

# Image Processing using Graph Laplacian Operator

David Wobrock

ALPINES Team - INRIA Paris  
Laboratoire Jacques-Louis Lions - Sorbonne Université  
KTH, Stockholm  
INSA Lyon

May 17, 2018

# Table of Contents

Introduction

Image processing using Laplacian operator

Parallel implementation

Conclusion

# Background

- ▶ Millions of pictures shared daily
- ▶ Image processing using spectral methods on smartphones (denoising, sharpening, ...)
- ▶ Involves eigenvalue problems, linear algebra and solving dense linear systems
- ▶ Opportunity for high-performance computing and parallelism

# Objective

- ▶ Not necessarily improving image processing
- ▶ Analyse the behaviour of solving large dense systems
- ▶ Large:  $N^2$ ,  $N$  the number of pixels in the input image
- ▶ Dense: affinity and Laplacian matrices are dense because global filter

# Image processing - Global filter algorithm

- ▶  $z$  output image,  $y$  input image,  $W$  data-dependent global filter of size  $N^2$
- ▶  $z_i = \sum_{j=1}^N W_{ij} y_j$
- ▶ A vector of weights for each pixel
- ▶ Global filter:  $z = Wy$

# Image processing - Image as graph

- ▶ Think of the image as a graph
- ▶ Each pixel is a node
- ▶ The graph is complete and the edges are weighted
- ▶ The weight represents the similarity between two pixels/nodes
- ▶ Similarity can be measured by weighting together spatial and color closeness of pixels
- ▶ Bilateral kernel:  $\exp\left(-\frac{\|x_i - x_j\|^2}{h_x^2}\right) \cdot \exp\left(-\frac{\|z_i - z_j\|^2}{h_z^2}\right)$

# Image processing - Filter and Laplacian

- ▶  $K$  affinity matrix,  $\mathcal{L}$  Laplacian matrix,  $W$  filter matrix and  $f$  a function
- ▶ Filter defined as  $W = I - f(\mathcal{L})$
- ▶ Let  $d_i = \sum_j K_{ij}$ ,  $D = \text{diag}\{d_i\}$  and  $\bar{d} = \frac{1}{N} \sum_i d_i$

Laplacian	Formula of $\mathcal{L}$	Symmetric	Spectral Range
Un-normalised	$D - K$	Yes	$[0, n]$
Normalised	$I - D^{-1/2} K D^{-1/2}$	Yes	$[0, 2]$
Random walk	$I - D^{-1} K$	No	$[0, 1]$
"Sinkhorn"	$I - C^{-1/2} K C^{-1/2}$	Yes	$[0, 1]$
Re-normalised	$\alpha(D - K)$ , $\alpha \approx \bar{d}^{-1}$	Yes	$[0, n]$

**Table:** Overview of different graph Laplacian operator definitions.

# Image processing - Sampling and Nyström extension

- ▶ Sample  $p$  pixels of the image ( $\leq 1\%$  of pixels)
- ▶ With  $K_A \in \mathbb{R}^{p \times p}$ ,  $K_B \in \mathbb{R}^{p \times N-p}$  and  $K_C \in \mathbb{R}^{N-p \times N-p}$ 
$$K = \begin{bmatrix} K_A & K_B \\ K_B^T & K_C \end{bmatrix}$$
- ▶ Approximate  $K$  by  $\tilde{K} = \tilde{\Phi} \tilde{\Pi} \tilde{\Phi}^T$
- ▶ With submatrix  $K_A = \Phi_A \Pi_A \Phi_A^T$ ,
$$\tilde{\Pi} = \Pi_A$$
$$\tilde{\Phi} = \begin{bmatrix} \Phi_A \\ K_B^T \Phi_A \Pi_A^{-1} \end{bmatrix}$$
- ▶ Finally,  $\tilde{K} = \begin{bmatrix} K_A & K_B \\ K_B^T & K_B^T K_A^{-1} K_B \end{bmatrix}$



# Image processing - Eigenvalues

- ▶ For the filter, only the largest eigenvalues are needed because the smallest tends to 0
- ▶ There is an equivalence between largest eigenvalues of  $W$  and the smallest ones of  $\mathcal{L}$
- ▶ The smallest eigenvalues can be computed with the inverse power method, which requires solving systems of linear equations
- ▶ But the Nyström extension only works for leading eigenvalues.

# Parallel implementation details

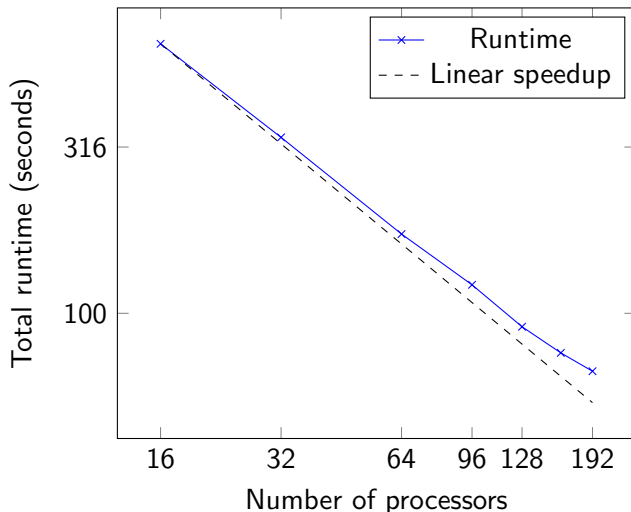
- ▶ C language
- ▶ PETSc (Toolkit for scientific computing in parallel)
- ▶ SLEPc, Elemental

# Full matrix computation result



Figure: Left: input image. Right: sharpened image.

## Runtime of full matrix computation



**Figure:** Total runtime of the algorithm with entire matrix computation (log scale).

# Inverse subspace iteration

---

**Algorithm 1** Inverse subspace iteration

---

**Input:**  $A$  the matrix of size  $p \times p$ ,  $m$  the number of required eigenvalues,  $\varepsilon$  a tolerance

**Output:**  $X_k$  the desired invariant subspace

Initialise  $m$  random orthonormal vectors  $X_0$  of size  $p$

$$R_0 \leftarrow (I - X_0 X_0^T) A X_0$$

For  $k=0, 1, 2, \dots$

**while**  $\|R_k\| > \varepsilon$  **do**

**for**  $i=1$  **to**  $m$  **do**

$$\text{Solve } A X_{k+1}^{(i)} = X_k^{(i)}$$

**end for**

$$X_{k+1} \leftarrow \text{Orthonormalise}(X_{k+1})$$

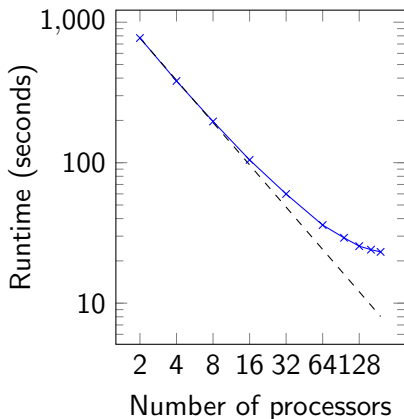
$$R_{k+1} \leftarrow (I - X_{k+1} X_{k+1}^T) A X_{k+1}$$

**end while**

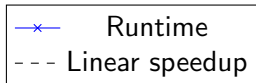
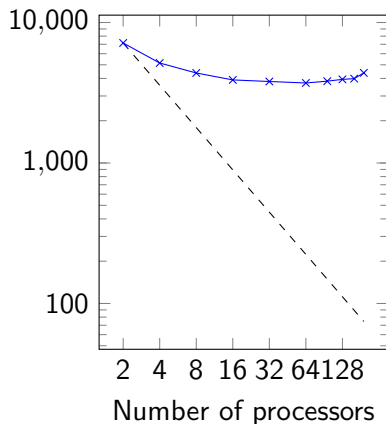
---

# Approximation runtime of approximated computation

50 eigenvalues.



500 eigenvalues.



# Runtime in the inverse subspace iteration algorithm

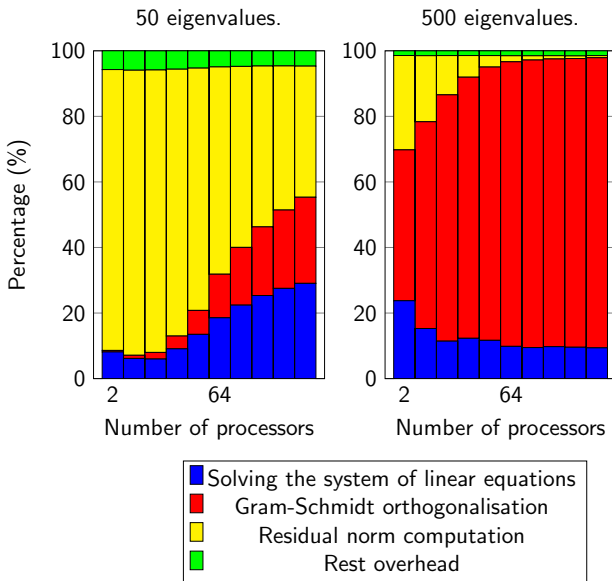
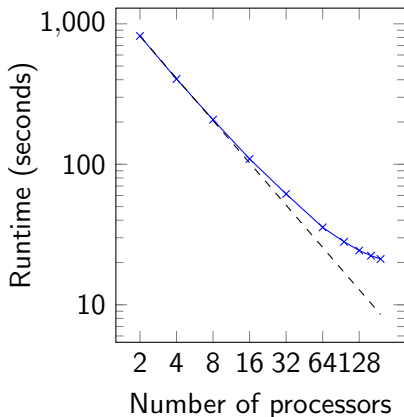


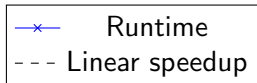
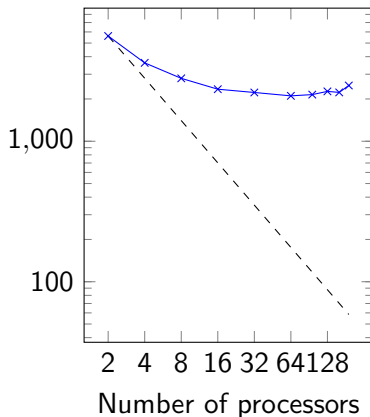
Figure: Proportion of each step in the inverse subspace iteration.

# Skipping the Gram-Schmidt procedure

50 eigenvalues.



500 eigenvalues.





# Skipping Gram-Schmidt more often

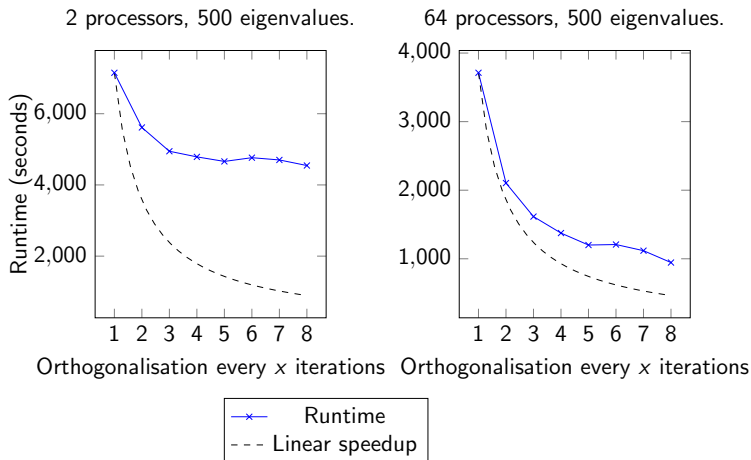


Figure: Runtime of the inverse subspace iteration depending on the amount of Gram-Schmidt procedures.

## Linear solver performances

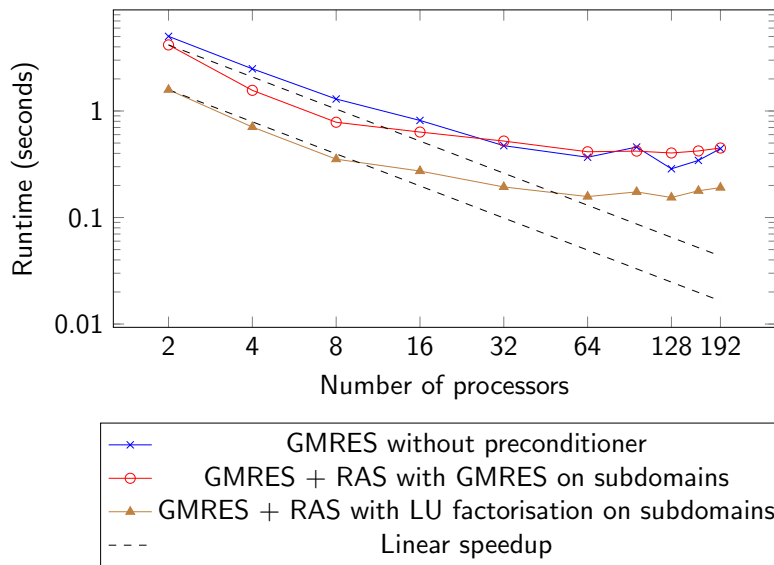


Figure: Runtime of the linear solver for 50 eigenvalues for a  $4000 \times 4000$  matrix (log scale).

## Linear solver performances - big image

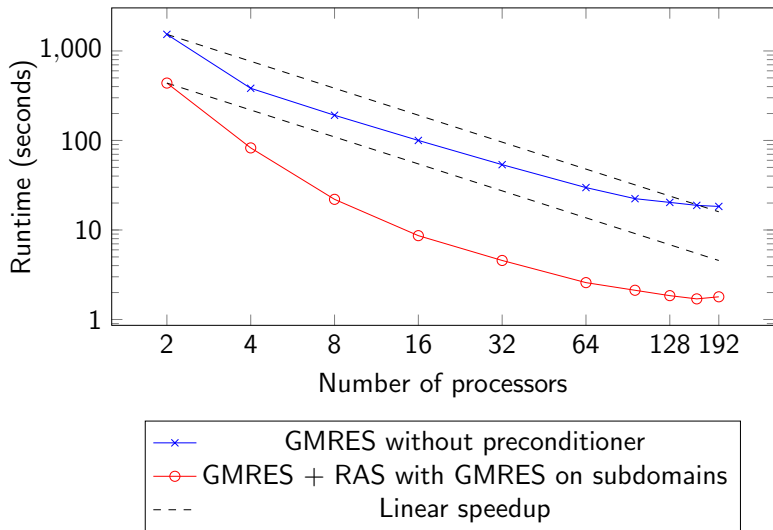


Figure: Runtime of the linear solver for 50 eigenvalues for a  $14400 \times 14400$  matrix (log scale).

# Discussions & perspectives

- ▶ Linear solver with domain decomposition methods as preconditioner
- ▶ Skipping Gram-Schmidt
- ▶ Finished image processing algorithm
- ▶ State-of-the-art performances with a similar method
- ▶ Explore more solvers, domain decomposition methods and other preconditioner