

A Block Structure Preserving Model Reduction for Linear Circuit with Large Number of Ports

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Abstract—We propose a block structure-preserving model reduction (BSMOR) for the highly structured VLSI system. The blocks can be derived based on specific applications such as block current characterization of the substrate, or power/ground grids. Compared to PRIMA, BSMOR can match more poles using the same Krylov subspace and also increases the sparse ratio of the state matrices of the resulting macro-model. Experiment shows that BSMOR has a 20X smaller reduction time than PRIMA does under the same error bound. To efficiently analyze the resulting macro-model with large number of ports, we further propose a bordered-block diagonal (BBD) partitioning with a bottom-up hierarchical clustering (BBDC) where the macro-model is partitioned into a number of subset-port models, each with a manageable model size. With a similar accuracy, BBDC obtains 30X speedup compared to the original macro-model.

I. INTRODUCTION

VLSI circuits contain a number of highly structured components such as bus, power ground grid and substrate. These components can be modeled by passive networks with tremendous amount of circuit elements and large numbers of ports. To analyze such network efficiently, model order reduction [1]–[5] has been studied extensively. Based on the Krylov subspace projection and congruence transformation, PRIMA [5] is widely used to generate the reduced macro-model with preserved passivity. However, the macro-model produced by PRIMA is not compact as the order is usually “too high” to achieve the specified accuracy. Furthermore, the macro-model is represented by a multiple-input-multiple-output (MIMO) transfer function, and is usually dense and inefficient to analyze for a large number of ports.

To improve upon PRIMA, a structure-preserving model reduction (SPRIM) is proposed in [6]. It partitions the state matrix in the MNA (modified nodal analysis) form into a natural 2×2 block matrices, i.e., conductance, capacitance, inductance, and adjacent (G, C, L, E_s) matrices. Accordingly the projection matrix is partitioned and the number of its columns is doubled. As a result, SPRIM matches the twice poles of the models by using the projection matrix given by PRIMA. In addition, the block structure of state matrices is preserved, which facilitates the realization of the reduced model. However, such a simple 2×2 partition does not leverage the regularity of the substrate network. In addition, the explicit hierarchical decomposition [7], [8] is proposed to handle a large number of ports. The capacity of these methods [6]–[8] need to be improved further.

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In this paper, we propose a block structure preserving model reduction (BSMOR) method, which generalizes SPRIM [6] in the sense that the G, C, L and E_s matrices are further partitioned into blocks. The blocks can be derived based on specific applications such as block current characterization of the substrate in this paper. We find that (i) for the system state matrices having *block diagonal structure*, BSMOR with m blocks can exactly match $(m - 1)q$ more poles than PRIMA does within q iterations. Therefore, increasing the block number leads to more matched poles using the same Krylov space; (ii) for the general structured state matrices, using Singular-value-decomposition (SVD) and K-means-clustering [9] based partitioning, the additional $(m - 1)q$ poles obtained by BSMOR can closely approximate the poles of original system. Therefore, BSMOR can lead to more efficient reduction under the same accuracy than PRIMA. In addition, BSMOR can also preserve the sparsity for reduced block matrices, which gives further efficiency boost to constructing a macro-model. The resulting macro-model consists of order-reduced blocks, each containing a subset of ports. To analyze a macro-model with a large number of ports, we further propose a bordered-block diagonal (BBD) partitioning and hierarchical clustering of reduced blocks. We call it BBDC analysis. The experiment shows that under the same accuracy, the reduction of our approach is 20X times faster than PRIMA to reduce a circuit with 1M elements, and the BBDC analysis is 30X faster compared to analyzing the original macro-model.

The rest of the paper is organized as follows. We review Grimme’s projection theorem [10] and SPRIM algorithm in Section II, present BSMOR method and its properties in Section III, and discuss the SVD and K-means-clustering based partitioning algorithm in Sections IV. We further present a hierarchical bordered-block-diagonal analysis for the reduced model in Section V. In Section VI, we apply our method to the substrate macro-modeling and noise analysis, and discuss how to find the block structure from the characterization of the block current. We present the experimental results in Section VII, and conclude the paper in Section VIII.

II. PRELIMINARY

Consider a modified nodal formulation (MNA) of the circuit equation in the frequency domain:

$$\begin{aligned} \mathcal{G}x(s) + s\mathcal{C}x(s) &= \mathcal{B}u_e(s) \\ y_e(s) &= \mathcal{B}^T x(s) \end{aligned} \quad (1)$$

where $x(s)$ is the state variable vector, \mathcal{G} and \mathcal{C} ($\in R^{N \times N}$) are state matrices. \mathcal{B} ($\in R^{N \times n_p}$) is

$$\mathcal{B} = [B \ 0]^T, \quad (2)$$

a port incident matrix. Eliminating $x(s)$ in (1) gives

$$\begin{aligned} y_e(s) &= H(s)u_e(s) \\ H(s) &= \mathcal{B}^T(\mathcal{G} + s\mathcal{C})^{-1}\mathcal{B}, \end{aligned} \quad (3)$$

where $H(s)$ is a multiple-input multiple-output (MIMO) transfer function. PRIMA finds a projection matrix $V \in R^{N \times q}$ such that its columns span the q -th block Krylov subspace $\mathcal{K}(\mathcal{A}, \mathcal{R}, q)$, i.e.,

$$\mathcal{K}(\mathcal{A}, \mathcal{R}, q) = \text{span}(V) = \{\mathcal{R}, \mathcal{A}\mathcal{R}, \dots, \mathcal{A}^{n-1}\mathcal{R}\}, \quad (4)$$

where $n = \lceil q/n_p \rceil$, $\mathcal{A} = (\mathcal{G} + s_0\mathcal{C})^{-1}\mathcal{C}$, $\mathcal{R} = (\mathcal{G} + s_0\mathcal{C})^{-1}\mathcal{B}$, and s_0 is the expansion point that ensures $\mathcal{G} + s_0\mathcal{C}$ is nonsingular. The resulting reduced transfer function is

$$\hat{H}(s) = \hat{\mathcal{B}}^T(\hat{\mathcal{G}} + s\hat{\mathcal{C}})^{-1}\hat{\mathcal{B}}, \quad (5)$$

where

$$\hat{\mathcal{G}} = V^T\mathcal{G}V, \quad \hat{\mathcal{C}} = V^T\mathcal{C}V, \quad \hat{\mathcal{B}} = V^T\mathcal{B}, \quad (6)$$

Theorem 1: If $\mathcal{K}(\mathcal{A}, \mathcal{R}, q) \subseteq \text{span}(V)$, then $\hat{H}(s)$ has the identical expanded first q -th moments with $H(s)$.

It is called the *Grimme's projection theorem* [10]. Note that $\hat{\mathcal{G}}, \hat{\mathcal{C}} \in R^{q \times q}$, and $\hat{\mathcal{B}} \in R^{q \times n_p}$.

In [6], a structure-preserving reduced model order reduction technique, SPRIM, is proposed. The primary observation is that instead of using the Krylov subspace $\mathcal{K}(\mathcal{A}, \mathcal{R}, q)$ for the projection matrix \tilde{V} , one can use any projection matrix such that the space spanned by the column in \tilde{V} contains the q -th block Krylov subspace. i.e.

$$\mathcal{K}(\mathcal{A}, \mathcal{R}, q) = \text{span}(V) \subseteq \text{span}(\tilde{V}) \quad (7)$$

In SPRIM, a 2×2 partition is naturally used as a linear state matrix in the MNA form shows a 2×2 block structure

$$\mathcal{G} = \begin{bmatrix} G & E_s^T \\ -E_s & 0 \end{bmatrix}, \mathcal{C} = \begin{bmatrix} C & 0 \\ 0 & L \end{bmatrix}, \quad (8)$$

where $G \in R^{n_1 \times n_1}$, $C \in R^{n_1 \times n_1}$, $L \in R^{n_2 \times n_2}$ are conductance, capacitance and inductance matrix, and $E_s \in R^{n_2 \times n_1}$ is the adjacent matrix indicating the branch current flow at the inductor. Note that $n_1 + n_2 = N$.

Therefore, a structured projection vector \tilde{V} can be constructed by partitioning the projection vector V obtained from the q -th PRIMA iteration

$$V = \begin{bmatrix} V_1 \\ V_2 \end{bmatrix} \rightarrow \tilde{V} = \begin{bmatrix} V_1 & 0 \\ 0 & V_2 \end{bmatrix}. \quad (9)$$

where $V_1 \in R^{n_1 \times q}$, $V_2 \in R^{n_2 \times q}$, and hence $\tilde{V} \in R^{N \times 2q}$. As a result, the number of columns in \tilde{V} is twice of that in V . Accordingly the new reduced state matrices are

$$\tilde{\mathcal{G}} = \begin{bmatrix} \tilde{G} & \tilde{E}_s^T \\ -\tilde{E}_s & 0 \end{bmatrix}, \tilde{\mathcal{C}} = \begin{bmatrix} \tilde{C} & 0 \\ 0 & \tilde{L} \end{bmatrix}, \quad (10)$$

where $\tilde{G} = V_1^T G V_1$, $\tilde{E}_s = V_2^T E_s V_1$ and $\tilde{C} = V_1^T C V_1$ and $\tilde{L} = V_2^T L V_2$. Similarly, the size of $\tilde{\mathcal{G}}, \tilde{\mathcal{C}} \in R^{2q \times 2q}$, and $\tilde{\mathcal{B}} \in R^{2q \times n_p}$ is twice than that of $\hat{\mathcal{G}}, \hat{\mathcal{C}}$, and $\hat{\mathcal{B}}$ reduced by using V . Therefore, the moments of the reduced model with state matrices: $\tilde{\mathcal{G}}$ and $\tilde{\mathcal{C}}$ are twice than those of the reduced model with state matrices: $\hat{\mathcal{G}}$ and $\hat{\mathcal{C}}$. In other words, the reduced model by \tilde{V} matches $2q$ moments of the original model instead of q moments as the reduced model by V .

Since the reduced model is written in the first order form in (10), the model reduced by SPRIM is twice larger than that produced by PRIMA. But the reduced model produced by SPRIM preserves the structure of the original model and can be further reduced into the second-order form using node elimination base on the Schur's decomposition [11]: $\tilde{\mathcal{H}}_{NA} = \tilde{G} + s\tilde{C} + \frac{1}{s}\tilde{E}_s^T \tilde{L}^{-1} \tilde{E}_s$ where $\tilde{\mathcal{H}}_{NA}$ is the reduced state matrix in NA form, which has the same size of the reduced matrix by using V . But the difference is that each element in $\tilde{\mathcal{H}}_{NA}$ becomes second-order rational function of s instead of first-order polynomial of s .

Hence SPRIM algorithm essentially consists of two reduction steps: the first step is the structure-preserving projection-based reduction and the second step is block node elimination based on Schur's decomposition. As a result, SPRIM can match more poles than PRIMA, which uses V as the projection matrix, but both result in a same *size* of the reduced model. If we just look at the first step, SPRIM simply matches more moments by using more columns in the projection matrix \tilde{V} , thus produces larger reduced state matrices in the first-order form.

III. BSMOR METHOD

SPRIM essentially is based on a 2×2 partitioning of the state matrices. If we use more partitions (*each partition called a block*), we can add more columns into the project matrix \tilde{V} , thus match more poles given the same Krylov space $\mathcal{K}(\mathcal{A}, \mathcal{R}, q)$.

A. Structured Block Projection

Specifically, we assume that the conductance matrix \mathcal{G} can be distinguished in m blocks

$$\mathcal{G} = \begin{bmatrix} \mathcal{G}_{1,1(n_1 \times n_1)} & \mathcal{G}_{1,2(n_1 \times n_2)} & \dots & \mathcal{G}_{1,m(n_1 \times n_m)} \\ \mathcal{G}_{2,1(n_2 \times n_1)} & \mathcal{G}_{2,2(n_2 \times n_2)} & \dots & \mathcal{G}_{2,m(n_2 \times n_m)} \\ \vdots & \vdots & \ddots & \vdots \\ \mathcal{G}_{m,1(n_m \times n_1)} & \mathcal{G}_{m,2(n_m \times n_2)} & \dots & \mathcal{G}_{m,m(n_m \times n_m)} \end{bmatrix}, \quad (11)$$

where each block has the size n_k ($\sum_{k=1}^m n_k = N$). A similar block structure can be found for \mathcal{C} matrix. Then, \mathcal{B} becomes

$$\mathcal{B} = [\mathcal{B}_{1(n_1 \times n_p)}, \mathcal{B}_{2(n_2 \times n_p)}, \dots, \mathcal{B}_{m(n_m \times n_p)}]^T \quad (12)$$

where each block contains user specified n_{p_k} ports ($n_p = \sum_{k=1}^m n_{p_k}$). Note that these blocks can be derived based on specific applications such as block current characterization of the substrate or the power/ground grid, called *natural basic blocks*, or determined after specific partitioning algorithms discussed later on.

Accordingly, we further partition the projection matrix V obtained from PRIMA according to the block structure in state matrices from (11)

$$\begin{aligned}
V &= \begin{bmatrix} V_1^{(n_1 \times q)} \\ V_2^{(n_2 \times q)} \\ \vdots \\ V_m^{(n_m \times q)} \end{bmatrix} \\
\rightarrow \tilde{V} &= \begin{bmatrix} V_1^{(n_1 \times q)} & 0 & \cdots & 0 \\ 0 & V_2^{(n_2 \times q)} & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & V_m^{(n_m \times q)} \end{bmatrix}. \quad (13)
\end{aligned}$$

where $\tilde{V} \in R^{N \times mq}$. We call this as *Block Structure-preserving Model Reduction* (BSMOR), where m is the number of blocks.

We can obtain the order reduced state matrices by projecting \tilde{V} :

$$\tilde{\mathcal{G}} = \tilde{V}^T \mathcal{G} \tilde{V}, \quad \tilde{\mathcal{C}} = \tilde{V}^T \mathcal{C} \tilde{V}, \quad \tilde{\mathcal{B}} = \tilde{V}^T \mathcal{B}. \quad (14)$$

Elementwise, we have

$$\tilde{\mathcal{G}}_{i,j} = V_i^T \mathcal{G}_{i,j} V_j, \quad \tilde{\mathcal{C}}_{i,j} = V_i^T \mathcal{C}_{i,j} V_j, \quad \tilde{\mathcal{B}}_i = V_i^T \mathcal{B}_i \quad (15)$$

where $\tilde{\mathcal{G}}_{i,j}$ represents the blocks at i block row and j block column in reduced matrix $\tilde{\mathcal{G}}$. So do $\tilde{\mathcal{C}}_{i,j}$ and $\tilde{\mathcal{B}}_i$. Let $V_i = V_i^{(q \times n_i)}$ to simplify notations. Note that such a $m \times m$ block projection preserves the structure and sparsity of the original \mathcal{G} , \mathcal{C} matrices. For example, when projected by \tilde{V} , the reduced $\tilde{\mathcal{G}}$ matrix is

$$\tilde{\mathcal{G}} = \begin{bmatrix} V_1^T \mathcal{G}_{1,1} V_1 & V_1^T \mathcal{G}_{1,2} V_2 & \cdots & V_1^T \mathcal{G}_{1,m} V_m \\ V_2^T \mathcal{G}_{2,1} V_1 & V_2^T \mathcal{G}_{2,2} V_2 & \cdots & V_2^T \mathcal{G}_{2,m} V_m \\ \vdots & \vdots & \ddots & \vdots \\ V_m^T \mathcal{G}_{m,1} V_1 & V_m^T \mathcal{G}_{m,2} V_2 & \cdots & V_m^T \mathcal{G}_{m,m} V_m \end{bmatrix}. \quad (16)$$

Clearly, if the original \mathcal{G} is sparse, the resulted $\tilde{\mathcal{G}}$ is sparse as well. In the contrast, when projected by V using PRIMA, the resulted $\hat{\mathcal{G}}$ is

$$\hat{\mathcal{G}} = \sum_{i=1}^m \sum_{j=1}^m V_i^T \mathcal{G}_{i,j} V_j, \quad (17)$$

which is dense in general. Similar observations can be found for \mathcal{C} and \mathcal{B} .

B. Passivity Preservation and Moment Matching

Similar to SPRIM, the reduced model of passive network obtained by Krylov-subspace projection preserves passivity:

Theorem 2: The reduced order model $\tilde{H}(s)$ by BSMOR is passive.

Proof: Because $\tilde{V}^T \tilde{V} = I$, and $\mathcal{G} + \mathcal{G}^T$, $\mathcal{C} + \mathcal{C}^T$ are symmetric positive definite, the congruence transformation based projections: $\tilde{V}^T \mathcal{G} \tilde{V}$, $\tilde{V}^T \mathcal{C} \tilde{V}$ preserve the passivity [5].

Using such a projection matrix \tilde{V} , we define a reduced-order model with the following transfer function

$$\tilde{H}(s) = \tilde{\mathcal{B}}^T (\tilde{\mathcal{G}} + s\tilde{\mathcal{C}})^{-1} \tilde{\mathcal{B}}, \quad (18)$$

and we have

Theorem 3: Let V be a matrix that satisfies $\mathcal{K}(\mathcal{A}, \mathcal{R}, q) \subseteq \text{span}(V)$ and \tilde{V} is defined in Eq.(13). $\tilde{H}(s)$ will match the first q moments in the expansion of $H(s)$ about s_0 . If the \mathcal{G} , \mathcal{C} matrices in the block diagonal form have m different blocks,

i.e. $\mathcal{A}_{1,1} \neq \mathcal{A}_{2,2} \neq \dots \neq \mathcal{A}_{m,m}$, $\tilde{H}(s)$ will match mq poles of $H(s)$.

Proof: Because $\mathcal{K}(\mathcal{A}, \mathcal{R}, q) \subseteq \text{span}(V) \subseteq \text{span}(\tilde{V})$, according to *Grimme's projection theorem* (Theorem 1), $\tilde{H}(s)$ will match the first q moments in the expansion of $H(s)$ about s_0 . Moreover, if \mathcal{G} , \mathcal{C} matrices are in the block diagonal form, then the state matrix $\mathcal{A} = \mathcal{G}^{-1}\mathcal{C}$ is in the block-diagonal form as well. Due to the structure-preserving projection, the resulted state matrix $\tilde{\mathcal{A}}$

$$\begin{aligned}
\tilde{\mathcal{A}} &= (\tilde{V}^T \mathcal{G} \tilde{V})^{-1} (\tilde{V}^T \mathcal{C} \tilde{V}) = \text{diag}[\mathcal{A}_{1,1}, \mathcal{A}_{2,2}, \dots, \mathcal{A}_{m,m}] \\
&= \begin{bmatrix} (\tilde{\mathcal{G}}_{1,1})^{-1} \tilde{\mathcal{C}}_{1,1} & 0 & \cdots & 0 \\ 0 & (\tilde{\mathcal{G}}_{2,2})^{-1} \tilde{\mathcal{C}}_{2,2} & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & (\tilde{\mathcal{G}}_{m,m})^{-1} \tilde{\mathcal{C}}_{m,m} \end{bmatrix}
\end{aligned}$$

have m sub-blocks each with size $q \times q$. Note that eigenvalues of $\tilde{\mathcal{A}}$ represent the reciprocal poles of the reduced model [5]. As a result, the partitioned projection matrix \tilde{V} leads to localized projection. In other words, the block projection matrix \tilde{V}_i is used only for the state matrix block $\mathcal{A}_{i,i}$ ($i = 1, \dots, m$). Each structured block projection matrix \tilde{V}_i will lead to the localized model order reduction for block i , which is represented by $(\mathcal{A}_{i,i}, \mathcal{R}_{i,i})$. Conceivably, the order reduced block $(\tilde{\mathcal{A}}_{i,i}, \tilde{\mathcal{R}}_{i,i})$ will match $(\mathcal{A}_{i,i}, \mathcal{R}_{i,i})$ with q poles. But the whole system consisting of the m blocks will match mq poles.

In general when $\mathcal{G}_{i,j}, \mathcal{C}_{i,j} \neq 0$, i.e., $\mathcal{A}_{i,j} \neq 0$, BSMOR obtains $(m-1)q$ times more poles than PRIMA within q iterations, where the $(m-1)q$ additional poles approximately match poles of the original system. If the coupling $\mathcal{A}_{i,j}$ between blocks $\mathcal{A}_{i,i}$ is weak, the original system can be well approximated by the block diagonal structure, and therefore, each block has a different pole distribution, and the $(m-1)q$ additional poles are close to the poles of the original system. As a result, introducing more partitions or blocks can archive the same reduction accuracy by using less iterations, which can in turn improve the reduction efficiency. On the other hand, if the coupling $\mathcal{A}_{i,j}$ between blocks $\mathcal{A}_{i,i}$ is strong, the accuracy of BSMOR depends on the partition algorithm. Note that in this case, blocks are strongly correlated. Below, we present a partitioning algorithm based on the the singular-value-decomposition and K-means clustering.

IV. SVD AND K-MEANS-CLUSTERING BASED PARTITIONING

When blocks are strongly correlated, the state matrix (\mathcal{A}) will be a low rank matrix. The true rank should reveal the number of uncorrelated rows in \mathcal{A} . Therefore, if we can find this true rank number m , and greedy group all similar row vectors of \mathcal{A} into m most uncorrelated blocks, then we can obtain a partitioning that each partitioned block has weak correlation with each other, and shows a different pole distribution. Using such a partitioned V to construct the structure projection matrix \mathcal{V} accordingly, and project \mathcal{G} , \mathcal{C} matrices, the obtained mq poles can closely match poles of the original system.

Based on this observation, SVD is applied on \mathcal{A} to determine the rank value m , which is used as the partition

number. With the known partition number, K-means-clustering method [9] can be applied to greedily add similar rows into one block, and find m disjoint blocks. As a result, the moment value, i.e., the pole distribution is similar in the same block, and is different between them. However, the dimension of \mathcal{A} ($\in R^{N \times N}$) is too high to directly apply SVD and K-means clustering. Observing that the projection matrix V composed by $(\mathcal{A}, \mathcal{R})$ as in (4) has a low-dimension $R^{q \times N}$, we apply SVD and K-means-clustering to V . The overall procedure is presented in Algorithm 1.

K-means-clustering classifies a given data set. The main idea is to define k centroids, one for each cluster. These centroids should be placed in a cunning way because of different location causes different result. As a result, the better choice is to place them as far away from each other as possible. The next step is to iteratively associate data points to the nearest centroid. When no point is left, the first step is completed and an initial clustering is obtained. At this point we need to recalculate k new centroids as bary-centers of the clusters resulting from the previous step. After we have these k new centroids, a new iteration binding has to be done. As a result the k centroids may change their locations and iterations stop when there is no more changes for the k centroids. This procedure minimizes the following squared error function

$$J = \sum_{j=1}^k \sum_{i=1}^n \|x_i^{(j)} - c_j\|^2, \quad (19)$$

where $\|x_i^{(j)} - c_j\|^2$ is the distance between data point $x_i^{(j)}$ and cluster centroid c_j .

Algorithm 1 BSMOR Partitioning 1

1. Find projection matrix by PRIMA

(1.1) *input*: $\mathcal{G}, \mathcal{C}, \mathcal{B}$;

(1.2) *form*: $\mathcal{A} = (\mathcal{G} + s_0\mathcal{C})^{-1}\mathcal{C}$, $\mathcal{R} = (\mathcal{G} + s_0\mathcal{C})^{-1}\mathcal{B}$;

(1.3) *construct*: $V = [V_1, V_2, \dots, V_q]$ such that $\text{span}(V) = \mathcal{K}(\mathcal{A}, \mathcal{R}, q)$, where $V \in R^{N \times mq}$;

2. SVD and K-means-clustering based partitioning

(2.1) *solve*: $m = \text{SVD}(V)$;

(2.2) *cluster*: $[V_{1(n_1 \times q)}, \dots, V_{m(n_m \times q)}]^T = \text{kmeans}(V, m)$;

(2.3) *construct*: $\mathcal{V} = \text{diag}[V_{1(n_1 \times q)}, \dots, V_{m(n_m \times q)}]$;

3. Structured block projection

(3.1) *project*: $\tilde{\mathcal{G}} = \tilde{V}^T \mathcal{G} \tilde{V}$, $\tilde{\mathcal{C}} = \tilde{V}^T \mathcal{C} \tilde{V}$, $\tilde{\mathcal{B}} = \tilde{V}^T \mathcal{B}$;

(3.2) *output*: $\tilde{\mathcal{G}}, \tilde{\mathcal{C}}, \tilde{\mathcal{B}}$.

In summary, using the partitioned V by the SVD and K-means-clustering, one can construct a structured-projection matrix \mathcal{V} , where each block has a different pole distribution, and the reduced model can closely match mq poles of the original system. As a result, one can obtain order reduced models with higher accuracy for each structured block by using the same sized Krylov subspace base vectors, or get the same order reduced model (same accuracy) using a smaller Krylov subspace. Therefore, BSMOR provides much more flexibility and a better trade-off between efficiency and model accuracy for reducing linear dynamic system models than PRIMA does.

V. BORDERED-BLOCK DIAGONAL PARTITIONING WITH HIERARCHICAL CLUSTERING

In this section, we first describe the flat macro-model generated by the reduced state matrices from Section 2.2. To efficiently handle the flat macro-model with large number of ports, we present a *bordered-block diagonal* (BBD) partitioning to solve each block individually. Moreover, we discuss a hierarchical clustering method to further improve the efficiency.

A. Flat Macro-model

For the frequency-dependent application in the analog/RF simulation like the substrate noise analysis, an Y -parameter based multiple port macro-model is widely used instead. An $n_p \times n_p$ MIMO admittance matrix $Y'(s)$ can be obtained by taking the eigen-decomposition of $\tilde{A} = (\tilde{\mathcal{G}} + s_0\tilde{\mathcal{C}})^{-1}\tilde{\mathcal{C}}$

$$Y'(s) = \begin{bmatrix} Y'_{1,1} & \cdots & Y'_{1,n_p} \\ \vdots & \ddots & \vdots \\ Y'_{n_p,1} & \cdots & Y'_{n_p,n_p} \end{bmatrix}, \quad (20)$$

with

$$Y'_{i,j} = c^{i,j} + \sum_{n=1}^q \frac{k_n^{i,j}}{s - p_n}, \quad (21)$$

where k_n and p_n are the residues and poles. Note that eigenvalues of $\tilde{A}^{(q)}$ represent the reciprocal poles of $Y'(s)$ [5]. Due to the preserved sparsity, the eigen-decomposition becomes more efficient when using the $\tilde{\mathcal{G}}$ and $\tilde{\mathcal{C}}$ from the BSMOR other than using those from PRIMA. Furthermore, as the reduction preserves the structure, it results in additional preservations: (i) the reciprocity of the network is also preserved, i.e., the $Y'(s)$ is symmetrical. In contrast, PRIMA does not preserve this property; (ii) the block structure is preserved as well. It means the reduced block can be distinguished by a subset of ports specified before BSMOR. As a result, we can further apply an additional port-partitioning, precisely, bottom-up port clustering to handle the large number of ports as discussed later on.

To partition a given network, we first transform the nodal admittance (20) into a branch admittance network:

$$Y_{ii} = \sum_{j=1}^{n_p} Y'_{ij}, \quad Y_{ij} = -Y'_{ij}. \quad (22)$$

Note that the flat macro-model consists of m order reduced blocks, where each *reduced block* contains n_{pk} ports with ground and coupling branch admittances. There also exist coupling branch admittances between any pair of reduced blocks. A transformed branch admittance network for a 4-port admittance matrix is shown in Fig. 1 (a). To partition the branch admittance network Y , one natural approach is to reserve each reduced block, and pack all the coupling branch admittances into one block, called as *coupling block* that connects all reduced blocks. An example of such a partitioning (or representation of the macro-model from BSMOR) is shown in Fig. 1 (b) for a 4-port admittance matrix.

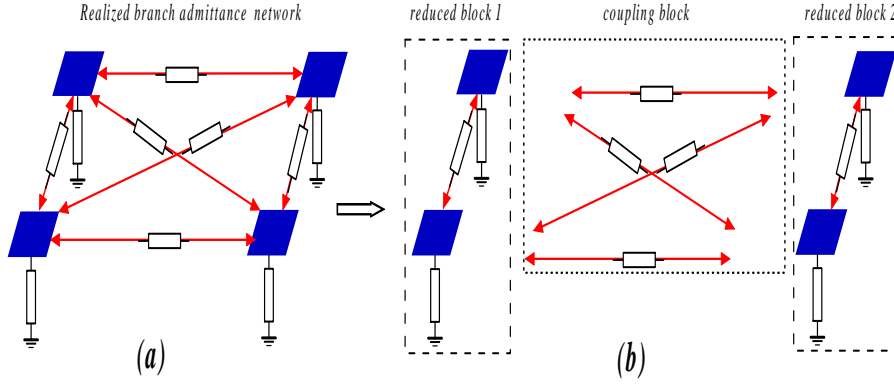


Fig. 1. An example of 4-port admittance with 2 reduced blocks. (a) realization in branch admittance network; (b) represented by 2 reduced blocks with an additional coupling block.

B. Bordered-Block Diagonal Matrix

For the k th reduced block, we have

$$\mathbf{Y}_k v_k = i_k + \tilde{i}_k, \quad (23)$$

where

$$(\mathbf{Y}_k)_{ii} = \sum_{j=1}^{n_{pk}} Y'_{ij}, \quad (\mathbf{Y}_k)_{ij} = -Y'_{ij} \quad (j \in n_{pk}), \quad (24)$$

and v_k , i_k are the port voltage and current vectors, where i_k is part of i_p : $i_k = i_p(\dots \underbrace{i_{k1} \dots}_{n_{pk}} \dots)$. Moreover, \tilde{i}_k is the

correlation current from the other reduced block through the coupling block.

The branch equation for the coupling block is

$$(\mathbf{Y}_0)^{-1} i_0 = v_0, \quad (25)$$

where \mathbf{Y}_0 is the branch admittance matrix describing the branches in the coupling block. It is a diagonal matrix such that its inversion is easily obtained as $1/(\mathbf{Y}_0)_{ii}$. Note that its size depends on the number of couplings among reduced blocks, and it can be efficiently implemented with the sparse matrix data structure. v_0 and i_0 are branch voltage and current vectors. They relate to the port voltage/current vectors v_k/i_k at k th block by

$$\tilde{i}_k = C_{k0} i_0, \quad v_0 = -\sum_{k=1}^m (C_{k0})^T v_k, \quad (26)$$

where C_{k0} is the cut matrix composed by $\{0, 1, -1\}$ to indicate the direction of branch currents between k th reduced block and the coupling block.

Combine (24) - (26), we have the following hybrid matrix equation

$$\begin{bmatrix} \mathbf{Y}_1 & 0 & \dots & 0 & C_{10} \\ 0 & \mathbf{Y}_2 & \dots & 0 & C_{20} \\ \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & \dots & \mathbf{Y}_m^{(l)} & C_{m0} \\ (C_{10})^T & (C_{20})^T & \dots & (C_{m0})^T & -(\mathbf{Y}_0)^{-1} \end{bmatrix} \begin{bmatrix} v_1 \\ v_2 \\ \vdots \\ v_m \\ i_0 \end{bmatrix} = \begin{bmatrix} i_1 \\ i_2 \\ \vdots \\ i_m \\ 0 \end{bmatrix}.$$

This hybrid matrix shows a *bordered-block-diagonal* (BBD) structure. It enables the below Algorithm 2 [12] to solve each reduced block individually without using the explicit inversion. Each reduced block matrix is first solved individually with LU factorization and substitution (1.1-1.5), the results from each reduced block are then used further to solve the coupling block (2.1-2.4), and the final v_k of each reduced block is updated (3.1-3.4) with the result from the coupling current i_0 .

Algorithm 2 Solve bordered-block-diagonal (BBD) matrix

1. Solve Y_k individually

for every k in m do

(1.1) *input*: Y_k, C_{k0}, i_k ;

(1.2) *factor*: $Y_k = L_k U_k$;

(1.3) *solve*: $L_k \Phi_k = C_{k0}$ for Φ_k , $\Psi_k U_k = (C_{k0})^T$ for Ψ_k , and $L_k \xi_k = i_k$ for ξ_k ;

(1.4) *form*: $F_k = \Phi_k^T \Psi_k$, and $G_k = \Psi_k^T \xi_k$

end for

2. Solve Y_0

(2.1) *form*: $F = Y_0^{-1} + \sum_{k=1}^m F_k$, $G = \sum_{k=1}^m G_k$;

(2.2) *solve*: $F i_0 = G$ for i_0 ;

3. Update Y_k individually

for every k in m do

(3.2) *form*: $\xi_k = \xi_k - \Phi_k i_0$;

(3.3) *solve*: $U_k v_k = \xi_k$ for v_k ;

(3.4) *output*: v_k .

end for

Typically, LU factorization requires $n^3/3$ multiplications and back/forward substitution requires $n^2/2$ multiplications. The computational cost of Algorithm 1 is therefore, $\sum_{k=1}^m (n_{pk}^3/3 + n_{pk}^2/2) + (n_0^3/3 + n_0^2/2)$, where n_{pk} is the port number (reduced block size) of each reduced block, and n_0 is the size of the coupling block. Note that if the parallel execution is used, the *summation* becomes the *maximum* execution time among blocks. To reduce the computational cost without using the parallel execution, we need control costs for reduced blocks and the coupling block as discussed below.

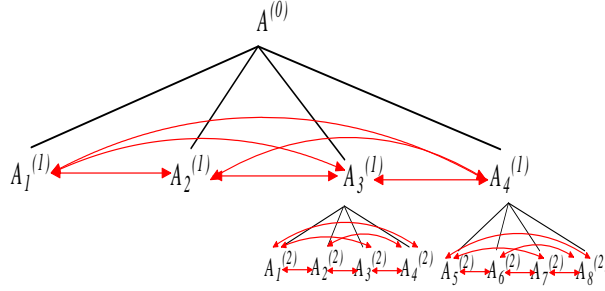


Fig. 2. The hierarchical tree structure of clustered blocks.

C. Hierarchical Clustering

As the factorization cost decreases with the size of the reduced block, apparently the computation cost will be small when the network is partitioned based on the reduced basic block from BSMOR. However, the size of \mathbf{Y}_0 increases with the reduced block number, and it increases the computation cost. To wisely arrange this trade-off, a hierarchical tree structure is used as shown in Fig. 2. In this tree, each node represents an *abstract block*. There are links connecting each pair of correlated blocks, representing inter-block couplings. The tree is constructed by iteratively *clustering* the reduced blocks from the bottom. The degree and the level is chosen to bound the size of the coupling block below a threshold. At the leaf level, a cluster of reduced blocks are siblings of a parent node, an abstract block. A *cluster-coupling block* is introduced to model the coupling between siblings. There is no direct coupling between abstract blocks not in a same cluster, but their coupling is modeled by cluster-coupling blocks for parent nodes. Therefore, we can maintain a constant link number (couplings) at each tree level. Note that the following *merge operation* is operated when two blocks k and l are clustered

$$i_{new} = [i_k, i_l], \quad v_{new} = [v_k, v_l],$$

and

$$(\mathbf{Y}_{new})_{ii} = \sum_{j \in n_{p_k} \cup n_{p_l}} Y'_{ij}, \quad (\mathbf{Y}_{new})_{ij} = -Y'_{ij}.$$

At the bottom level, we solve each clustered block using Algorithm 1. It is inefficient to calculate v_k directly on the higher levels since the block size get larger and larger. Fortunately this is not necessary, because one can use the already calculated v_k of the children, same as to attach the voltage sources to the coupling block at parent node. To do this we need to update i_0 from $(l-1)$ th level to l th level by

$$v_0^{(l-1)} = - \sum_{k=1}^{m^{(l-1)}} (G_{k0}^{(l)})^T v_k^{(l-1)}, \quad i_0^{(l-1)} = Y_0^{(l)} v_0^{(l-1)}, \quad (27)$$

and then solve v_k at l th level by (3.1)-(3.4) in Algorithm 1. Moreover, with the hierarchical tree structure, v_k is recursively updated by a bottom-up depth-first traversal of the tree, where we assume that the cut matrices and block branch admittance are pre-computed and stored hierarchically. For simplicity of presentation, we call BBD analysis with hierarchical clustering

as BBDC analysis. Note that when the factorization cost of large matrix at the top level is large, we further apply an error-bounded sparsification technique similar as [7] to the branch admittance matrix. As the sparsification is performed at the top level, this error is bounded.

VI. BLOCK SPECIFICATION IN SUBSTRATE NOISE ANALYSIS

In this section, we discuss the application of BSMOR and BBDC analysis to the substrate macro-modeling and noise analysis. The substrate outside of active/contact areas can be treated as a uniformly doped layer, where an electrostatic Maxwell's equation is:

$$\epsilon \frac{\partial}{\partial t} (\nabla \cdot E) + \frac{1}{\rho} (\nabla \cdot E) = 0. \quad (28)$$

The Eddy current term (the primary cause of substrate loss) can be ignored if the substrate is highly doped, where the conduction current is dominant. Note that (28) can be discretized in differential form using finite-difference [13] or integral form using boundary element (BEM) methods. Because the BEM method needs to find a numerically stable multi-layer Green's function [14], it is not trivial to construct in general when the layout geometry becomes arbitrary. In this paper, the finite-difference based discretization is used to generate the RC mesh/grid as the substrate circuit model. As the electric field varies nonlinearly as a function of the distance, the finite-difference method approximates this variation as a piecewise constant function by carefully choosing the pitch of the mesh according to the current density, i.e., the strength of the electrical field.

For leading-edge integrated circuits, the count of gates is typically in millions. The number of possible locations to place contacts of sensitive analog/RF circuits is large as well. Therefore, a flat multi-port description of each individual substrate noise injector and receptors is impractical. We assume that the chip is partitioned into smaller circuit, i.e., blocks based on the switching current density. As a result, within a block all noise current injections can be clustered into one independent current source at one single injection port. Such a block maximum current spectrum envelope is studied in [15] to characterize the injection noise sources in a bottom-up fashion. The noise current injected by the gate G at frequency f_p is denoted $i_G(f_m)$, and $f_m = m \times f_0$ ($m = 0, 1, 2, \dots, M$), where f_0 is the clock frequency and M is the sampling bound.

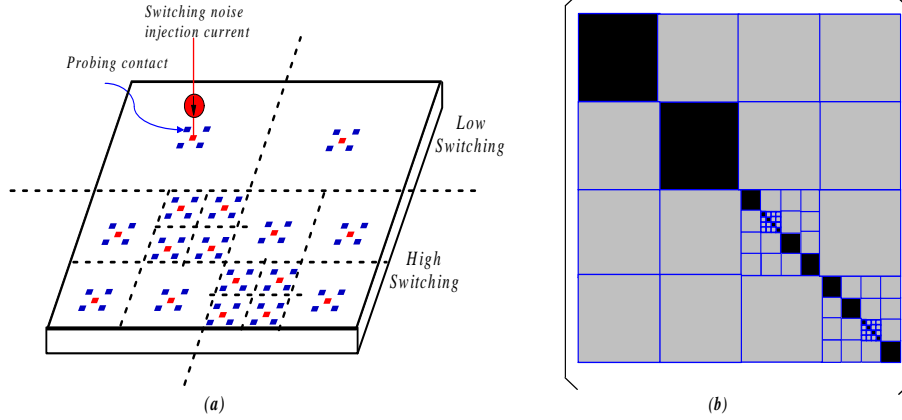


Fig. 3. (a) The non-uniform substrate mesh network characterized by the switching current density; (b) The corresponding block structure of conductance/capacitance matrices.

Then, the total noise current of c_N gates in k th block is $i_{Ck} = \sum_{k=1}^{C_N} i_{Gk}(f_m)$, and by a library-based characterization of the primary input transition v_p , the block current envelope spectrum is found by $i_k^{max}(f_m) = \max_{v_p} |i_{Ck}(f_m)|$.

Therefore, if there are m characterized blocks, each block would contain n_{pk} user specified ports, including one input port representing the injecting current noise source according to the above block current assumption, and $(n_{pk} - 1)$ output ports representing all possible contact locations for analog/RF modules. There are total n_p ($n_p = \sum_{k=1}^m n_{pk}$) specified ports. The port current vector i_p becomes: $i_p = [i_1^{max} \dots i_k^{max} \dots i_m^{max} \dots 0]$, where all omitted entries are zeros standing for probing output ports. Note that the propagated noise is observed from v_p .

However, with the use of the power management technique like the clock gating, the $i_{Ck}(f_m)$ can be very non-uniform for each block across the chip. For the block with the high current density, the electric field tends to vary largely, and a finer grids are necessary for the accurate approximation. Otherwise, coarse grid is used instead. For example, the substrate plane in Fig. 3 (a) has 4 parts with different switching current densities and it results in a non-uniform mesh structure. As a result, it demonstrates a block structure according to the block current density. For example, Fig. 3 (b) shows such a block structure for the block current distribution in Fig. 3 (a).

VII. NUMERICAL RESULTS

We implement the BSMOR on a Linux workstation (P4 2.66GHz, 1G RAM). The mesh structures of the substrate are generated from the typical mixed signal circuit application. In this section, we first investigate the accuracy of BSMOR, then study their scalabilities by increasing the circuit size and number of ports. As an example, we also present the noise map generation for a 256-contact array injected by a frequency-varying ring oscillator at dc and 10GHz.

A. Sparsity Preservation

Due to the structured construction of \tilde{V} by (13), BSMOR preserves the structure and sparsity of \tilde{G} , \tilde{C} matrices even after the reduction. For example, for the 256x256 RC-mesh, Fig. 4 shows the structure of these two state matrices before and after a 16×16 BSMOR reduction. The \tilde{G} , \tilde{C} matrices show 72% and 93% sparsification ratio, respectively. It is another advantage to use BSMOR other than PRIMA, as PRIMA generates a fully dense state matrices after the reduction. Moreover, the sparsification ratio increases when increasing the block number. It is not surprising as conceptually when a block contains only one element, the “reduced” state matrices become exactly the same as the original sparse state matrices.

B. m q-pole Matching

To verify the partitioning algorithm in Section IV, a non-uniform RC mesh with 32 basic blocks is used. Each block has different magnitude RC value in the range 1 1000X. The number of connections between any pair of blocks are also different. The non-zero pattern of elements in G matrix (512×512) is shown in Fig. 5 (a), and the C matrix has the similar structure. SVD is used to determine the rank number of V matrix as 8, and K-means-clustering is used to partitioning the V into 7 blocks (128, 64, 26, 38, 64, 64, 64) as shown in Fig. 5 (b). Based on this partitioning, the structure projection matrix \mathcal{V} is obtained to project the original system matrices. The poles of the original model, reduced model by PRIMA (8th-order), and the reduced model by BSMOR (8th-order) are compared in Fig. 6. Clearly, PRIMA can only approximate 8 poles of the original model, but BSMOR can match 64 poles.

C. Frequency Response Comparison

For a uniform 256x256 RC-mesh (320K circuit elements), Fig. 7 compares frequency responses at one port between the original circuit and reduced models by PRIMA, 2×2 BSMOR, and 8×8 BSMOR. The partitioning algorithm in Section IV found that the partition number is 8 and each block has same size. Clearly, with 10th iteration the 8×8 BSMOR is

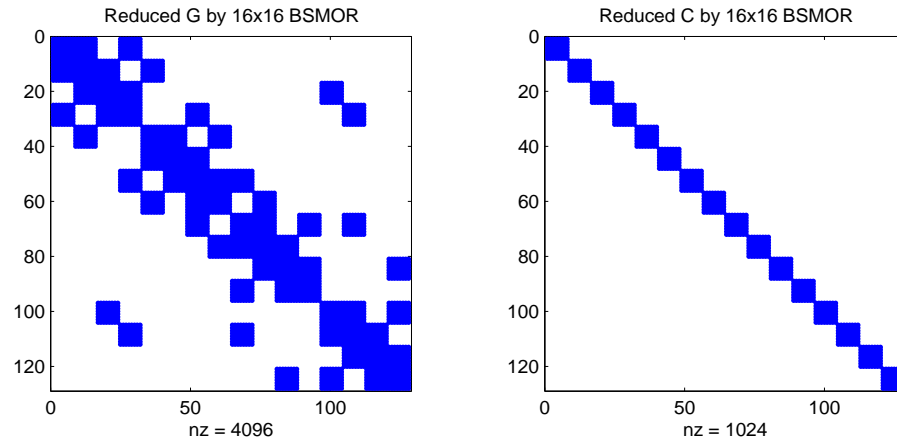


Fig. 4. Non-zero patterns for G , C matrices of a uniform RC-mesh (256×256) after a 16×16 BSMOR reduction with 8 iterations, where NZ is the number of non-zero.

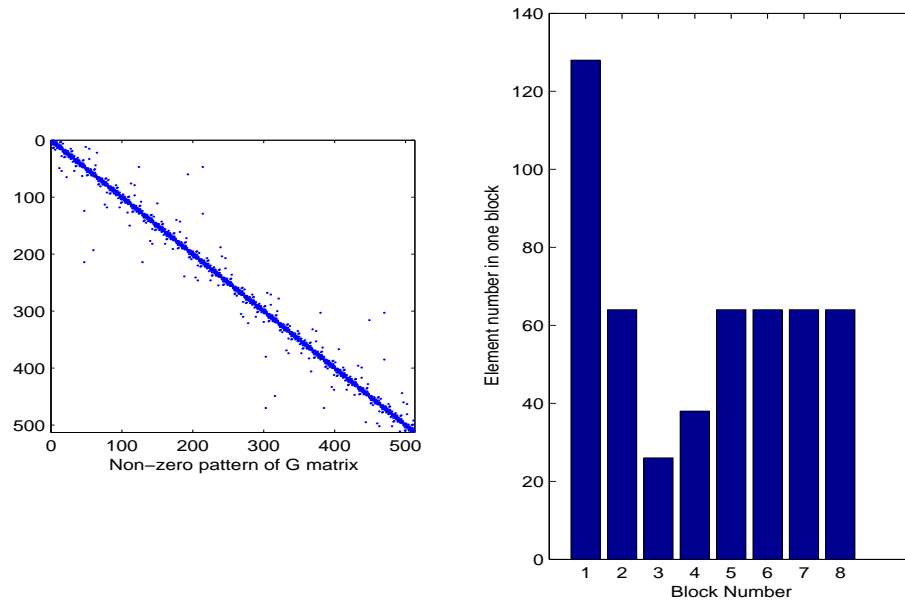


Fig. 5. (a) The nonzero pattern of input G matrix. (b) The partitioning result from SVD and K-means-clustering.

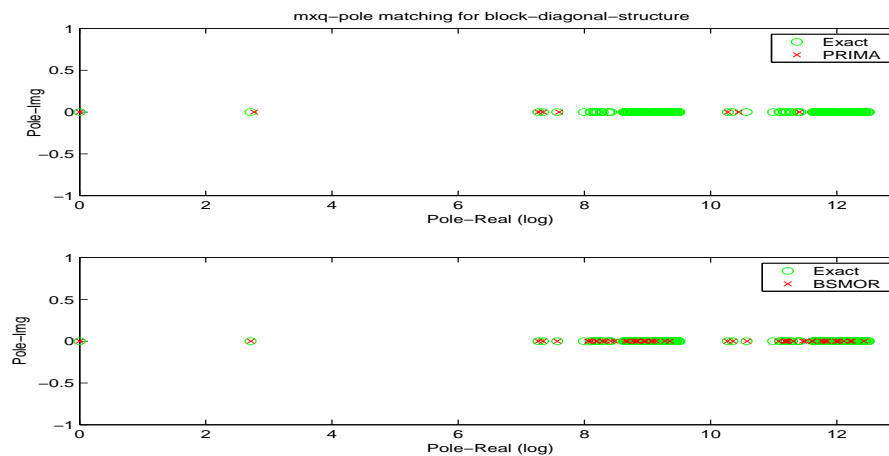


Fig. 6. m q poles matching comparison of PRIMA and BSMOR.

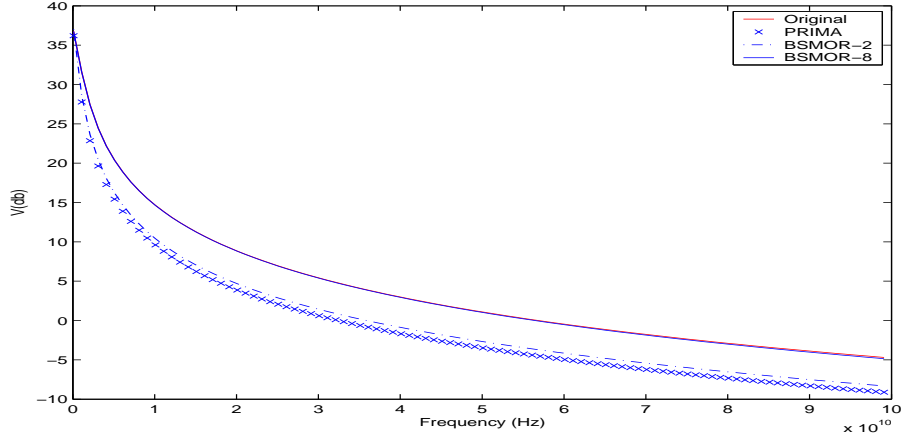


Fig. 7. Frequency responses of the BSMOR, PRIMA, and original model at one port of a uniform mesh (256x256) after 10 iterations.

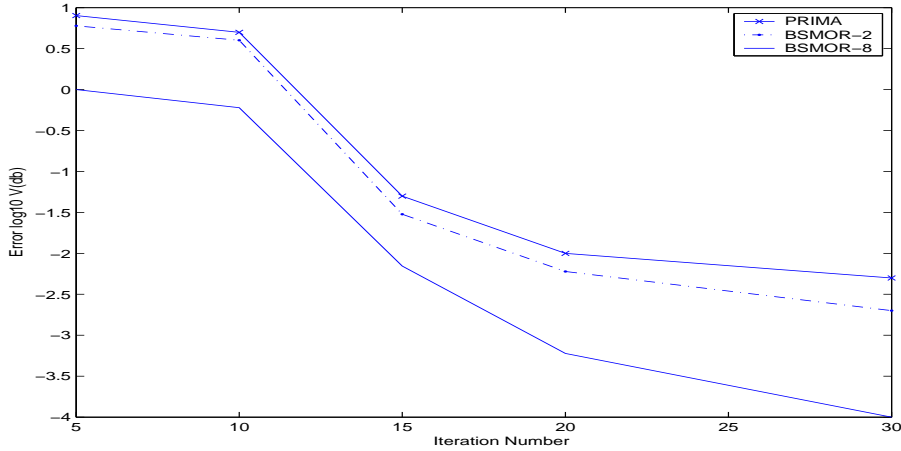


Fig. 8. Maximum errors of the frequency response of the BSMOR and PRIMA for increasing order models of a uniform mesh (256x256) up to 20GHz.

identical to the original circuit response but PRIMA and 2×2 BSMOR are still not converged. Fig. 8 further compares the maximum error of frequency responses by PRIMA, 2×2 , and 8×8 BSMOR vs. the iteration number during the reduction. In the same iteration, it shows that using more partitions (block number) to construct the projection matrix can have better accuracy than using less partitions as PRIMA does. In other words, BSMOR can generate more compact model with improved pole matching ability.

We further present the result of a reduced non-uniform mesh composed by 4 submeshes with different sizes (64x64-64x64-256x256-256x256). We use three partitioning approaches: (i) a fine-uniform partitioning with block size 16 and total 40 blocks; (ii) a coarse-uniform partitioning with block size 40 and total 16 blocks; and (iii) SVD and K-means-clustering based partitioning, which results in a non-uniform partition with two block size (16, 64) and total 16 blocks. As shown in Fig. 9, after 10 iterations, the responses are visually identical for the original model, the reduced model from (i) and (iii), but the coarse-uniform partitioning (ii) does not converge. It shows that the accurate reduced model needs to be generated from a projection matrix with the partition according to the structure of the original matrix, rather than a general 2×2

partition as SPRIM does. Moreover, the reduction time of (ii) and (iii) is similar, and is 4X (4.17s vs. 20.38s) faster than the fine-uniform partitioning (i) does.

D. Scalability Study

We study the efficiency of the reduction convergence by BSMOR and PRIMA. Different block numbers are used according to the different circuit size. We set an error bound as shown in Table I, defined by the maximum error of the frequency response at one port up to 20GHz. We then perform reductions of BSMOR and PRIMA by increasing their iterations until that their accuracies meet the bound. As shown in Table I, BSMOR uses less iterations (≤ 8) to meet the error bound than PRIMA does. As a result, the reduction time of BSMOR is also smaller than that of PRIMA. For example, for a largest mesh circuit with 1M elements, BSMOR achieves 20X (240.22s vs. 4982.76s) speedup under the error bound $1e-4$. Note that a relative small block number (64) is chosen for the largest circuit (1M) here. This is due to the fact that BSMOR needs additional steps to construct the projection matrix, and it results in a little bit larger state matrix that introduce the cost of matrix-vector multiplication. Hence the increase of the speedup is slowed if we choose large

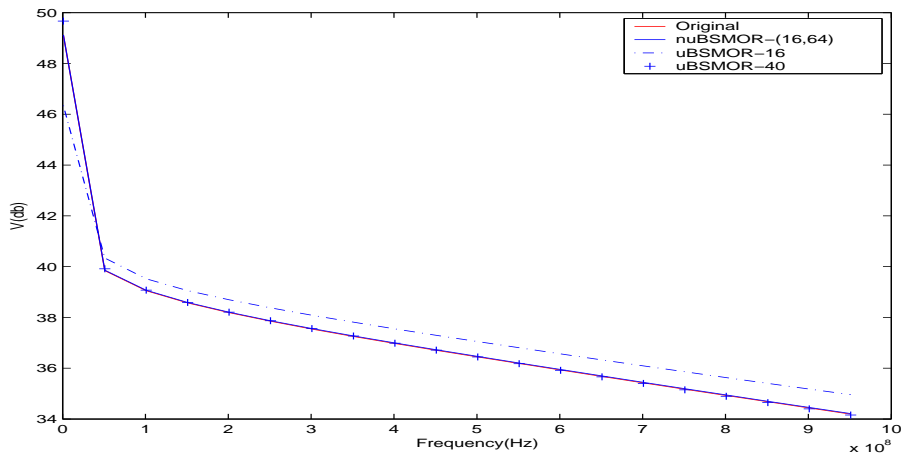


Fig. 9. Frequency responses of the BSMOR by the non-uniform partition, uniform partition, and original model at one port of a nonuniform mesh (64x64-64x64-256x256-256x256).

block number. In general, the result shows that with more partitions to construct a project matrix, BSMOR can match more poles than PRIMA does and hence the reduction time can be significantly reduced under the same accuracy.

We further study the simulation time scalability of the partitioned macro-model by BBDC. PRIMA is used to generate the flat macro-model, BSMOR is used to generate the partitioned macro-model with hierarchy, and different block numbers are used to generate the macro-model according to the port number. Each reduced block contains 10 ports. The original, flat and partitioned models are all simulated in frequency domain up to 20GHz. The maximum error of the frequency response (relative to the original model) up to 20GHz at a selected port is used for comparison.

We observe that when the port number is less than 50 ports the simulation time of the partitioned macro-model is up to 30X times faster than the flat macro-model with a similar accuracy. This speedup comes from two aspects: (i) the cost of the eigen-decomposition to construct flat macro-model is reduced by BSMOR as the sparsity of reduced state matrices is reserved; On the other hand, PRIMA produces a dense reduced state matrices that are computation expensive during the eigen-decomposition; (ii) the partitioned solution further reduces the simulation time as no expensive computation is involved for the large system matrix. To achieve a similar efficiency for the circuits with the large number of ports (≥ 100), we further use the hierarchical clustering (degree 10) with the sparsification (5% error bound) to control the size and sparsity of the coupling blocks. For 1-level and 2-level hierarchical solution, we sparsify the admittance matrices at bottom level, and second level, respectively. Since the error at local matrix can propagate up, we find the solution by sparsification at 1-level partition is less accurate than that at 2-level partition. Moreover, we find that the flat macro-model can not be completed for a 400-port circuit. A clear scalability trend is shown in Fig. 10. We find that the simulation time of the flat macro-model grows up quickly. It shows the similar trend as the original model. This is due to the fact that the dense matrix structure degrades the overall performance when

compared to the original larger but sparser matrix. In contrast, with the use of the BBDC analysis, the simulation time grows much slower than the flat macro-model.

E. Map of Substrate Noise Spectrum

We then apply the partitioned macro-model to generate a map of substrate noise spectrum. The injection current of a frequency-varying ring oscillations is characterized at $f_0 = 100MHz, 1GHz$. The maximum currents are characterized in time domain and then FFT (2048 samplings) is used to obtain the current envelope in frequency domain. The substrate considered here is a $3mm \times 3mm$ plane with a $200um$ thick p-type substrate ($\sigma = 0.1[\Omega cm]^{-1}$). We assume that the contacts are in a 16×16 array, and all the noise-current injection sources (ring oscillators) are placed diagonally in the array. The original substrate circuit is a 256×256 RC-mesh with 320K elements, and we apply 32×32 BSMOR to obtain a 256-port macro-model, representing a 16×16 contact array. The reduction time is about 120s. Fig. 11 shows the map of the noise envelope (voltage bounce magnitude) at dc and $10GHz$. Clearly, reducing the central clock frequency from 1GHz to 100MHz can reduce 25db peak noise at the high frequency (10GHz), but the noise envelope at dc is not reduced. Moreover, the substrate noise coupling is localized at dc but it can diffuse across the contact array at 10GHz. As we assume a high conductivity substrate, the use of the guard ring is effective for this type of substrate. A p^+ -guard ring is used for the isolation with the conductivity $\sigma = 100.0[\Omega cm]^{-1}$. We model the effect of this isolation by changing the surrounding resistance of the contact for each ring oscillator. As shown in Fig. 12, by using a guard ring at 10GHz for $f_0 = 1GHz$, the substrate noise is confined around the injection sources at the diagonal of the contact array.

VIII. CONCLUSION

In this paper, we have proposed a block structure preserving model reduction (BSMOR). We found that compared to PRIMA, BSMOR can match $(m-1)q$ more poles for the *block diagonal structure* system, and for the general system,

Ckt	elements	err-bound	BSMOR			PRIMA	
			block#	iter#	time	iter#	time
mesh1	1K	1e-8	2x2	4	0.03s	10	0.09s
mesh2	10K	1e-8	8x8	6	0.07s	20	0.28s
mesh3	80K	1e-6	16x16	6	0.42s	30	3.82s
mesh4	160K	1e-6	16x16	6	5.14s	40	46.98s
mesh5	320K	1e-4	32x32	6	10.27s	60	104.62s
mesh6	1M	1e-4	64x64	8	240.22s	80	4982.76s

TABLE I

COMPARISON OF THE REDUCTION TIME OF BSMOR AND PRIMA UNDER THE SAME ACCURACY UP TO 20GHZ.

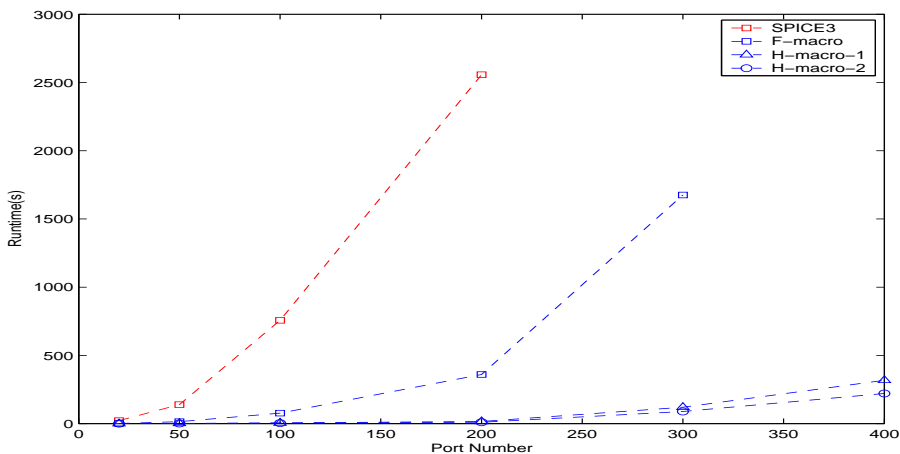
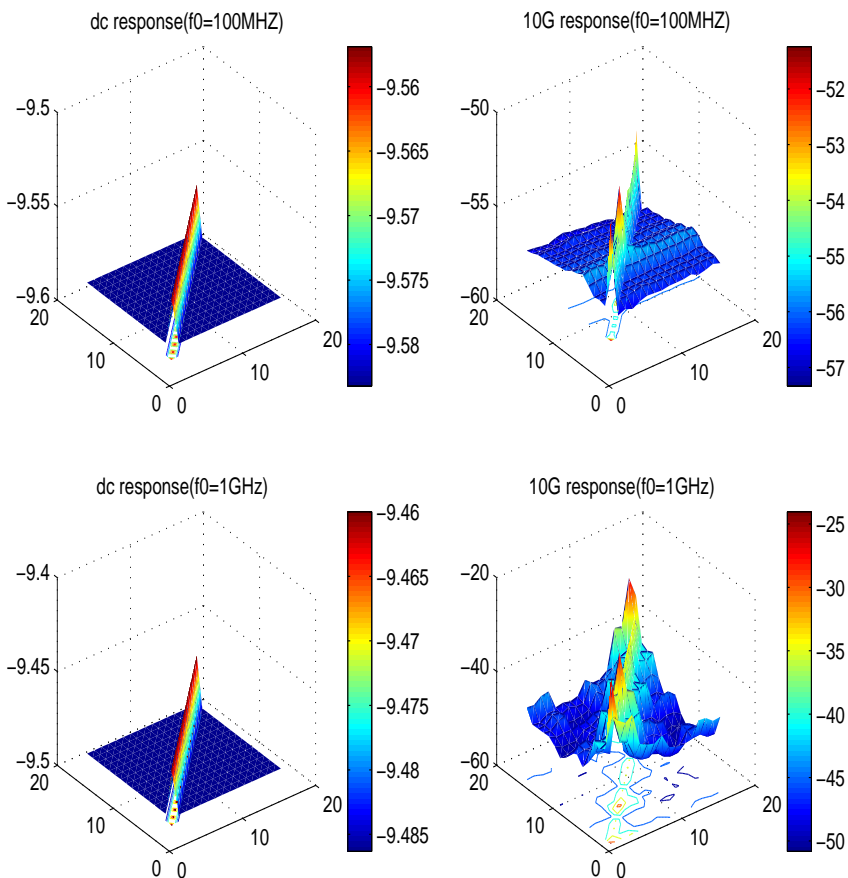


Fig. 10. The scalability trend of simulation time for the original model, flat macro-model, partitioned models with different hierarchical levels.

Fig. 11. A noise map with 16x16 contacts array injected by frequency-controllable ring oscillators at $f_0=100\text{MHz}$ and $f_0=1\text{GHz}$.

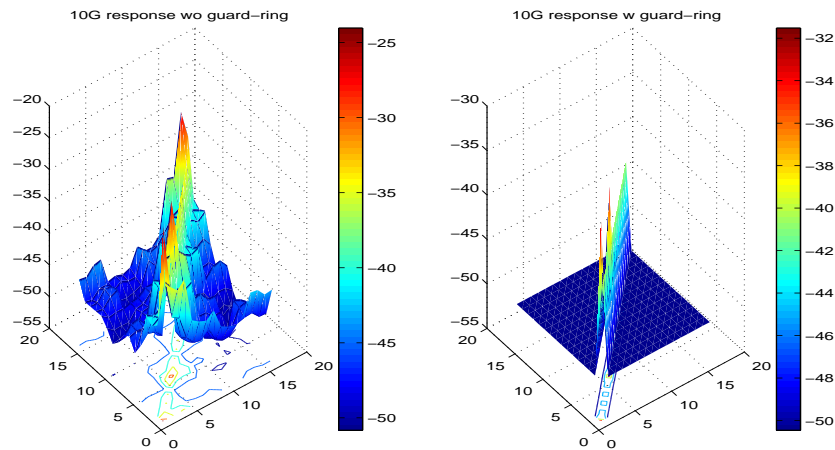


Fig. 12. A noise map at high frequency 10GHz ($f_0=1\text{GHz}$) with/without guard rings.

the additional $(m - 1)q$ poles can closely approximate the poles of original system by using the partitioning based on SVD and K-means-clustering. This property in turn improves the model reduction efficiency compared to PRIMA under the same error bound. For a circuit with 1M elements, BSMOR has a 20X smaller reduction time than PRIMA does. As BSMOR preserves the structure of state matrices, it generates *sparse* reduced state matrices. For a circuit with 320K elements, the reduced state matrices (\mathcal{G} , \mathcal{C}) has 72% and 93% sparsification ratio after a 16×16 BSMOR reduction. It leads to an efficient construction of a MIMO macro-model when using eigen-decomposition. To be able to handle the resulting macro-model with large number of ports, we further used bordered-block diagonal partition with hierarchical clustering (BBDC) to decompose the macro-model into blocks with the manageable size. The experiment shows that BBDC reduces 30X simulation time than the original macro-model.

To construct the structured-projection matrix \mathcal{V} , we in this paper first obtain V by applying PRIMA to the entire \mathcal{G} , \mathcal{C} matrices, and then partition V using SVD and K-means-clustering algorithm. As all operations are on the entire matrix, its computation effort could be too big for large scale circuits, although fewer iterations are needed to achieve higher accurate reduced model compared to PRIMA. To reduce the computation cost, we plan to study how to construct the structured block projection \mathcal{V} more efficiently.

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