# Fast algorithms and numerical methods for the solution of Boundary Element Methods

Session 6: Low rank approximations

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# Outline of the boundary element method

#### Illustration with the EFIE with Dirichlet Boundary Condition

Step 1: Solve the boundary integral equation

$$\int_{\Gamma} G(\mathbf{x} - \mathbf{y}) p(\mathbf{y}) dS_{\mathbf{y}} = -u^{inc}(\mathbf{x}), \quad \mathbf{x} \in \Gamma$$

- · Linear system to solve
- · Unknowns only on the boundary

Step 2: Invoke the boundary integral representation for the evaluation of the quantities at interior points (boundary excluded)

$$u^+(\mathbf{x}) = \int_{\Gamma} G(\mathbf{x} - \mathbf{y}) p(\mathbf{y}) dS_{\mathbf{y}}, \quad \mathbf{x} \in \Omega^+/\Gamma.$$

• Cost reduced matrix-vector product: p(y) already known on  $\Gamma$ 

#### How to reduce the costs of the BEM?

BIE to solve (EFIE): 
$$\int_{\Gamma} G(x-y)p(y)dS_y = -u^{inc}(x), \quad x \in \Gamma$$

BEM discretization  $\Rightarrow$  fully-populated system  $\mathbb{A} p = b$ 

Assembly of the matrix system and matrix-vector product:  $O(N^2)$ 

#### How can we reduce the costs?

- Not possible to speed-up the solution of the initial system
- But it is possible for an approximate system

$$\mathbf{A} := \begin{bmatrix} -2 & 4 & 6 & -3 \\ 4 & -8 & -12 & 6 \\ -6 & 12 & 18 & -9 \\ -8 & 16 & 24 & -12 \end{bmatrix} = \begin{bmatrix} 1 \\ -2 \\ 3 \\ 4 \end{bmatrix} \begin{bmatrix} -2 & 4 & 6 & -3 \end{bmatrix}$$

 BEM system is not low-rank but it can be approximated by a low-rank system: reduction of storage and solution time

#### Algebraic fast BEM

Example with 1 Gauss point per element and a  $\mathbb{P}^0$  interpolation

$$\left[\begin{array}{c}b_1\\ \vdots\\ b_N\end{array}\right] = \left[\begin{array}{c}w_{\Gamma_1}\\ & \ddots\\ & w_{\Gamma_N}\end{array}\right] \left[\begin{array}{c}G(\mathbf{x}_i,\mathbf{y}_j)\\ & \end{array}\right] \left[\begin{array}{c}w_{\Gamma_1}\\ & \ddots\\ & w_{\Gamma_N}\end{array}\right] \left[\begin{array}{c}p_1\\ \vdots\\ p_N\end{array}\right]$$

If we can find  $\mathbb{U}$   $(N \times r \text{ with } r \ll N)$  and  $\mathbb{V}$   $(N \times r \text{ with } r \ll N)$  such that  $\mathbb{G} \simeq \mathbb{U} \mathbb{V}^T$ , it follows a similar approximation for  $\mathbb{A} \simeq \mathbb{U}_A \mathbb{V}_A^T$ .

$$\begin{bmatrix} b_1 \\ \vdots \\ b_N \end{bmatrix} = \underbrace{ \begin{bmatrix} w_{\Gamma_1} \\ & \ddots \\ & w_{\Gamma_N} \end{bmatrix}}_{\mathbb{U}_A \text{ of size } N \times r} \underbrace{ \begin{bmatrix} w_{\Gamma_1} \\ & \ddots \\ & w_{\Gamma_N} \end{bmatrix}}_{\mathbb{V}_A^T \text{ of size } r \times N} \begin{bmatrix} p_1 \\ \vdots \\ p_N \end{bmatrix}$$

Advantages of this algebraic approach?

- Factorization of matrix-vector product  $\mathbb{A} \boldsymbol{p} = \mathbb{U}_A(\mathbb{V}_A^T \boldsymbol{p})$
- · Possibility to combine with a direct solver

# Singular Value Decomposition

#### Rank of a matrix:

Column rank: max # of linearly independent column vectors Column and row ranks are equal ⇒ rank of the matrix

**Definition (Singular Value Decomposition)**  $\mathbb{M} \in \mathbb{C}^{m \times n}$  with rank( $\mathbb{M}$ ) = r. The Singular Value Decomposition

(SVD) of M is the choice of two orthogonal basis

- $\mathbf{v}_1, \dots, \mathbf{v}_r$  of row space of M (right singular vectors)
- and  $\mathbf{u}_1, \dots, \mathbf{u}_r$  of column space of M (left singular vectors)
- such that  $\mathbb{M} v_i = \sigma_i \mathbf{u}_i, \ \sigma_1 \geq \sigma_2 \geq ... \geq \sigma_r \geq 0$  (singular values)

#### Link with the eigendecomposition

The left singular vectors of M are eigenvectors of MM\* The right-singular vectors of M are eigenvectors of M\*M The non-zero singular values of M are the square roots of the non-zero eigenvalues of MM\* and M\*M

### Singular Value Decomposition: matrix form

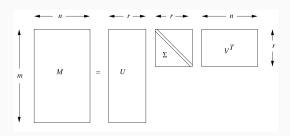
**Theorem (Singular Value Decomposition)**  $\mathbb{M} \in \mathbb{C}^{m \times n}$ , there exists a factorization of  $\mathbb{M}$  of the form  $\mathbb{M} = \mathbb{U}\Sigma\mathbb{V}^*$ 

- $\mathbb{U}$  and  $\mathbb{V}$  are unitary matrices:  $\mathbb{U}^*\mathbb{U} = \mathbb{I}_m$  and  $\mathbb{V}^*\mathbb{V} = \mathbb{I}_n$
- Σ is a diagonal matrix (singular values)

#### The storage is reduced to O(mr + r + nr)



G.H. Golub and C.F. Van Loan. Matrix computations. JHU Press, 2012.



# **SVD** and low-rank approximations

The SVD does not give an approximation but only a factorization

#### **Definition (Truncated SVD)**

 $\mathbb{M}_r$  is the SVD of  $\mathbb{M}$  truncated to the r largest singular values

$$\mathbb{M}_r = \sum_{i=1}^r \mathbb{U}_i \Sigma_{ii} \mathbb{V}_i^*$$

The numerical rank depends on the used norm

$$k(\varepsilon) := \min\{r \mid ||\mathbb{M} - \mathbb{M}_r|| \le \varepsilon ||\mathbb{M}||\}$$

Unitary invariant norm ||UMV|| = ||M|| for all unitary matrices U and V

- Frobenius norm:  $||\mathbf{M}||^2_F = \sum_{i,j} |\mathbf{M}_{ij}|^2$ 
  - Easy to compute (if M is known)
- Spectral or 2-norm:  $||M||_2 = \sigma_1$  ( $\sigma_1$  largest singular value)
  - Need to compute the SVD
- Frobenius norm is always at least as large as the spectral radius  $||M||_2 \le ||M||_F \le \sqrt{r} ||M||_2$

#### **Best low-rank approximation**

# Theorem (Eckart-Young, Best low rank approximation) $\mathbb{M} \in \mathbb{C}^{m \times n}$ with $m \ge n$ , and ||.|| a unitary invariant norm. The best rank-k approximation $\mathbb{M}_r$ of $\mathbb{M}$ defined such that

$$\mathbb{M}_r := \min \Big\{ ||\mathbb{M} - \mathbb{R}|| \quad | \quad \mathbb{R} \in \mathbb{C}^{m \times n}, rank(\mathbb{R}) \le r \Big\} = ||\mathbb{M} - \mathbb{M}_r||$$

$$is \quad \mathbb{M}_r = \sum_{i=1}^r \mathbb{U}_i \Sigma_{ii} \mathbb{V}_i^*, \quad \textit{with} \quad \mathbb{M} = \mathbb{U} \Sigma \mathbb{V}^*.$$

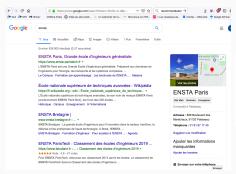
In addition, 
$$||\mathbb{M} - \mathbb{M}_r||_F^2 = \sum_{i=r+1}^n \sigma_i^2$$
 and  $||\mathbb{M} - \mathbb{M}_r||_2 = \sigma_{r+1}$ .

Truncated SVD is the best low-rank approximation for  $L^2$ -norm.

# Low-rank approximations: finding the main information

#### Representing concepts hidden in massive datasets: matrices are used to

- Evaluate the importance of Web pages: Pagerank algorithm (number of occurrences is easy to fool, add the links between pages)
- Community detection: social networks, protein interaction network
- Recommendation systems: Amazon, Netflix





# Finding concepts underlying movies

	M1	M2	МЗ	M4	M5
Jill	3	1	1	3	1
Jane	1	2	4	1	3
Joe	3	1	1	3	1
Jack	4	3	5	4	4

# Finding concepts underlying movies

	M1	M2	МЗ	M4	M5
Jill	3	1	1	3	1
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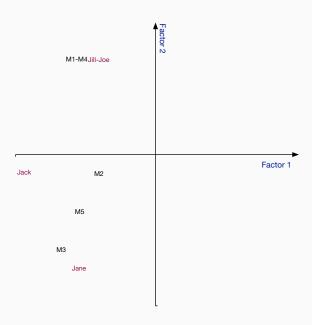
=	U	S	V'	=

-0.3460	0.5294
-0.4190	-0.6515
-0.3460	0.5294
-0.7649	-0.1221

11.822	0	
0	3.9039	

-0.4698	-0.3235	-0.5238	-0.4698	-0.4237
0.5217	-0.1564	-0.5527	0.5217	-0.3545

# **SVD Maps Users and Items Into Latent Space**

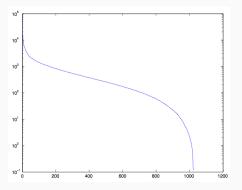


# Low-rank approximations: finding the main information

Image Compression: the goal is to reduce the storage

- Images represented as matrices of size n times m pixels
- · Gray scale images: 1 number per pixel
- Color images: 3 numbers per pixel (red, green and blue)

SVD: form the best rank-r approximations for the matrix



# Low-rank approximations: finding the main information

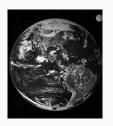
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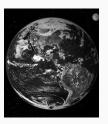
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SVD: form the best rank-r approximations for the matrix









$$r = 10$$

r = 50

k = 200

r = 1024

Truncated SVD to remove the redundant information

#### Low-rank matrices

If we have a low-rank representation of the matrix:  $M = AB^T$ 

- with  $\mathbb{A} \in \mathbb{R}^{m \times r}$  and  $\mathbb{B} \in \mathbb{R}^{n \times r}$
- Then the storage is reduced from mn to r(m+n)

Acceleration of the matrix-vector multiplication:  $\mathbf{M}x = \mathbf{y}$ 

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- Then the storage is reduced from mn to r(m+n)

Acceleration of the matrix-vector multiplication:  $\mathbf{M}\mathbf{x} = \mathbf{v}$ 

- Step 1:  $\mathbf{w} \leftarrow \mathbb{B}^T \mathbf{x}$
- Step 2: y ← Aw
- The number of operations is reduced from O(mn) to O(r(m+n))

# **Definition (Low-rank matrices)** $\mathbb{M} \in \mathbb{R}^{m \times n}$ of rank r is called low-rank if

$$r(m+n) \ll m.n$$

We will always use this representation for low-rank matrices

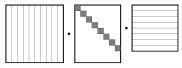
# Computing a low-rank approximation

The truncated SVD gives the best low-rank approximation But computing SVD too expensive:  $O(rN^3)$  (r: rank of approximation)

- If we know the SVD:  $\mathbb{M} = \mathbb{U}\Sigma\mathbb{V}^*$ 
  - Direct solver: compute the pseudo-inverse  $\mathbb{M}^+ = \mathbb{V}\Sigma^+\mathbb{U}^*$
  - Iterative solver: compute the approximation  $\mathbb{M}=\mathbb{AB}^*=\mathbb{U}\Sigma\mathbb{V}^*$  to accelerate the matrix-vector product

#### Is the SVD the only way to compute a low-rank approximation?

Need of a factorization but not of all the properties of the SVD SVD requires all entries of a matrix to construct low-rank approx.



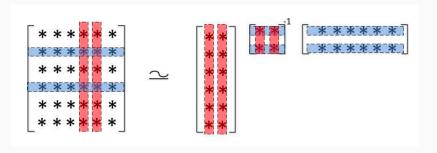
Savings if we use only a small part of the entries

# Skeleton decomposition

**Definition (Skeleton decomposition)**  $\mathbb{A} \in \mathbb{R}^{m \times n}$ , rank  $\mathbb{A}$ =r. There exists a non-singular submatrix  $\hat{\mathbb{A}} \in \mathbb{R}^{r \times r} \hat{\mathbb{A}} = \mathbb{A}(\hat{I}, \hat{J}) \text{ with } \mathbb{A} = \mathbb{C}\hat{\mathbb{A}}^{-1}\mathbb{R}, \ \mathbb{C} = \mathbb{A}(I, \hat{J}), \ \mathbb{R} = \mathbb{A}(\hat{I}, J)$ 



Goreinov, Tyrtyshnikov and Zamarashkin. A Theory of Pseudoskeleton Approximations. Linear Algebra and its Applications, 1997.



# **Skeleton decomposition**

#### Sketch of the proof

- By definition of the rank, since A is of rank r there exists an invertible submatrix of A, Â of size r x r
- It follows the definition of  $\hat{I}$ ,  $\hat{J}$ ,  $\mathbb{R}$  and  $\mathbb{C}$

Noting 
$$\mathbb{A} = \begin{bmatrix} \alpha \mathbb{A}_2 & \mathbb{A}_2 & \beta \mathbb{A}_2 \\ \alpha \hat{\mathbb{A}} & \hat{\mathbb{A}} & \beta \hat{\mathbb{A}} \\ \alpha \mathbb{A}_7 & \mathbb{A}_7 & \beta \mathbb{A}_7 \end{bmatrix}$$
, it follows  $\mathbb{C}\hat{\mathbb{A}}^{-1}\mathbb{R} = \begin{bmatrix} \mathbb{A}_2\hat{\mathbb{A}}^{-1}\alpha\hat{\mathbb{A}} & \mathbb{A}_2 & \mathbb{A}_2\hat{\mathbb{A}}^{-1}\beta\hat{\mathbb{A}} \\ \hat{\mathbb{A}}\hat{\mathbb{A}}^{-1}\alpha\hat{\mathbb{A}} & \hat{\mathbb{A}} & \hat{\mathbb{A}}\hat{\mathbb{A}}^{-1}\beta\hat{\mathbb{A}} \\ \mathbb{A}_7\hat{\mathbb{A}}^{-1}\alpha\hat{\mathbb{A}} & \mathbb{A}_7 & \mathbb{A}_7\hat{\mathbb{A}}^{-1}\beta\hat{\mathbb{A}} \end{bmatrix}$ 

 Finally, use the fact that rows and columns are linear combinations of the rows and columns of Â

Verify that it is correct on a small matrix of rang 3

# **Fully-pivoted Cross Approximation**

Starting point: Every rank r matrix is the sum of r matrices of rang 1 Principle: iteratively removing a row and a column of the matrix

• Successive approximations applied to the remainder  $\mathbb{A} = \mathbb{A}_k + \mathbb{R}_k, \quad \mathbb{R}_k = \mathbb{A} - \sum_{\ell=1}^k \mathbf{u}_\ell \mathbf{v}_\ell^T$ 

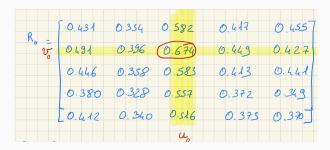
$$A = A_k + R_k, \quad R_k = A - \sum_{\ell=1}^k \mathbf{u}_\ell \mathbf{v}_\ell^T$$

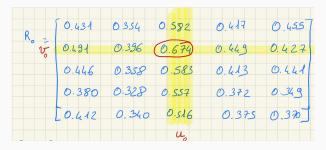
- Similarly to the Gaussian elimination, the pivot is the largest entry of the matrix (to define a stable algorithm)
- At each iteration, we nullify in the remainder the rows and columns dependent from the pivot row and column

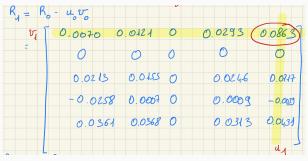
$$\mathbb{A} = \left[ \begin{array}{ccc} a_{11} & a_{12} & \alpha a_{11} \\ a_{21} & a_{22} & \alpha a_{21} \\ a_{31} & a_{32} & \alpha a_{31} \end{array} \right] \quad \mathbb{R}_1 = \mathbb{A} - \mathbb{A}(:,1) \mathbb{A}(2,:)/a_{21} = \left[ \begin{array}{ccc} 0 & r_{12} & 0 \\ 0 & 0 & 0 \\ 0 & r_{32} & 0 \end{array} \right]$$

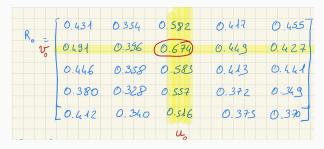
#### At iteration *k*:

- Find the pivot  $(i^*, j^*)$  such that  $(i^*, j^*) = \operatorname{argmax}_{ij} |(\mathbb{R}_k)_{ij}|$
- Compute vectors:  $\mathbf{u}_{k+1} := \frac{(\mathbb{R}_k)_{ij^*}}{(\mathbb{R}_k)_{i^*i^*}}, \mathbf{v}_{k+1} := (\mathbb{R}_k)_{i^*j}$
- Update the approximation:  $\mathbb{A}_{k+1} = \mathbb{A}_k + \mathbf{u}_{k+1} \mathbf{v}_{\iota_{-1}}^T$

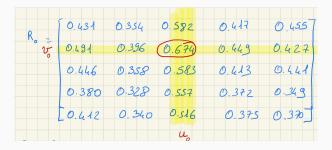








R <sub>2</sub> =	R, - W, V,				/1
	0	0	0	0	0
-	0	0	0	0	0
	0,0155	0,0055	0	0.0003	0
	-0.0255	0.0013	0	0.0023	0
	0.0326)	0.0308	0	0.0166	0



- -

# Fully-pivoted Cross Approximation: pseudo-code

```
Initialization: \mathbb{R}_0 := A, \mathscr{P}_r = \emptyset; \mathscr{P}_c = \emptyset, k = 0 repeat k := k+1 Find the pivot (i^*, j^*) := \operatorname{argmax}_{i,j} |\mathbb{R}_{k-1}(i, j)| \mathscr{P}_r = \mathscr{P}_r \cup \{i^*\}, \quad \mathscr{P}_c = \mathscr{P}_c \cup \{j^*\} \delta_k := \mathbb{R}_{k-1}(i^*, j^*) \mathbf{u}_k := \mathbb{R}_{k-1}(i, j^*) \mathbf{v}_k := \mathbb{R}_{k-1}(i^*, i)/\gamma \mathbb{R}_k = \mathbb{R}_{k-1} - \mathbf{u}_k \mathbf{v}_k until \|\mathbb{R}_k\|_F \le \varepsilon \|A\|_F
```

It requires

steps to generate an approximation of rank  $\emph{r}$ 

It requires

to compute the pivot indices

# Fully-pivoted Cross Approximation: pseudo-code

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Initialization: \mathbb{R}_0 := A, \mathscr{P}_r = \emptyset; \mathscr{P}_c = \emptyset, k = 0 repeat k := k+1 Find the pivot (i^*, j^*) := \operatorname{argmax}_{i,j} |\mathbb{R}_{k-1}(i, j)| \mathscr{P}_r = \mathscr{P}_r \cup \{i^*\}, \quad \mathscr{P}_c = \mathscr{P}_c \cup \{j^*\} \delta_k := \mathbb{R}_{k-1}(i^*, j^*) \mathbf{u}_k := \mathbb{R}_{k-1}(i, j^*) \mathbf{v}_k := \mathbb{R}_{k-1}(i, j^*) \mathbf{v}_k := \mathbb{R}_{k-1}(i, j^*) \mathbb{R}_k = \mathbb{R}_{k-1} - \mathbf{u}_k \mathbf{v}_k until \|\mathbb{R}_k\|_F \le \varepsilon \|A\|_F
```

- It requires O(rmn) steps to generate an approximation of rank r
- It requires all the entries of A to compute the pivot indices

### **Exact reproduction of rank** r matrices

#### **Lemma (Exact reproduction of rank** r **matrices)** Let $\mathbb{A}$ be matrix of rank exactly r. Then the matrix $\mathbb{A}_r$ is equal to $\mathbb{A}$ .

$$\mathbb{A}_r := \sum_{\ell=1}^r \mathbf{u}_\ell \mathbf{v}_\ell^T$$

If rank(A)=r, the algorithm terminates in r steps.

Consistent with the Skeleton decomposition:  $\mathbb{A} \in \mathbb{R}^{m \times n}$ , rank A = r. There exists a non-singular submatrix  $\hat{\mathbb{A}} \in \mathbb{R}^{r \times r}$   $\hat{\mathbb{A}} = \mathbb{A}(\hat{I}, \hat{J})$  with  $\mathbb{A} = \mathbb{C}\hat{\mathbb{A}}^{-1}\mathbb{R}$ ,  $\mathbb{C} = \mathbb{A}(I, \hat{J})$ ,  $\mathbb{R} = \mathbb{A}(\hat{I}, J)$ 

How can we reduce the complexity?

# **Principle of the Partially-pivoted Cross Approximation**

- Fully-pivoted: pivot is the largest entry in the residual
- Partially-pivoted: maximize only for 1 of the 2 indices (the other one is fixed) → only one row or one column is assembled

# Partially-pivoted CA: pseudo-code

```
Initialization: \mathbb{R}_0 := A, i^* = 1, \mathscr{P}_r = \emptyset; \mathscr{P}_c = \emptyset, k = 1
repeat
     Find the pivot column j^* := \operatorname{argmax}_{i} |\mathbb{R}_{k-1}(i^*, j)|
     \delta_k := \mathbb{R}_{k-1}(i^*, j^*)
     if \delta_k == 0 then
          if \#\mathscr{P}_r = n-1 then
               STOP
          end if
     else
          \mathbf{u}_k := \mathbb{R}_{k-1}(:, j^*)
          \mathbf{v}_k := \mathbb{R}_{k-1}(i^*,:)/\gamma
          \mathbb{R}_k = \mathbb{R}_{k-1} - \mathbf{u}_k \mathbf{v}_k
          k := K + 1
     end if
     \mathscr{P}_r = \mathscr{P}_r \cup \{i^*\}, \quad \mathscr{P}_c = \mathscr{P}_c \cup \{j^*\}
     i^* := \operatorname{argmax}_{i \notin \mathscr{P}_r} |\mathbf{u}_k(i)|
until Stopping criterion is fulfilled
```

# **Adaptive Cross Approximation (ACA)**

 $\mathbb{R}_k$  is never explicitly formed

$$\mathbb{R}_k(i,j) = \mathbb{A}(i,j) - \sum_{\ell=1}^k \mathbf{u}_\ell(i) \mathbf{v}_\ell(j)$$

Can we determine the rank k adaptively for a given approximation accuracy  $\varepsilon$ ?

- Fully-pivoted ACA:  $||A A_k||_F \le \varepsilon ||A||_F$
- Partially-pivoted ACA: A is not formed, stagnation-based error estimator

$$||\mathbf{u}_k||_2||\mathbf{v}_k||_2 \le \varepsilon ||\mathbb{A}_k||_F$$

· Optimal computation of the Frobenius norm

$$||\mathbf{A}_k||_F^2 = ||\mathbf{A}_{k-1}||_F^2 + 2\sum_{\ell=1}^{k-1} \mathbf{u}_k^T \mathbf{u}_\ell \mathbf{v}_\ell^T \mathbf{v}_k + ||\mathbf{u}_k||_2^2 ||\mathbf{v}_k||_2^2$$

$$\mathbb{A} = \left[ \begin{array}{ccccc} 6.5 & 31 & -14 & -43 \\ 9.1 & -3 & 11 & 31 \\ 17.6 & -16 & 28 & 80 \\ 26.2 & 50 & -7 & -26 \end{array} \right]$$

First iteration, we set  $i^* = 1$ ,  $\mathscr{P}_r = \{1\}$ 

$$\mathbb{R}_0 = \begin{bmatrix} 6.5 & 31 & -14 & -43 \\ 9.1 & -3 & 11 & 31 \\ 17.6 & -16 & 28 & 80 \\ 26.2 & 50 & -7 & -26 \end{bmatrix}$$

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First iteration, we set  $i^* = 1$ ,  $\mathscr{P}_r = \{1\}$  and find  $j^* = 4$ ,  $\mathscr{P}_c = \{4\}$ 

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We have 
$$u_1 = [1 - 0.7209 - 1.8605 \ 0.6047]^T$$
  $v_1 = [6.5 \ 31 \ -14 \ -43]$ 

Next pivot is  $i^* = 3$  ( $i^* = 1$  already used)

$$\mathbb{A} = \left[ \begin{array}{ccccc} 6.5 & 31 & -14 & -43 \\ 9.1 & -3 & 11 & 31 \\ 17.6 & -16 & 28 & 80 \\ 26.2 & 50 & -7 & -26 \end{array} \right]$$

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We have 
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  $v_1 = [6.5 \ 31 - 14 - 43]$ 

Next pivot is  $i^* = 3$  ( $i^* = 1$  already used)

If we compute the residual (not performed in practice):

$$\mathbb{R}_1 = \left[ \begin{array}{cccc} 0 & 0 & 0 & 0 \\ 13.7860 & 19.3488 & 0.9070 & 0 \\ 29.6930 & 41.6744 & 1.9535 & 0 \\ 22.2698 & 31.25581 & 1.4651 & 0 \end{array} \right]$$

 $||A_1||_F = 127.6636$ ,  $||v_1||_2 ||u_1||_2 = 127.6636$ , convergence not achieved

$$\mathbb{A} = \left[ \begin{array}{ccccc} 6.5 & 31 & -14 & -43 \\ 9.1 & -3 & 11 & 31 \\ 17.6 & -16 & 28 & 80 \\ 26.2 & 50 & -7 & -26 \end{array} \right] \quad \mathbb{R}_1 = \left[ \begin{array}{cccccc} 0 & 0 & 0 & 0 \\ 13.7860 & 19.3488 & 0.9070 & 0 \\ \hline 29.6930 & 41.6744 & 1.9535 & 0 \\ 22.2698 & 31.25581 & 1.4651 & 0 \end{array} \right]$$

 $\mathcal{P}_r = \{1, 3\}. \text{ New row } \mathbb{R}_1: [29.693 \ 41.6744 \ 1.9536 \ 0]$ 

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 $\mathcal{P}_r = \{1, 3\}.$  New row  $\mathbb{R}_1$ : [29.693 41.6744 1.9536 0]

We find  $j^* = 2$ ,  $\mathcal{P}_c = \{4, 2\}$ . New column [0 19.3488 41.6744 31.2558]<sup>T</sup>

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$$u_2 = [0 \ 0.4643 \ 1 \ 0.75]^T$$
 and  $v_2 = [29.6930 \ 41.6744 \ 1.9535 \ 0]$ 

Next pivot is  $i^* = 4$  ( $i^* = 1$  or 3 already used)

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$$||A_2||_F = 113.7962$$
 et  $||v_2||_2 ||u_2||_2 = 68.2826$ 

End of the algorithm: We cannot find a new pivot