

Hierarchical interpolative factorization

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Joint work with Lexing Ying

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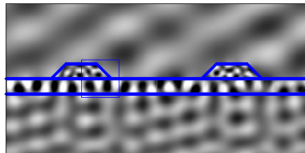
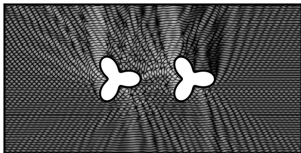
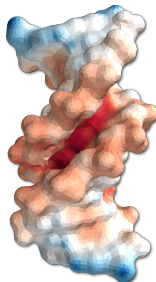
Introduction

Elliptic PDEs in differential or integral form:

$$-\nabla \cdot (a(x)\nabla u(x)) + b(x)u(x) = f(x)$$

$$a(x)u(x) + \int_{\Omega} K(x,y)u(y) d\Omega(y) = f(x)$$

- Fundamental to science and engineering
- Interested in 2D/3D, complex geometry
- Discretize $\rightarrow$ **structured** linear system $Au = f$



Goal: fast, accurate, and robust algorithms to compute $u = A^{-1}f$

Introduction

How to solve? Let $A \in \mathbb{C}^{N \times N}$.

- ▶ If N is small, use **direct** methods (Gaussian elimination, matrix factorization)
 - Stable, robust, rapid updates
 - $O(N^3)$ complexity, infeasible for $N \gtrsim 10^4$
- ▶ If N is large, use **iterative** methods (CG, GMRES, multigrid)
 - Can be accelerated to $O(n_{\text{iter}}N)$ complexity, highly scalable
 - Extremely successful, workhorse of modern scientific computing

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- ▶ What if n_{iter} is large (high contrasts, geometric singularities)?
- ▶ What if there are many RHS's (time stepping, inverse problems)?

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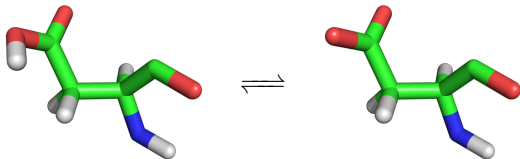
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But ...

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- ▶ What if there are many RHS's (time stepping, inverse problems)?

In certain important environments, there is a need for fast direct solvers.

Example: protein pK_a calculations

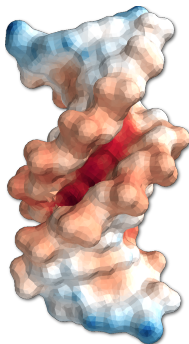


$$pK_a \equiv -\log_{10} \frac{[A][H]}{[AH]} = \frac{\beta}{\ln 10} \Delta G_{AH \rightarrow A+H}^p$$

$$\begin{aligned} \Delta G_{AH \rightarrow A+H}^p &= \Delta G_{AH \rightarrow A+H}^s + \Delta G_A^{s \rightarrow p} - \Delta G_{AH}^{s \rightarrow p} \\ &= \underbrace{\Delta G_{AH \rightarrow A+H}^s}_{\text{experiment}} + \underbrace{\Delta G_A^{s \rightarrow p} - \Delta G_{AH}^{s \rightarrow p}}_{\text{electrostatic only}} \end{aligned}$$

- Ionization behavior is important for enzymatic and structural properties

Example: protein pK_a calculations



- ▶ Linearized Poisson-Boltzmann equation
- ▶ Discretize: $A(\Sigma)u = f(q)$
 - Σ : molecular geometry
 - q : atomic partial charges
- ▶ **Fixed** matrix, one solve for each of n_{titr} titrating sites
- ▶ Conformational flexibility: $O((n_{\text{titr}}n_{\text{rot}})^p)$ perturbed solves

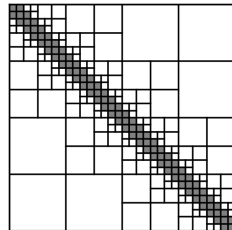
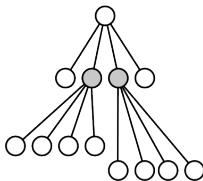
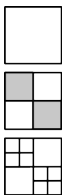
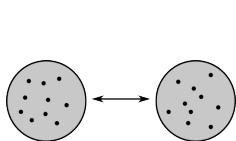
Potential for massive acceleration using fast direct methods.

Related problems:

- ▶ Protein structure prediction, protein-protein docking, protein design
- ▶ Inverse scattering, time stepping, materials design

Previous work

- ▶ **PDEs**: exploit sparsity (multifrontal), reduce to IE-like systems
- ▶ **IEs**: exploit FMM-type hierarchical low-rank structure
- ▶ $\mathcal{H}$ -matrices: $O(N \log^\alpha N)$ but with a large constant
- ▶ HSS matrices/recursive skeletonization: $O(N)$ in 1D, $O(N^{3/2})$ in 2D, $O(N^2)$ in 3D
- ▶ HSS/RS/MF with structured matrix algebra: $O(N)$ in 2D, $O(N^{4/3})$ in 3D



Hierarchical interpolative factorization

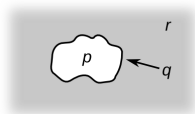
- ▶ RS/MF + recursive dimensional reduction
- ▶ Same idea as using structured algebra but much simpler
- ▶ Explicit matrix **sparsification**, generalized LU decomposition
- ▶ Linear or quasilinear complexity, small constants
- ▶ Unified formalism for IEs and PDEs
- ▶ Works for 2D/3D, adaptive and complex geometry

Tools: sparse elimination, interpolative decomposition, skeletonization

Sparse elimination

Let

$$A = \begin{bmatrix} A_{pp} & A_{pq} \\ A_{qp} & A_{qq} & A_{qr} \\ & A_{rq} & A_{rr} \end{bmatrix}.$$



(Think of A as a **sparse** matrix.) If A_{pp} is nonsingular, define

$$R_p^* = \begin{bmatrix} I & & \\ -A_{qp}A_{pp}^{-1} & I & \\ & & I \end{bmatrix}, \quad S_p = \begin{bmatrix} I & -A_{pp}^{-1}A_{pq} & \\ & I & \\ & & I \end{bmatrix}$$

so that

$$R_p^* A S_p = \begin{bmatrix} A_{pp} & & \\ & * & A_{qr} \\ & A_{rq} & A_{rr} \end{bmatrix}.$$

- ▶ DOFs p have been eliminated
- ▶ Interactions involving r are unchanged

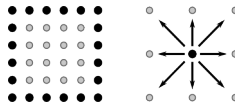
Interpolative decomposition

If $A_{:,q}$ is numerically low-rank, then there exist

- ▶ **skeleton** ($\hat{q}$) and **redundant** ($\check{q}$) columns partitioning $q = \hat{q} \cup \check{q}$
- ▶ an interpolation matrix T_q

such that

$$A_{:, \check{q}} \approx A_{:, \hat{q}} T_q.$$



- ▶ Essentially a pivoted QR written slightly differently
- ▶ Rank-revealing to any specified precision $\epsilon > 0$

Interactions between separated regions are low-rank.

Skeletonization

- ▶ Let $A = \begin{bmatrix} A_{pp} & A_{pq} \\ A_{qp} & A_{qq} \end{bmatrix}$ with A_{pq} and A_{qp} low-rank
- ▶ Apply ID to $\begin{bmatrix} A_{qp} \\ A_{pq}^* \end{bmatrix}$: $\begin{bmatrix} A_{q\check{p}} \\ A_{\check{p}q}^* \end{bmatrix} \approx \begin{bmatrix} A_{q\hat{p}} \\ A_{\hat{p}q}^* \end{bmatrix} T_p \implies \begin{aligned} A_{q\check{p}} &\approx A_{q\hat{p}} T_p \\ A_{\check{p}q} &\approx T_p^* A_{\hat{p}q} \end{aligned}$
- ▶ Reorder $A = \begin{bmatrix} A_{\check{p}\check{p}} & A_{\check{p}\hat{p}} & A_{\check{p}q} \\ A_{\hat{p}\check{p}} & A_{\hat{p}\hat{p}} & A_{\hat{p}q} \\ A_{q\check{p}} & A_{q\hat{p}} & A_{qq} \end{bmatrix}$, define $Q_p = \begin{bmatrix} I & & \\ -T_p & I & \\ & & I \end{bmatrix}$
- ▶ **Sparsify** via ID: $Q_p^* A Q_p \approx \begin{bmatrix} * & * & \\ * & A_{\hat{p}\hat{p}} & A_{\hat{p}q} \\ & A_{q\hat{p}} & A_{qq} \end{bmatrix}$
- ▶ Eliminate: $R_{\check{p}}^* Q_p^* A Q_p S_{\check{p}} \approx \begin{bmatrix} * & & \\ & * & A_{\hat{p}q} \\ & A_{q\hat{p}} & A_{qq} \end{bmatrix}$
- ▶ Reduces to a subsystem involving **skeletons** only

Integral equations

- ▶ Old algorithm (RS) in new **factorization** form
- ▶ New algorithm: HIF-IE

Algorithm: recursive skeletonization factorization

Build quadtree/octree.

for each level $\ell = 0, 1, 2, \dots, L$ from finest to coarsest **do**

Let C_ℓ be the set of all cells on level ℓ .

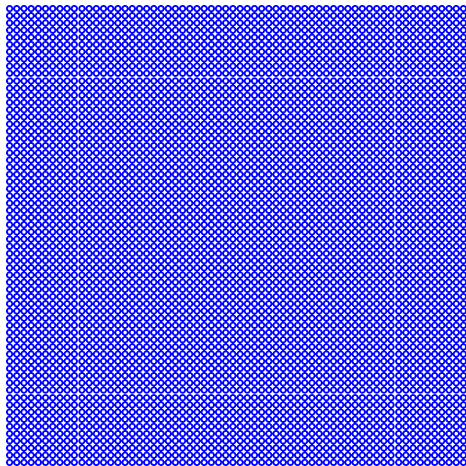
for each cell $c \in C_\ell$ **do**

Skeletonize remaining DOFs in c .

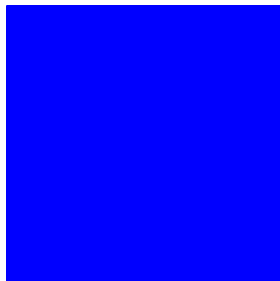
end for

end for

RSF in 2D: level 0

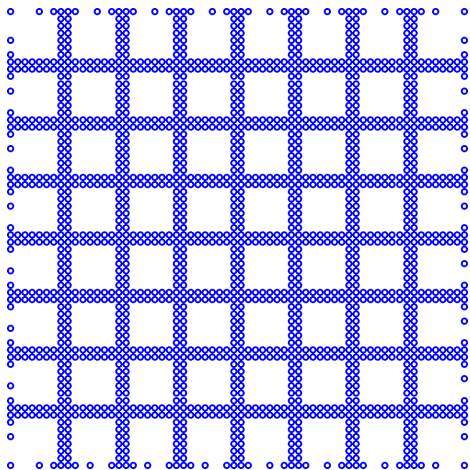


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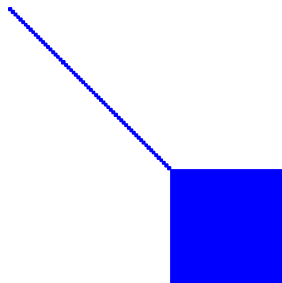


matrix

RSF in 2D: level 1

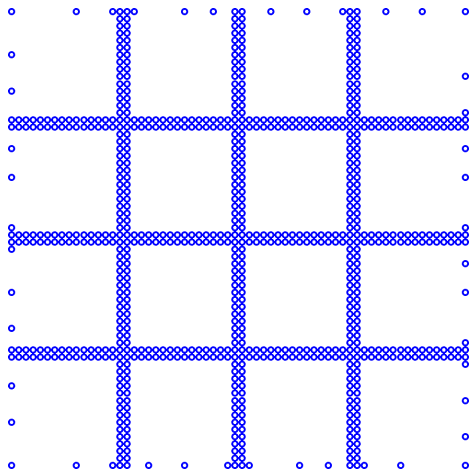


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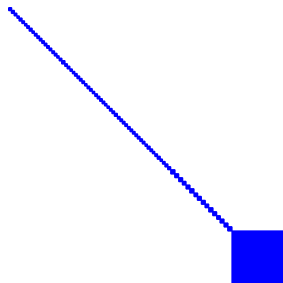


matrix

RSF in 2D: level 2

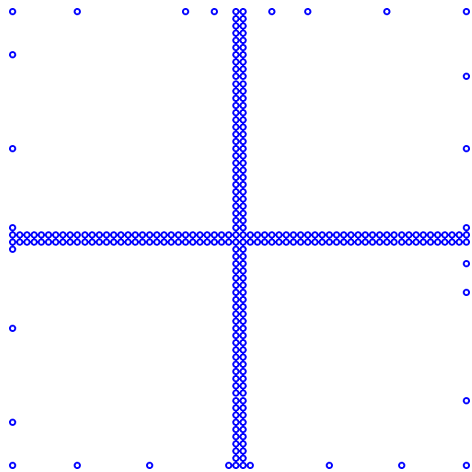


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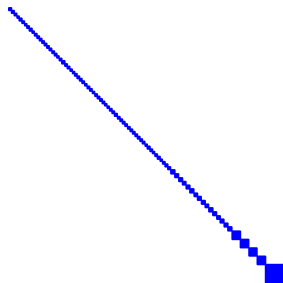


matrix

RSF in 2D: level 3

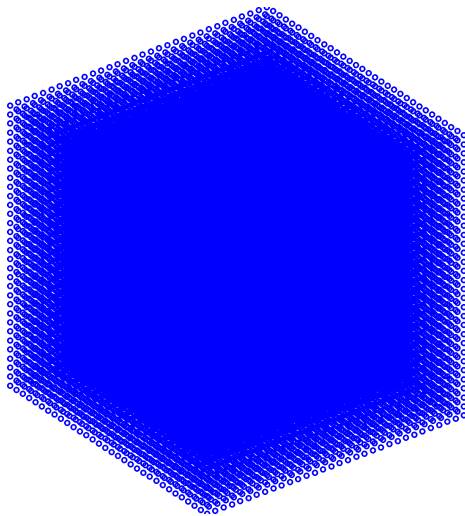


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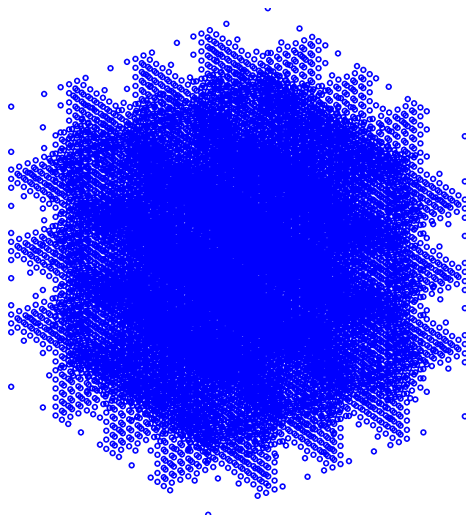
matrix

RSF in 3D: level 0



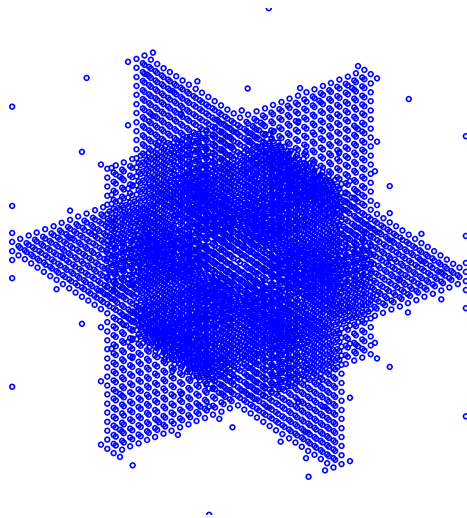
domain

RSF in 3D: level 1



domain

RSF in 3D: level 2



domain

- Skeletonization operators:

$$U_\ell = \prod_{c \in C_\ell} Q_c R_c, \quad V_\ell = \prod_{c \in C_\ell} Q_c S_c$$

$$Q_c = \begin{bmatrix} I & & \\ * & I & \\ & & I \end{bmatrix}, \quad R_c, S_c = \begin{bmatrix} I & * & \\ & I & \\ & & I \end{bmatrix}$$

- Block diagonalization:

$$D \approx U_{L-1}^* \cdots U_0^* A V_0 \cdots V_{L-1}$$

- Generalized LU decomposition:

$$A \approx U_0^{-*} \cdots U_{L-1}^{-*} D V_{L-1}^{-1} \cdots V_0^{-1}$$

$$A^{-1} \approx V_0 \cdots V_{L-1} D^{-1} U_L^* \cdots U_0^*$$

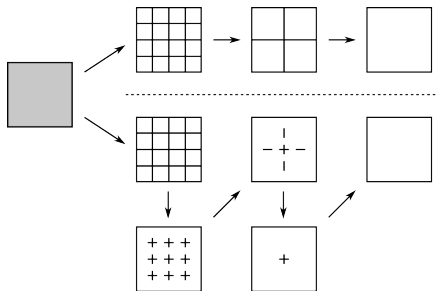
- Fast direct **solver** or preconditioner

RSF analysis

The cost is determined by the skeleton size.

	1D	2D	3D
Skeleton size	$O(\log N)$	$O(N^{1/2})$	$O(N^{2/3})$
Factorization cost	$O(N)$	$O(N^{3/2})$	$O(N^2)$
Solve cost	$O(N)$	$O(N \log N)$	$O(N^{4/3})$

Question: How to reduce the skeleton size in 2D and 3D?



Question: How to reduce the skeleton size in 2D and 3D?

- ▶ Skeletons cluster near cell interfaces (Green's theorem)
- ▶ Exploit skeleton geometry by further skeletonizing **along interfaces**
- ▶ Dimensional reduction

Algorithm: hierarchical interpolative factorization for IEs in 2D

Build quadtree.

for each level $\ell = 0, 1, 2, \dots, L$ from finest to coarsest **do**

Let C_ℓ be the set of all **cells** on level ℓ .

for each cell $c \in C_\ell$ **do**

Skeletonize remaining DOFs in c .

end for

Let $C_{\ell+1/2}$ be the set of all **edges** on level ℓ .

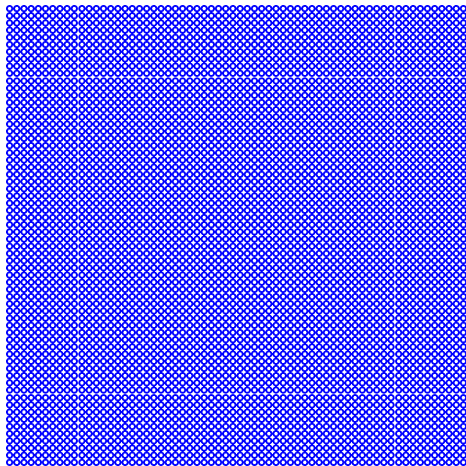
for each cell $c \in C_{\ell+1/2}$ **do**

Skeletonize remaining DOFs in c .

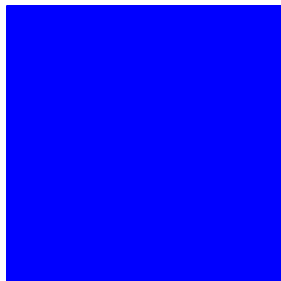
end for

end for

HIF-IE in 2D: level 0

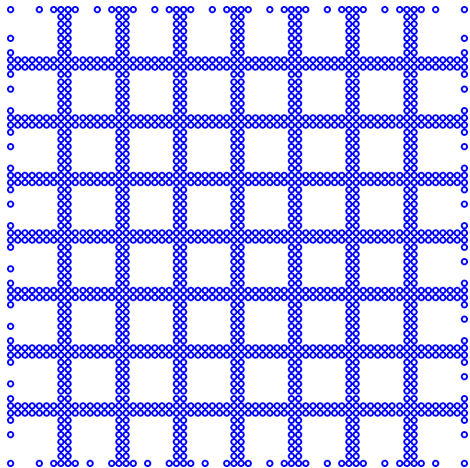


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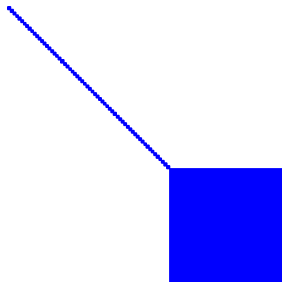


matrix

HIF-IE in 2D: level 1/2

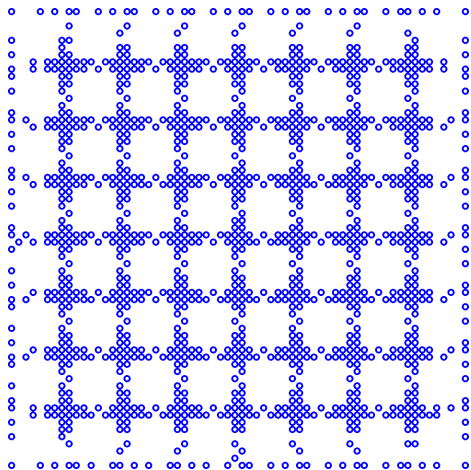


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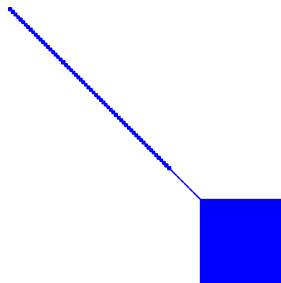


matrix

HIF-IE in 2D: level 1

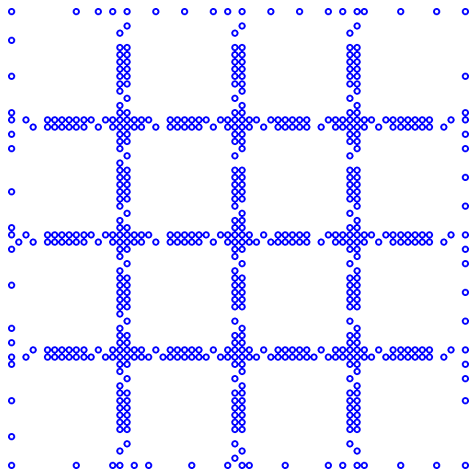


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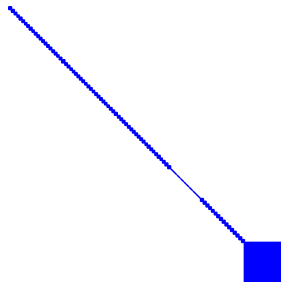


matrix

HIF-IE in 2D: level 3/2

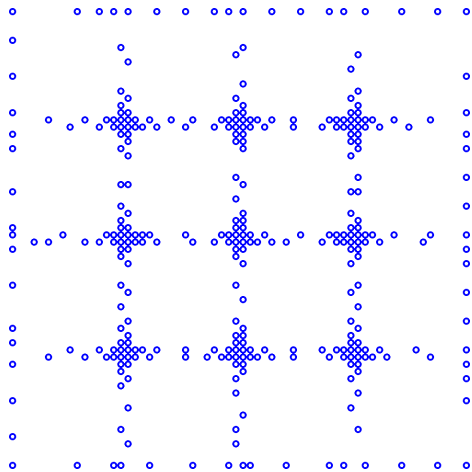


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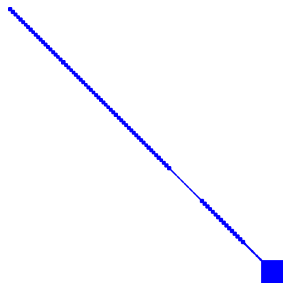


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HIF-IE in 2D: level 2

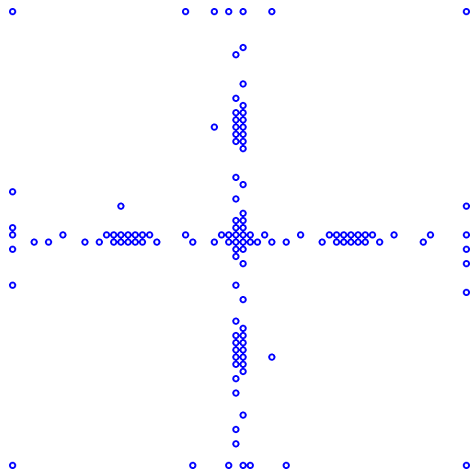


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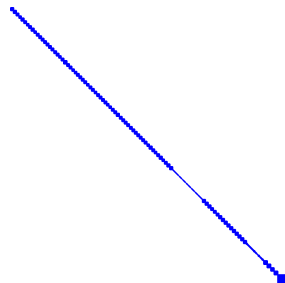


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HIF-IE in 2D: level 5/2

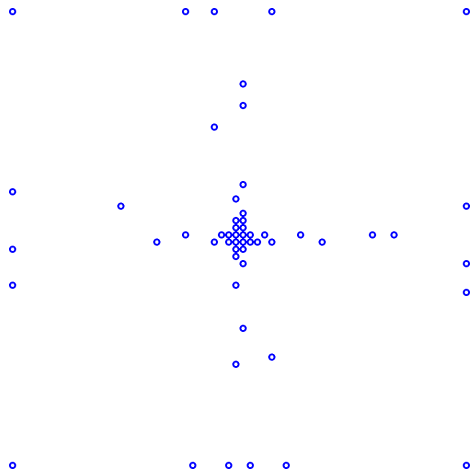


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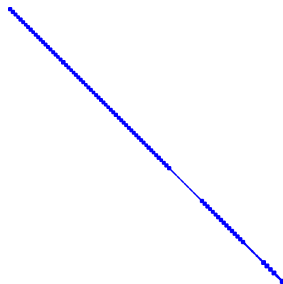


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HIF-IE in 2D: level 3

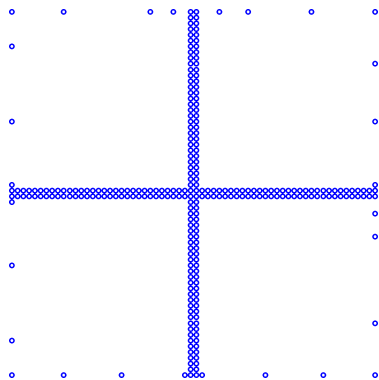


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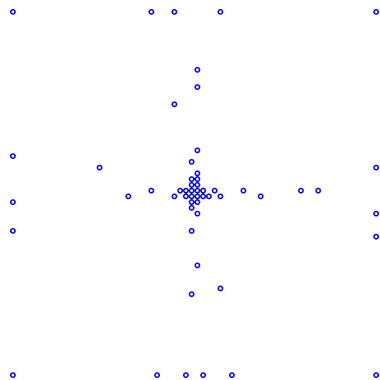


matrix

RSF vs. HIF-IE in 2D

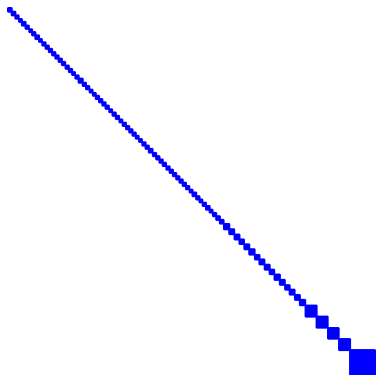


RSF

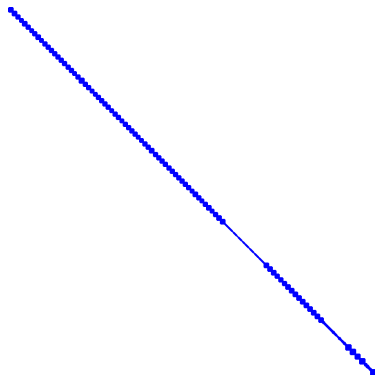


HIF-IE

RSF vs. HIF-IE in 2D



RSF



HIF-IE

Algorithm: hierarchical interpolative factorization for IEs in 3D

Build octree.

for each level $\ell = 0, 1, 2, \dots, L$ from finest to coarsest **do**

Let C_ℓ be the set of all **cells** on level ℓ .

for each cell $c \in C_\ell$ **do**

Skeletonize remaining DOFs in c .

end for

Let $C_{\ell+1/3}$ be the set of all **faces** on level ℓ .

for each cell $c \in C_{\ell+1/3}$ **do**

Skeletonize remaining DOFs in c .

end for

Let $C_{\ell+2/3}$ be the set of all **edges** on level ℓ .

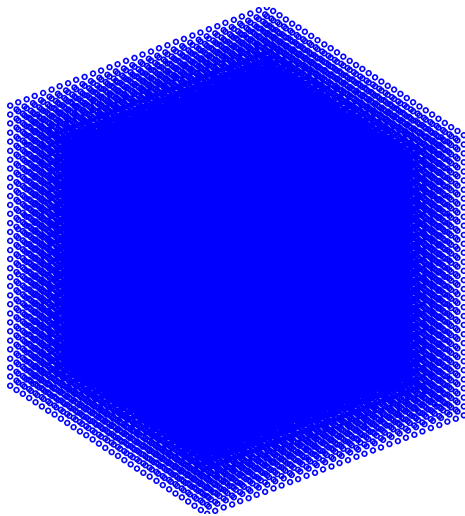
for each cell $c \in C_{\ell+2/3}$ **do**

Skeletonize remaining DOFs in c .

end for

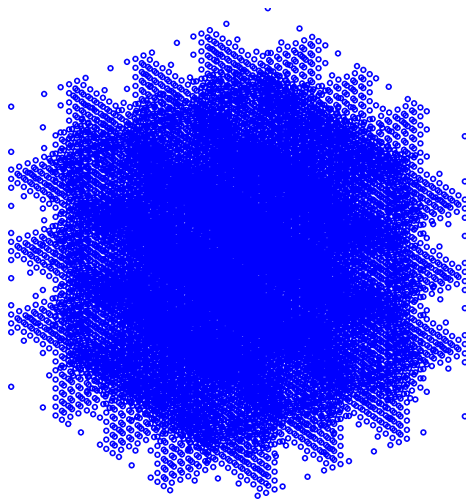
end for

HIF-IE in 3D: level 0



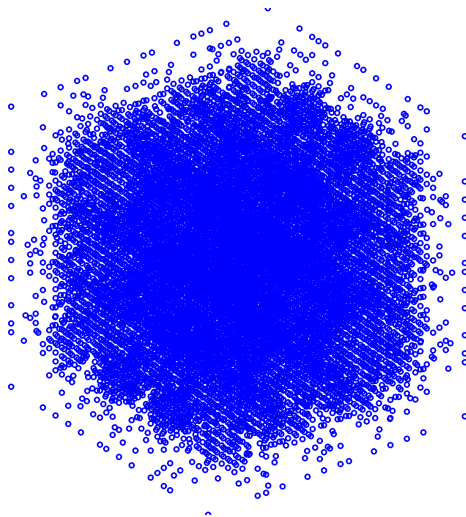
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HIF-IE in 3D: level 1/3



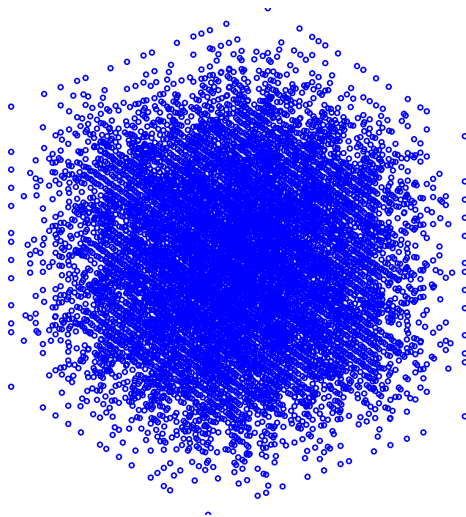
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HIF-IE in 3D: level 2/3



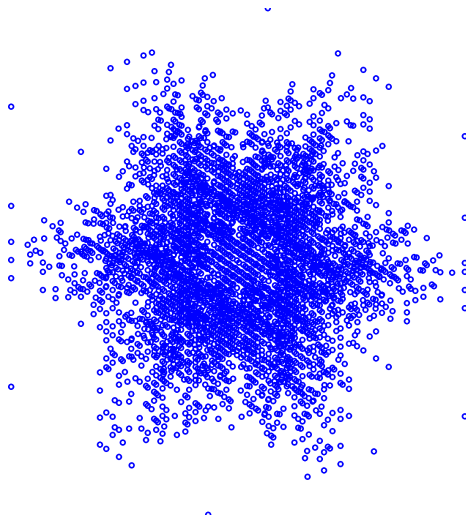
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HIF-IE in 3D: level 1



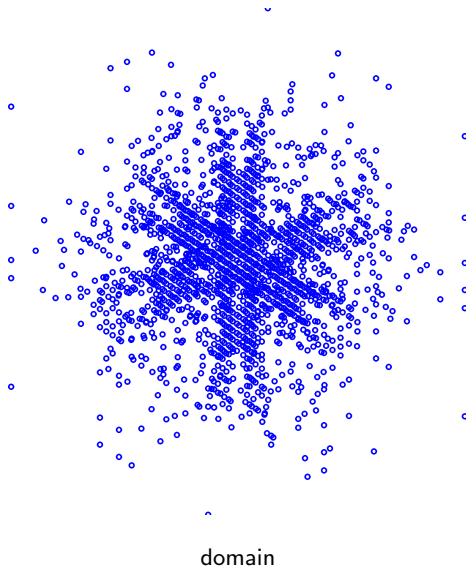
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HIF-IE in 3D: level 4/3

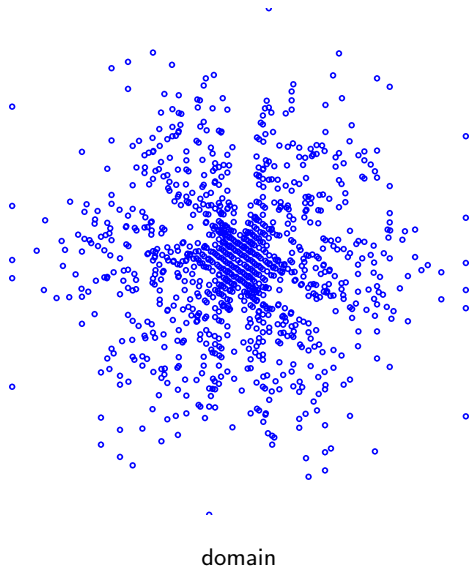


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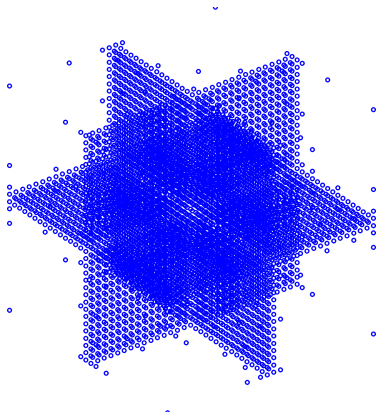
HIF-IE in 3D: level 5/3



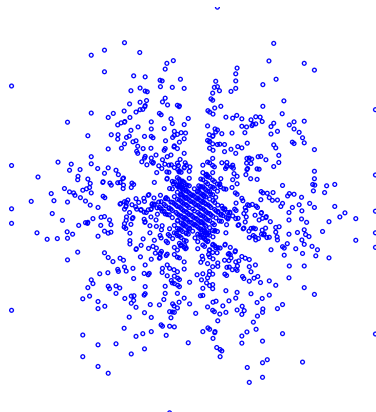
HIF-IE in 3D: level 2



RSF vs. HIF-IE in 3D



RSF



HIF-IE

► 2D:

$$A \approx U_0^{-*} U_{1/2}^{-*} \cdots U_{L-1/2}^{-*} D V_{L-1/2}^{-1} \cdots V_{1/2}^{-1} V_0^{-1}$$

$$A^{-1} \approx V_0 V_{1/2} \cdots V_{L-1/2} D^{-1} U_{L-1/2}^* \cdots U_{1/2}^* U_0^*$$

► 3D:

$$A \approx U_0^{-*} U_{1/3}^{-*} U_{2/3}^{-*} \cdots U_{L-1/3}^{-*} D V_{L-1/3}^{-1} \cdots V_{2/3}^{-1} V_{1/3}^{-1} V_0^{-1}$$

$$A^{-1} \approx V_0 V_{1/3} V_{2/3} \cdots V_{L-1/3} D^{-1} U_{L-1/3}^* \cdots U_{2/3}^* U_{1/3}^* U_0^*$$

Conjecture:

Skeleton size:	$O(\log N)$
Factorization cost:	$O(N)$
Solve cost:	$O(N)$

► 2D:

$$A \approx U_0^{-*} U_{1/2}^{-*} \cdots U_{L-1/2}^{-*} D V_{L-1/2}^{-1} \cdots V_{1/2}^{-1} V_0^{-1}$$

$$A^{-1} \approx V_0 V_{1/2} \cdots V_{L-1/2} D^{-1} U_{L-1/2}^* \cdots U_{1/2}^* U_0^*$$

► 3D:

$$A \approx U_0^{-*} U_{1/3}^{-*} U_{2/3}^{-*} \cdots U_{L-1/3}^{-*} D V_{L-1/3}^{-1} \cdots V_{2/3}^{-1} V_{1/3}^{-1} V_0^{-1}$$

$$A^{-1} \approx V_0 V_{1/3} V_{2/3} \cdots V_{L-1/3} D^{-1} U_{L-1/3}^* \cdots U_{2/3}^* U_{1/3}^* U_0^*$$

Conjecture:

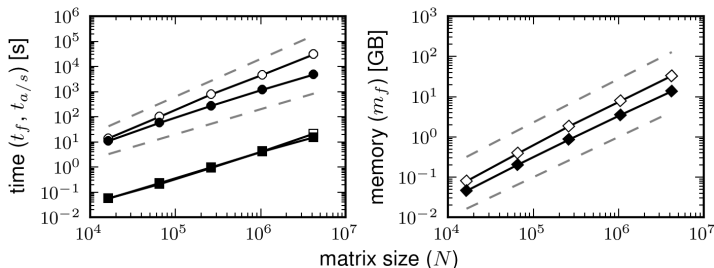
Skeleton size:	$O(\log N)$
Factorization cost:	$O(N)$
Solve cost:	$O(N)$

Actually slightly more complicated . . .

Numerical results in 2D

First-kind **volume** IE on the unit **square** with

$$a(x) \equiv 0, \quad K(x, y) = -\frac{1}{2\pi} \log \|x - y\|.$$

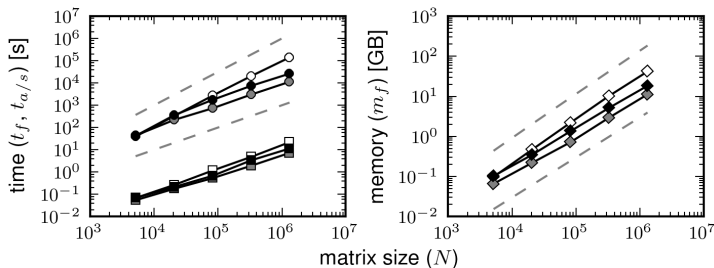


- ▶ **rskelf2** (white), **hifie2** (black)
- ▶ Factorization time (○), solve time (□), memory (◇) at precision $\epsilon = 10^{-6}$
- ▶ Reference scalings (gray dashes):
 - Left: $O(N)$ and $O(N^{3/2})$
 - Right: $O(N)$ and $O(N \log N)$

Numerical results in 3D

Second-kind **boundary** IE on the unit **sphere** with

$$a(x) \equiv -\frac{1}{2}, \quad K(x, y) = \frac{\partial}{\partial \nu(y)} \frac{1}{4\pi \|x - y\|}.$$

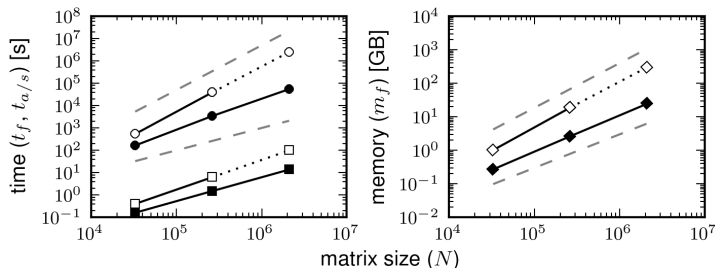


- ▶ **rskelf3** (white), **hifie3** (gray), **hifie3x** (black)
- ▶ Factorization time (○), solve time (□), memory (◇) at precision $\epsilon = 10^{-3}$
- ▶ Reference scalings (gray dashes):
 - Left: $O(N)$ and $O(N^{3/2})$
 - Right: $O(N)$ and $O(N \log N)$

Numerical results in 3D

First-kind **volume** IE on the unit **cube** with

$$a(x) \equiv 0, \quad K(x, y) = \frac{1}{4\pi \|x - y\|}.$$



- ▶ **rskelf3** (white), **hifief3** (black)
- ▶ Factorization time (○), solve time (□), memory (◇) at precision $\epsilon = 10^{-3}$
- ▶ Reference scalings (gray dashes):
 - Left: $O(N)$ and $O(N^2)$
 - Right: $O(N)$ and $O(N^{4/3})$

Differential equations

- ▶ Old algorithm (MF)
- ▶ New algorithm: HIF-DE
- ▶ Exploit **sparsity**: trivial “skeletonization”, thin separators

Algorithm: multifrontal

Build quadtree/octree.

for each level $\ell = 0, 1, 2, \dots, L$ from finest to coarsest **do**

Let C_ℓ be the set of all cells on level ℓ .

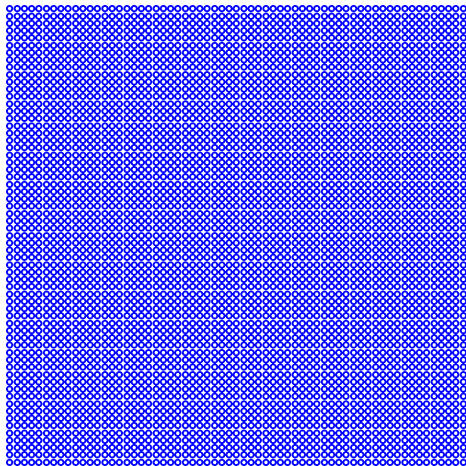
for each cell $c \in C_\ell$ **do**

Eliminate remaining interior DOFs in c .

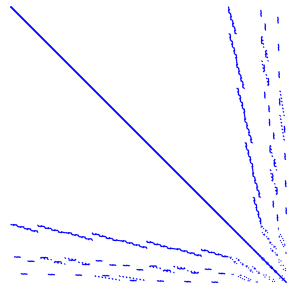
end for

end for

MF in 2D: level 0

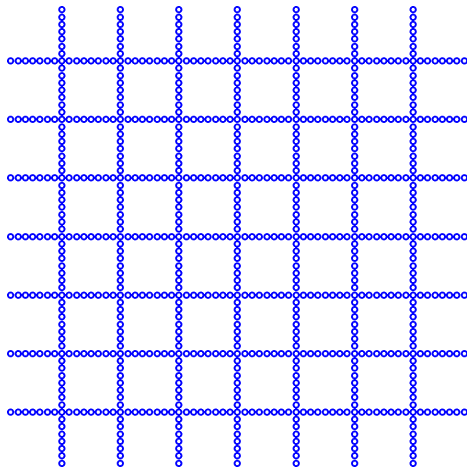


domain

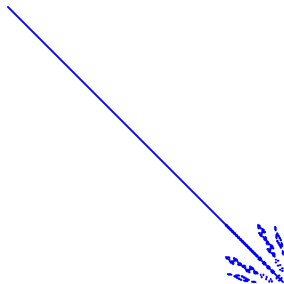


matrix

MF in 2D: level 1

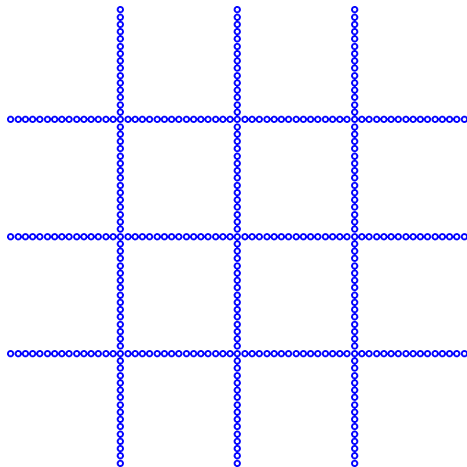


domain

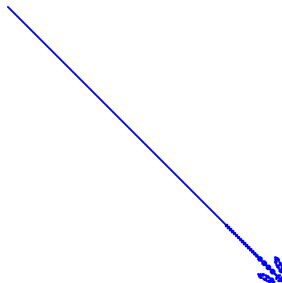


matrix

MF in 2D: level 2

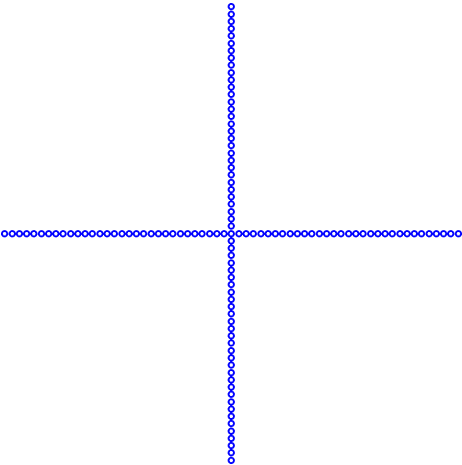


domain

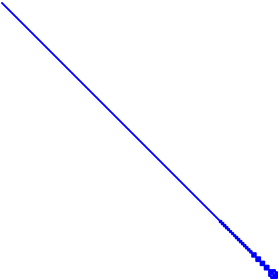


matrix

MF in 2D: level 3

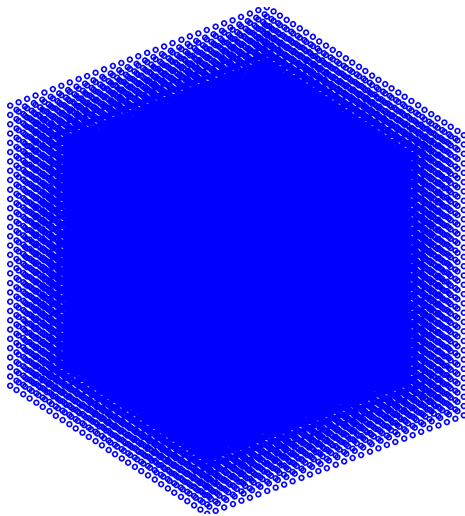


domain



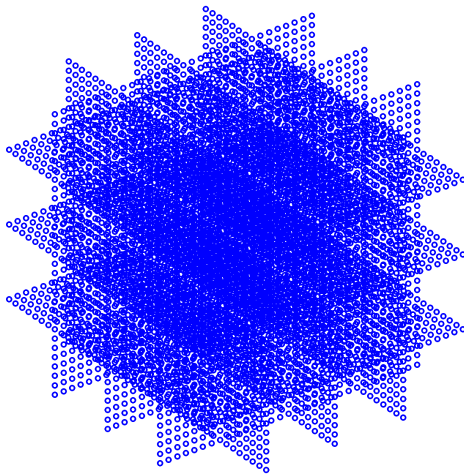
matrix

MF in 3D: level 0



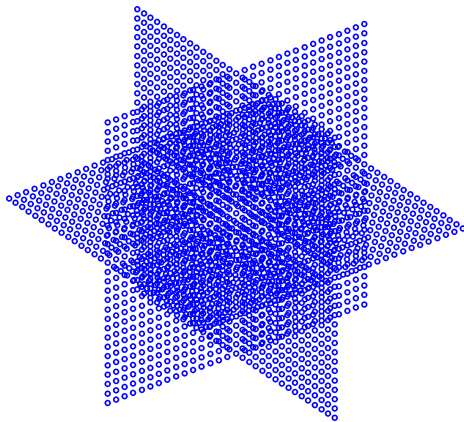
domain

MF in 3D: level 1



domain

MF in 3D: level 2



domain

Algorithm: hierarchical interpolative factorization for PDEs in 2D

Build quadtree.

for each level $\ell = 0, 1, 2, \dots, L$ from finest to coarsest **do**

Let C_ℓ be the set of all **cells** on level ℓ .

for each cell $c \in C_\ell$ **do**

Eliminate remaining interior DOFs in c .

end for

Let $C_{\ell+1/2}$ be the set of all **edges** on level ℓ .

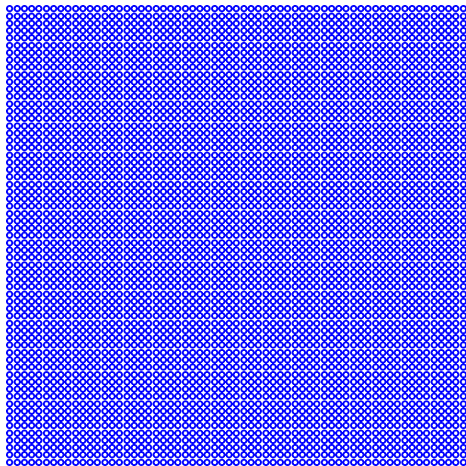
for each cell $c \in C_{\ell+1/2}$ **do**

Skeletonize remaining DOFs in c .

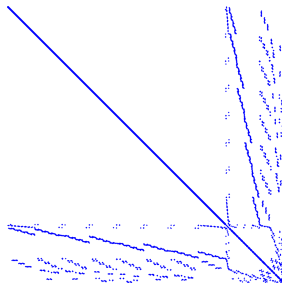
end for

end for

HIF-DE in 2D: level 0

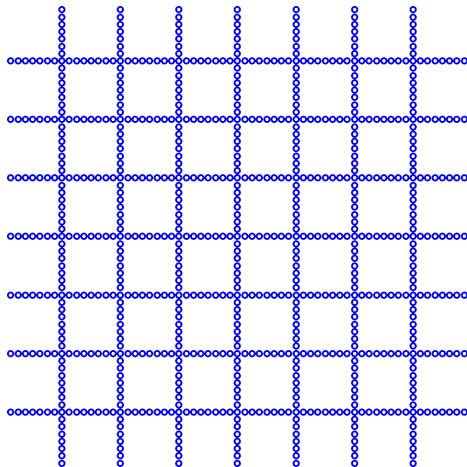


domain

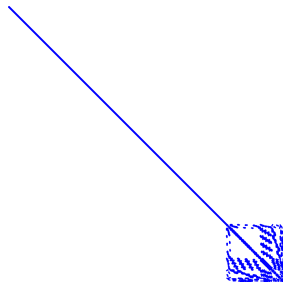


matrix

HIF-DE in 2D: level 1/2

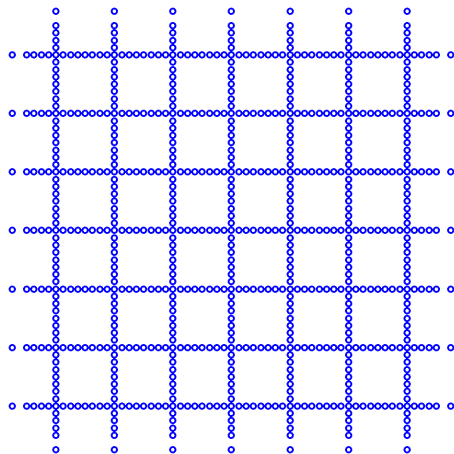


domain

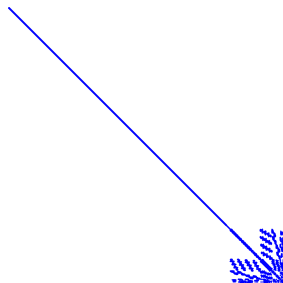


matrix

HIF-DE in 2D: level 1

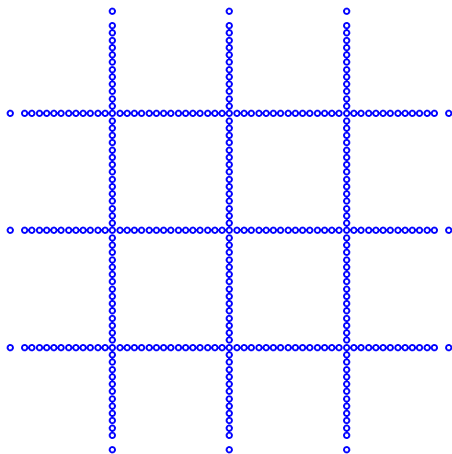


domain

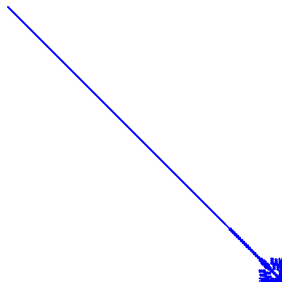


matrix

HIF-DE in 2D: level 3/2

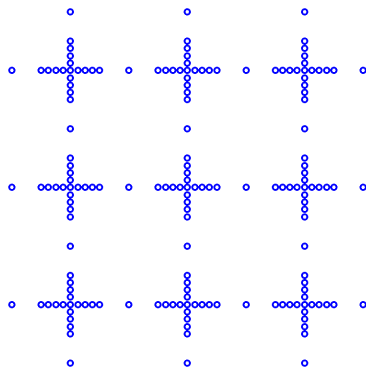


domain

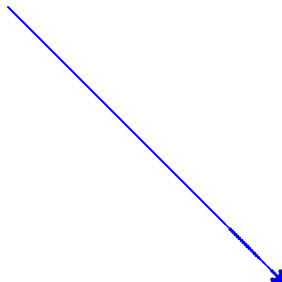


matrix

HIF-DE in 2D: level 2

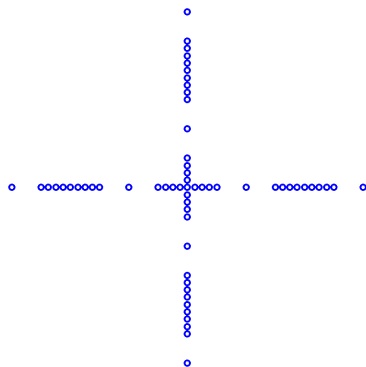


domain

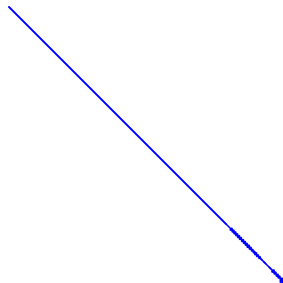


matrix

HIF-DE in 2D: level 5/2

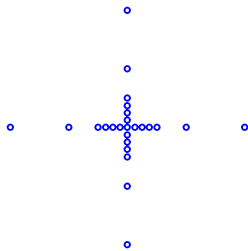


domain

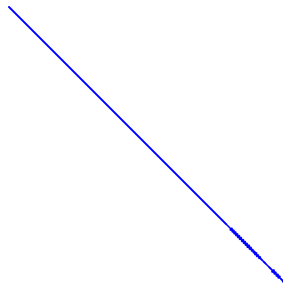


matrix

HIF-DE in 2D: level 3

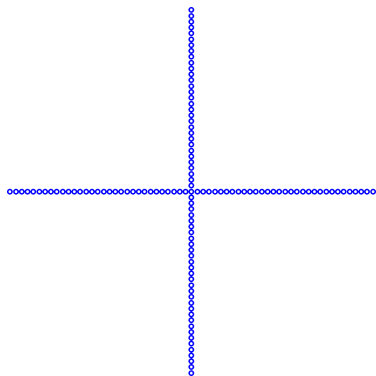


domain

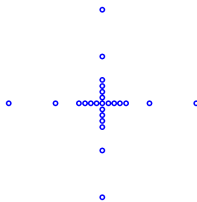


matrix

MF vs. HIF-DE in 2D

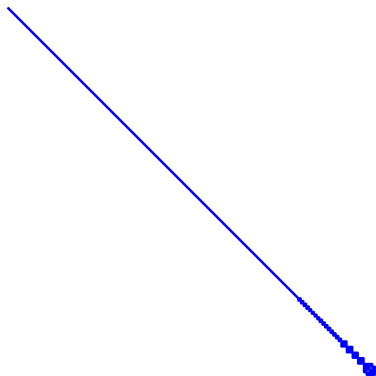


MF

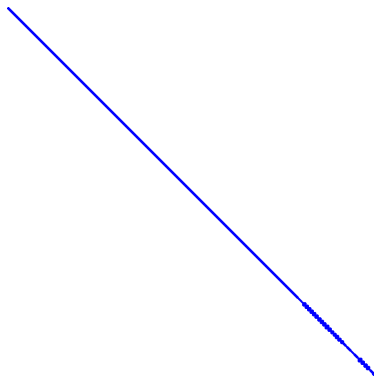


HIF-DE

MF vs. HIF-DE in 2D



MF



HIF-DE

Algorithm: hierarchical interpolative factorization for PDEs in 3D

Build octree.

for each level $\ell = 0, 1, 2, \dots, L$ from finest to coarsest **do**

Let C_ℓ be the set of all **cells** on level ℓ .

for each cell $c \in C_\ell$ **do**

Eliminate remaining interior DOFs in c .

end for

Let $C_{\ell+1/3}$ be the set of all **faces** on level ℓ .

for each cell $c \in C_{\ell+1/3}$ **do**

Skeletonize remaining DOFs in c .

end for

Let $C_{\ell+2/3}$ be the set of all **edges** on level ℓ .

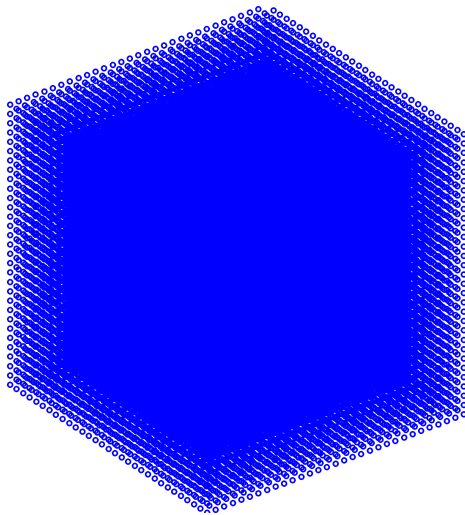
for each cell $c \in C_{\ell+2/3}$ **do**

Skeletonize remaining DOFs in c .

end for

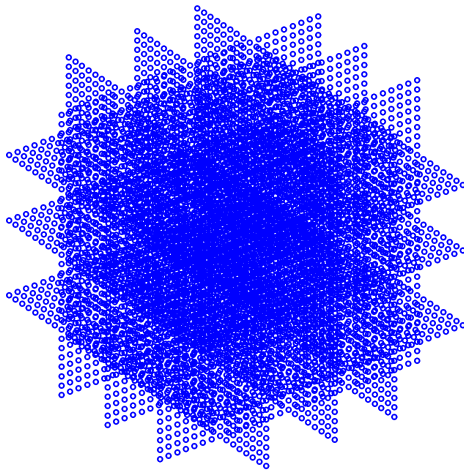
end for

HIF-DE in 3D: level 0



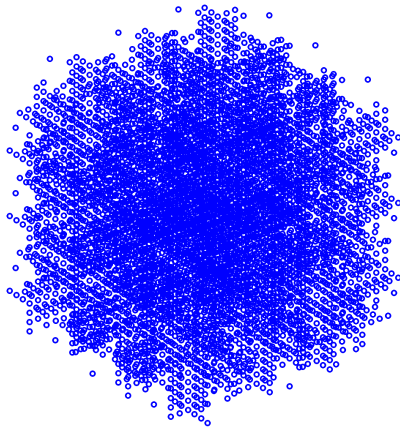
domain

HIF-DE in 3D: level 1/3



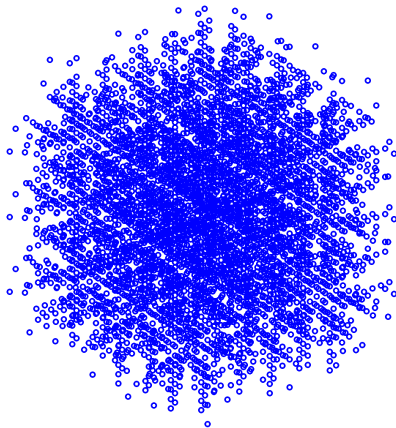
domain

HIF-DE in 3D: level 2/3



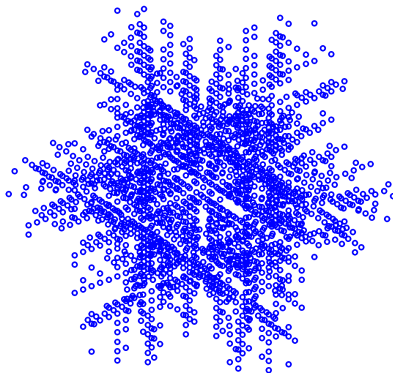
domain

HIF-DE in 3D: level 2



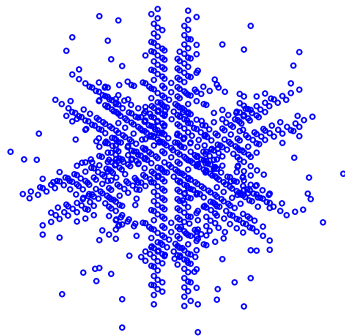
domain

HIF-DE in 3D: level 4/3



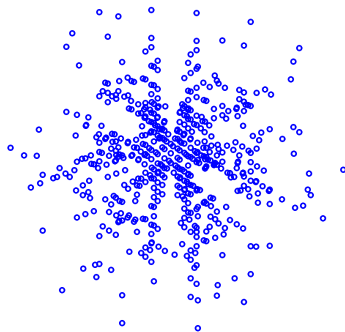
domain

HIF-DE in 3D: level 5/3



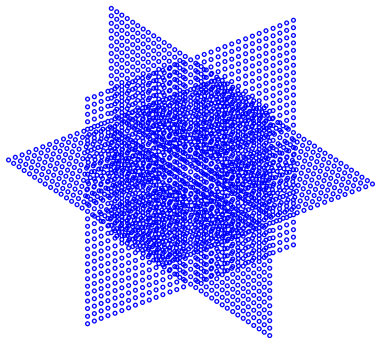
domain

HIF-DE in 3D: level 2

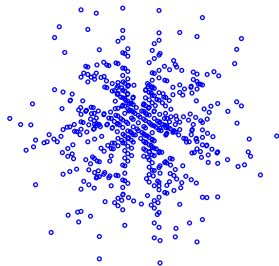


domain

MF vs. HIF-DE in 3D



MF

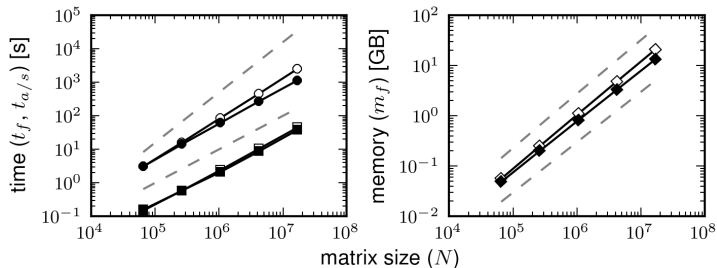


HIF-DE

Numerical results in 2D

Five-point stencil on the unit **square** with

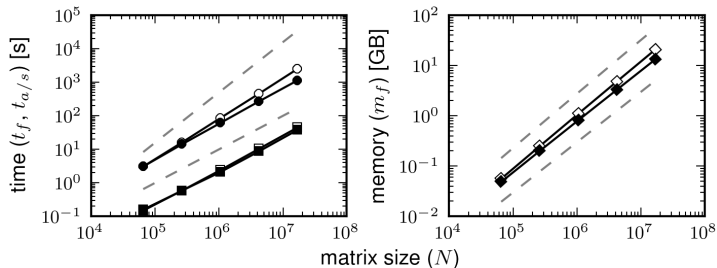
$$a(x) \equiv 1, \quad b(x) \equiv 0.$$



- ▶ **mf2** (white), **hifde2** (black)
- ▶ Factorization time ($\circ$), solve time ($\square$), memory ($\diamond$) at precision $\epsilon = 10^{-9}$
- ▶ Reference scalings (gray dashes):
 - Left: $O(N)$ and $O(N^{3/2})$
 - Right: $O(N)$ and $O(N \log N)$

Numerical results in 2D

Five-point stencil on the unit **square** with $a(x)$ a quantized high-contrast random field, $b(x) \equiv 0$.

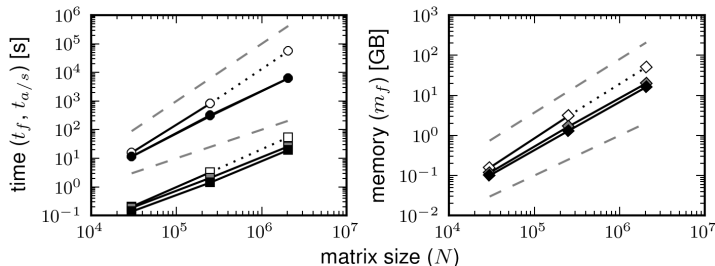


- ▶ **mf2** (white), **hifde2** (black)
- ▶ Factorization time ($\circ$), solve time ($\square$), memory ($\diamond$) at precision $\epsilon = 10^{-9}$
- ▶ Reference scalings (gray dashes):
 - Left: $O(N)$ and $O(N^{3/2})$
 - Right: $O(N)$ and $O(N \log N)$

Numerical results in 3D

Seven-point stencil discretization on the unit **cube** with

$$a(x) \equiv 1, \quad b(x) \equiv 0.$$



- ▶ **mf3** (white), **hifde3** (gray), **hifde3x** (black)
- ▶ Factorization time ($\circ$), solve time ($\square$), memory ($\diamond$) at precision $\epsilon = 10^{-6}$
- ▶ Reference scalings (gray dashes):
 - Left: $O(N)$ and $O(N^2)$
 - Right: $O(N)$ and $O(N^{4/3})$

Conclusions

- ▶ Efficient **factorization** of structured operators in 2D and 3D
 - Fast matrix-vector multiplication
 - Fast direct solver at high accuracy, preconditioner otherwise
 - Empirical **linear complexity** but no proof yet
- ▶ Sparsification and elimination (**skeletonization**) via the ID
- ▶ Dimensional reduction by alternating between cells, faces, and edges
- ▶ Can be viewed as adaptive numerical coarsening
- ▶ **Extensions:** $A^{1/2}$, $\log \det A$, $\text{diag } A^{-1}$
- ▶ Naturally parallelizable, block-sweep structure

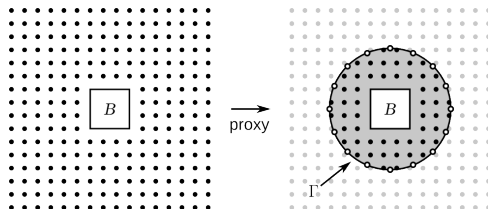
- ▶ *Perspective:* structured dense matrices can be sparsified very efficiently
- ▶ Can borrow directly from sparse algorithms, e.g., RSF = MF
- ▶ What other features of sparse matrices can be exploited?

MATLAB codes available at <https://github.com/klho/FLAM/>.

Additional slides

Proxy compression

- ▶ Main cost of algorithm is computing IDs of tall-and-skinny matrices
- ▶ **Global** operation can be reduced to **local** operation using Green's theorem
- ▶ Suffices to compress against neighbors plus “proxy” surface
- ▶ Crucial for beating $O(N^2)$ complexity



Second-kind IEs

- ▶ IEs of the form $u(x) + \int_{\Omega} K(x, y) u(y) d\Omega(y) = f(x)$
- ▶ **High contrast** in diagonal vs. off-diagonal entries
- ▶ Mixing of cell, face, edge in HIF-IE leads to error
- ▶ Need to use effective precision $O(\epsilon/N)$
- ▶ Quasilinear complexity estimates:

	2D	3D
Factorization cost	$O(N \log N)$	$O(N \log^6 N)$
Solve cost	$O(N \log \log N)$	$O(N \log^2 N)$