Image Processing using Graph Laplacian Operator

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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

- ightharpoonup 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: $exp(-\frac{||x_i-x_j||^2}{h_x^2}) \cdot exp(-\frac{||z_i-z_j||^2}{h_z^2})$

Image processing - Filter and Laplacian

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

Laplacian	Formula of ${\cal L}$	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"	$I - C^{-1/2}KC^{-1/2}$	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)
- $\begin{array}{l} \quad \text{With } K_A \in \mathbb{R}^{p \times p}, K_B \in \mathbb{R}^{p \times N p} \text{ and } K_C \in \mathbb{R}^{N p \times N p} \\ K = \begin{bmatrix} K_A & K_B \\ K_B^T & K_C \end{bmatrix} \end{array}$
- Approximate K by $\tilde{K} = \tilde{\Phi} \tilde{\Pi} \tilde{\Phi}^T$
- With submatrix $K_A = \Phi_A \Pi_A \Phi_A^T$, $\tilde{\Pi} = \Pi_A$ $\tilde{\Phi} = \begin{bmatrix} \Phi_A \\ K_P^T \Phi_A \Pi_A^{-1} \end{bmatrix}$
- $\blacktriangleright \text{ Finally, } \tilde{K} = \begin{bmatrix} K_A & K_B \\ K_B^T & K_B^T K_A^{-1} K_B \end{bmatrix}$

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.

Image processing - Algorithm recap

 $\begin{tabular}{ll} \textbf{Algorithm 1} & \textbf{Image processing using approximated graph Laplacian} \\ \textbf{operator} \\ \end{tabular}$

```
Input: y an image of size N, f the function applied to \mathcal{L}
Output: \tilde{z} the output image by the approximated filter
   {Sampling}
  Sample p pixels, p \ll N
   {Kernel matrix approximation}
  Compute K_A (size p \times p) and K_B (size p \times (N-p))
  Compute the Laplacian submatrices \mathcal{L}_A and \mathcal{L}_B
   {Eigendecomposition}
   Compute the m smallest eigenvalues \Pi_A and the associated eigen-
  vectors \Phi_A of \mathcal{L}_A
   {Nyström extension and compute the filter}
  See methods of solution proposed by Fowlkes, 2004
  \tilde{z} \leftarrow \tilde{W}v
```

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

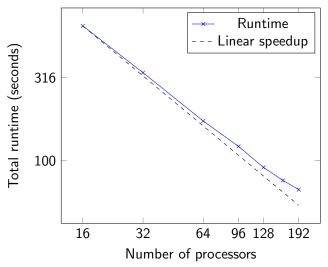


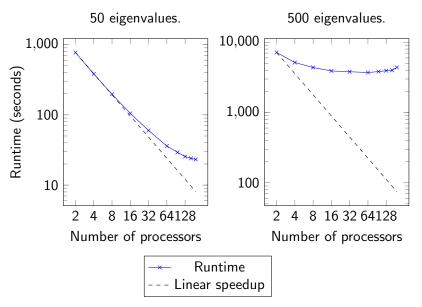
Figure: Total runtime of the algorithm with entire matrix computation (log scale).

Inverse subspace iteration

Algorithm 2 Inverse subspace iteration

```
Input: A the matrix of size p \times p, m the number of required eigen-
  values, \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, ...
  while ||R_k|| > \varepsilon do
      for i=1 to m do
        Solve AX_{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



Runtime in the inverse subspace iteration algorithm

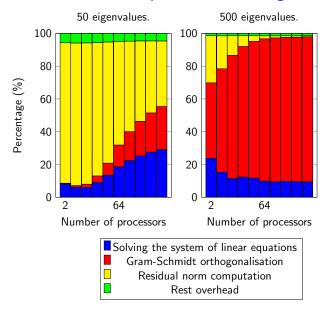
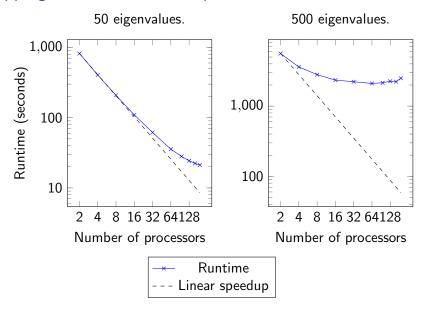


Figure: Proportion of each step in the inverse subspace iteration.

Skipping the Gram-Schmidt procedure



Skipping Gram-Schmidt more often

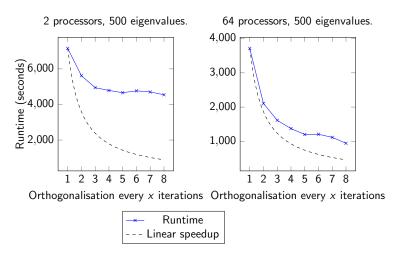


Figure: Runtime of the inverse subspace iteration depending on the amount of Gram-Schmidt procedures.

Linear solver performances

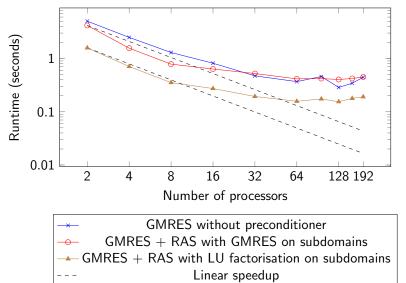


Figure: Runtime of the linear solver for 50 eigenvalues for a 4000×4000 matrix (log scale).

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Linear solver performances - big image

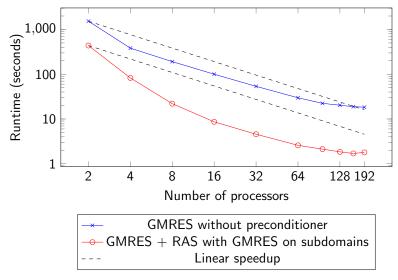


Figure: Runtime of the linear solver for 50 eigenvalues for a 14400×14400 matrix (log scale).

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