7.0 \times 10^{-5}$ | $8.3 \times 10^{-4}$ | $6.2 \times 10^{-3}$ |

TABLE VI
The Impact of $f_{0}\left(e_{0}\right.$ IS Fixed and $\left.e_{0}=10^{-6}\right)$

| $f_{0}$ | $10^{-6}$ | $10^{-5}$ | $10^{-3}$ | $10^{-3}$ | $10^{-2}$ | $10^{-1}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| NEE | $9.4 \times 10^{-4}$ | $1.3 \times 10^{-3}$ | $1.1 \times 10^{-3}$ | $1.7 \times 10^{-3}$ | $2.3 \times 10^{-3}$ | $3.9 \times 10^{-3}$ |

TABLE VII
The Impact of $g_{0}\left(h_{0}\right.$ IS Fixed and $\left.h_{0}=10^{-3}\right)$

| $g_{0}$ | $10^{-6}$ | $10^{-5}$ | $10^{-3}$ | $10^{-3}$ | $10^{-2}$ | $10^{-1}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| NEE | $1.4 \times 10^{-3}$ | $1.7 \times 10^{-3}$ | $1.4 \times 10^{-3}$ | $1.1 \times 10^{-3}$ | $1.4 \times 10^{-3}$ | $1.1 \times 10^{-3}$ |

TABLE VIII
the Impact of the Initial $K$

| Initial rank $K$ | 10 | 15 | 20 | 25 | 30 |
| :---: | :---: | :---: | :---: | :--- | :--- |
| NEE | $1.1 \times 10^{-3}$ | $1.1 \times 10^{-3}$ | $1.1 \times 10^{-3}$ | $1.4 \times 10^{-3}$ | $1.1 \times 10^{-3}$ |
| estimated rank | 8 | 9 | 11 | 12 | 13 |

TABLE IX
The Impact of Hankel Parameter $n_{2}$

| $n_{2}$ | 1 | 5 | 10 | 15 | 20 | 25 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| NEE | 0.075 | $6.5 \times 10^{-3}$ | $3.0 \times 10^{-3}$ | $1.4 \times 10^{-3}$ | $1.1 \times 10^{-3}$ | $1.2 \times 10^{-3}$ |

of B1. One can see that the recovery error and uncertainty index increase when the missing/bad data percentage increases. In Table IV, the recovery error is large when the missing data percentage is $45 \%$, and the corresponding uncertainty index is significantly larger than the values at other missing data percentages when the recovery errors are small. This verifies the effectiveness of our proposed uncertainty index.
3) The Impact of Parameter Selections: We evaluated the impact of parameter selections in recovering 5\% M2 missing and 5\% B2 bad data. The bad data are randomly selected from $(0.3,0.5)$. As discussed in Section III-D, we fixed $e_{0}$ and vary $f_{0}$ to show the impact of $\left(e_{0}, f_{0}\right)$. One can see from Table VI that Ba-NSDR maintains a very small recovery error with a wide range of $f_{0}$. We also fixed $h_{0}$ and varied $g_{0}$ to show the impact of $\left(g_{0}, h_{0}\right)$. Table VII shows that Ba-NSDR is not sensitive to the selection of $g_{0}$.

Table VIII shows the recovery performance when the initial rank varies. The recovery error NEE of Ba-NSDR remains very small with different ranks. Moreover, the estimated final ranks are consistent and much smaller than the initial rank, indicating that Ba-NSDR prunes the rank effectively.

The Hankel parameter $n_{2}$ is increased from 1 to 25 and the results are shown in Table IX. When $n_{2}=1$, the Hankel matrix reduces to the original data matrix. One can see from Table IX that increasing $n_{2}$ indeed leads to more accurate recovery results.

TABLE X
The Impact of Gaussian Kernel Parameter $c_{2}=c_{3}$

| $c_{2}=c_{3}$ | 40 | 50 | 60 | 70 | 80 |
| :---: | :---: | :---: | :---: | :---: | :---: |
| NEE | $1.3 \times 10^{-3}$ | $1.2 \times 10^{-3}$ | $1.1 \times 10^{-3}$ | $1.4 \times 10^{-3}$ | $1.4 \times 10^{-3}$ |

Table X shows the performance when the Gaussian kernel parameters $c_{2}$ and $c_{3}$ increase. The numerical results indicate that the proposed method is not sensitive to the Gaussian kernel parameters $c_{2}$ and $c_{3}$.

## V. Conclusion

This paper proposes a Bayesian high-rank Hankel matrix recovery ( $\mathrm{Ba}-\mathrm{NSDR}$ ) method to recover the synchrophasor measurements with missing and bad data. The proposed method maps the constructed Hankel matrix into a higher dimensional space by employing the kernel method and exploits the lifted low-rank Hankel property in recovering synchrophasor data under significant nonlinear dynamics. Ba-NSDR clearly outperforms the existing methods, especially when the data contain long consecutive missing or bad data. The distinctive features of Ba-NSDR include an uncertainty index that reflects the reliability of recovery results and the robustness to the initial rank selection. One future direction is to explore the effect of different kernels so that the method can pick the best kernel automatically for different scenarios.

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[^1]:    ${ }^{1}$ We provide an additional case study on the recorded PMU data of a transformer failure event in Central New York in the supplementary materials.

