

# AMG Preconditioners based on Parallel Hybrid Coarsening and Bi-objective Matching



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### Overview

Solve  $Ax = b$

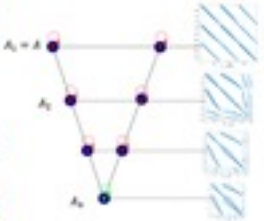
- $A$  is massive but sparse
- Need to use cluster/parallel computers
- Solve using Conjugate Gradient (CG) using some preconditioner  $B$ 
  - Solve  $B^{-1}Ax = B^{-1}b$

How to compute  $B$ ?

- $B$  is precomputed
- Use **Aggressive Multigrid Method (AMG)**

What is AMG?

- Coarse and Project  $k$
- Use **graphs** to build the Multigrid hierarchy



### A bi-objective matching framework

$$\max \sum w(e)x(e)$$

Subject to,  $x$  is a matching

$$\max \sum w(e)x(e) + \lambda \sum x(e)$$

Subject to,  $x$  is a matching

$x$  is a binary incident vector over edges

$\lambda = 0 \rightarrow$  max weight matching  
 $\lambda = \infty \rightarrow$  max cardinality matching

1. Overview

3. Bi-objective matching


### Experiments

- Metrics
  - Operator Complexity
    - Measure of the memory footprint of the multigrid and estimated cost of a V cycle
    - $\kappa_{PC} = \frac{\sum_i \max(A_i)}{\min(A_0)} > 1$
  - Number of Iterations
    - Number of iterations of the preconditioned CG solver
  - Setup time
    - Building time the preconditioner using bi-objective matching
  - Solving time
    - Total solution time of preconditioned CG

Smaller is better

### Coarsening

- How to do Coarsening?
  - From Matrix  $A$  to a graph  $G$
  - Coarse using successive **graph matching**
    - Matching set of non-overlapping edges in graph
  - Each level reduces the size of the matrix
- What would be a "good" coarsening?
  - Reduce the size of the matrix (ideally by **half** at every level)
  - The reduced matrix should be close to **diagonally dominant**



### From Optimal to Approximate matching

**Optimal Matching**

- Expensive
- Complicated
- No parallelism

**Approximate Matching**

- Fast, easy to implement and often parallel
- Groves algorithm (1/2 approximate)
  - Sort the edges from high to low
  - Construct a maximal matching in that order
- 1/2 -  $\epsilon$ 
  - Repeated short augmenting paths from random vertex

5. Empirical results

2. Coarsening

4. Approximate Matching

# Overview

## Solve

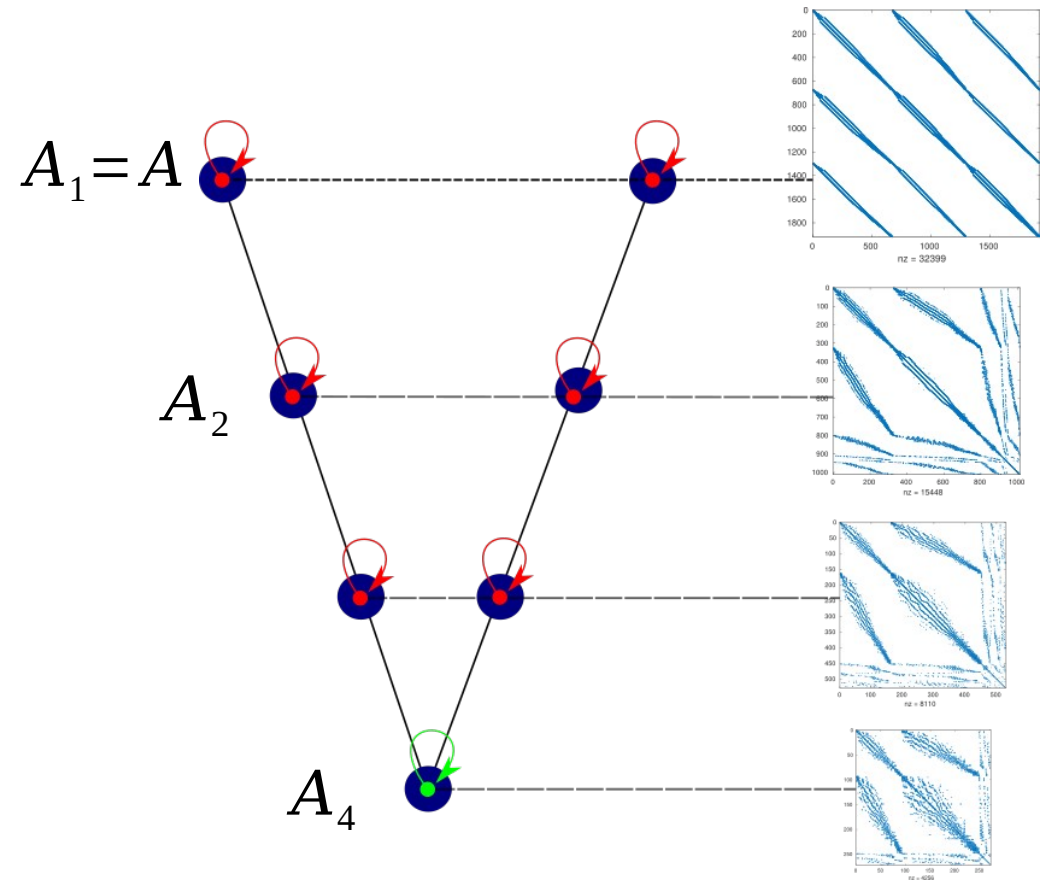
- is massive but sparse
- Need to use cluster/parallel computers
- Solve using Conjugate Gradient (CG) using some preconditioner .
  - Solve

## How to compute B?

- B is precomputed.
- Use **Algebraic Multigrid Method (AMG)**

## What is AMG?

- Coarse and Project A
- Use **graphs** to build the Multigrid Hierarchy.



# Contributions



Developed a bi-objective matching framework



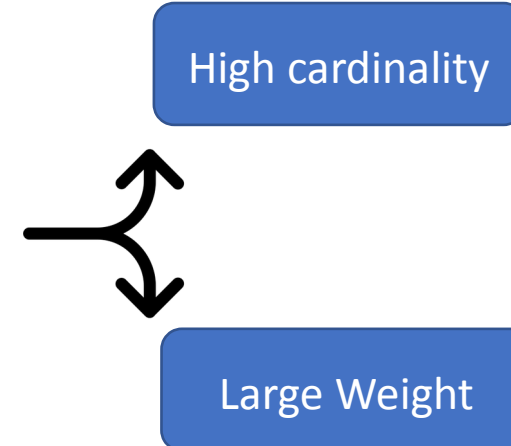
Employed the bi-objective matching to parallel coarsening in AMG



Experimented on solving anisotropic linear system in multiprocessor.

# Coarsening

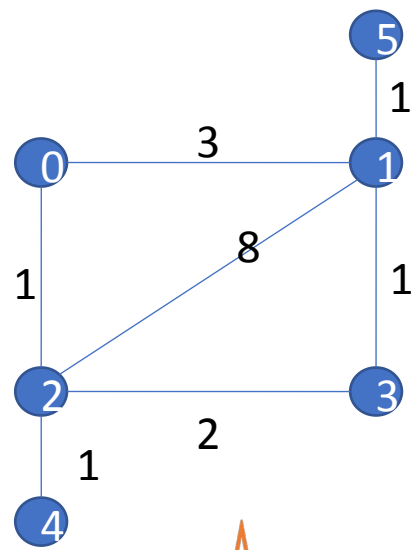
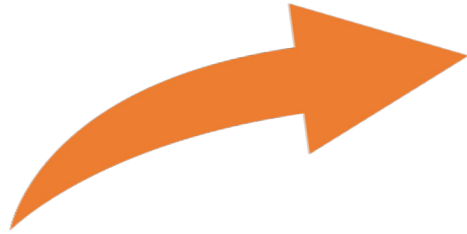
- How to do Coarsening?
  - From Matrix to a graph
  - Coarse using successive **graph matching**
    - Matching: set of non-overlapping edges in Graph.
  - Each level reduces the size of the matrix
- What would be a **“good”** coarsening?
  - Reduce the size of the matrix (ideally by **half** at every level)
  - The reduced matrix should be close to **diagonally dominant**



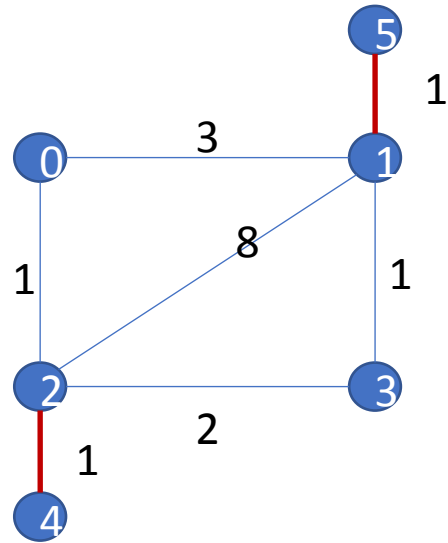
A matching is a set of non-overlapping edges.

$$\begin{pmatrix} 0 & 3 & 1 & 0 & 0 & 0 \\ 3 & 0 & 8 & 0 & 0 & 1 \\ 1 & 8 & 0 & 2 & 1 & 0 \\ 0 & 2 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 & 0 & 0 \end{pmatrix}$$

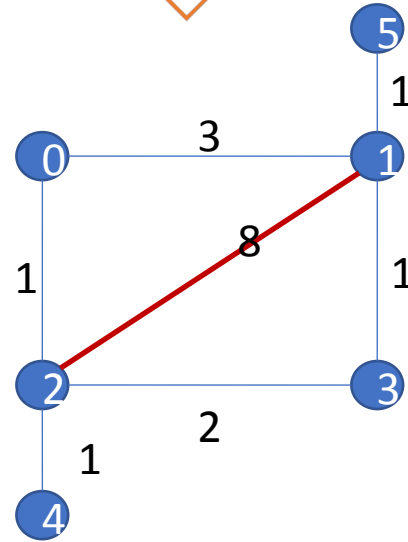
Matrix: A



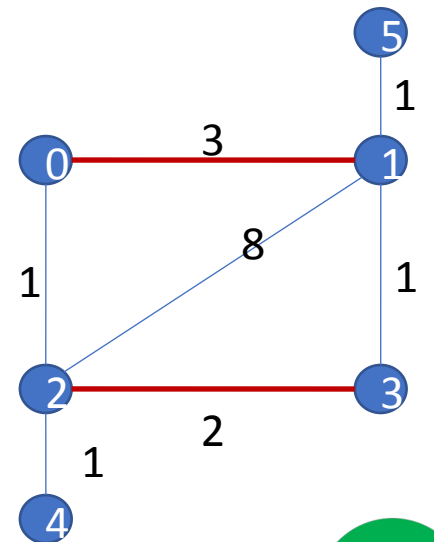
Graph: G



Card: 2  
Weight: 2



Card: 1  
Weight: 8



Card: 2  
Weight: 5



# A bi-objective matching framework

Subject to, is a  
matching

is a binary incident vector over edges

Subject to, is a  
matching

max weight matching  
max cardinality matching

# Properties of $k$ -matching

## Pareto Optimality

- Both **weights** and the **cardinality** are optimal for a parameter  $\lambda > 0$ .
- Need to solve matching optimally

$\lambda$  is parametric to the desired size of matching

- can be a function of max weight of the graph
- set such that matching has certain cardinality guarantee

Subject to, is a matching

$$\lambda = \max \left\{ \frac{k-1}{2} \gamma - \frac{k+1}{2} \delta, \epsilon \right\}$$

Max weight matching where there is **no Augmenting path** of length  $k$ .  
max weight, min weight



# From Optimal to Approximate matching

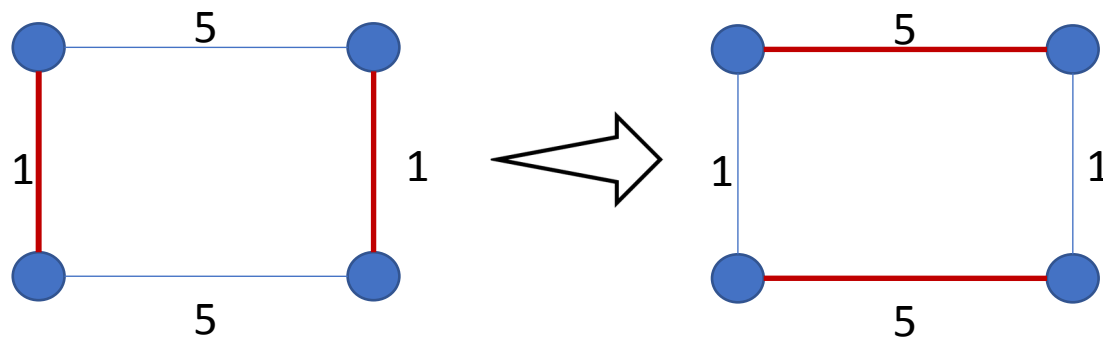
## Optimal Matching

- Expensive
- Complicated
- No parallelism

## Approximate Matching

- Fast, easy to implement and often parallel.
- Greedy Algorithm (1/2-approximate)
  - Sort the edges from high to low
  - construct a maximal matching in that order
- Repeated short augmenting paths from random vertex

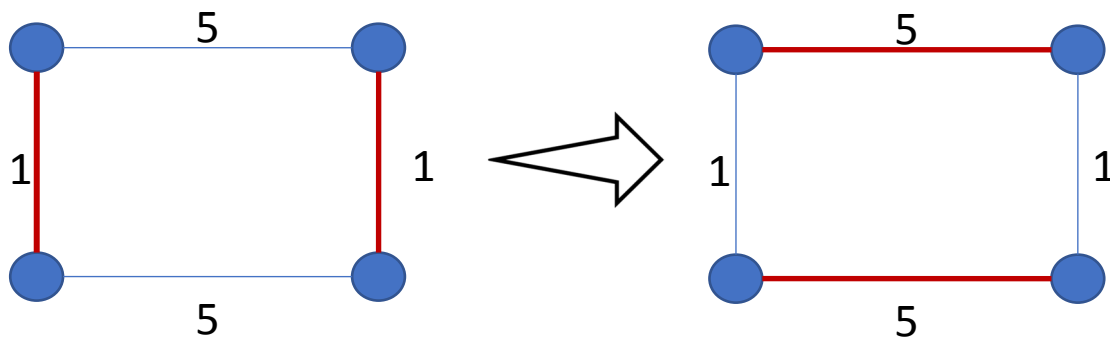
# The Randomized 2/3-approximate Algorithm\*



A 2-augmentation

\*Sanders, Peter, and Seth Pettie. "A simpler linear time 2/3-epsilon approximation for maximum weight matching." (2004).

# The Randomized 2/3-approximate Algorithm\*



A 2-augmentation

## 2-augmentation P

- With respect to a matching  $M$
- Weight increasing alternating path
- Number of edges in

\*Sanders, Peter, and Seth Pettie. "A simpler linear time 2/3-epsilon approximation for maximum weight matching." (2004).

# The Random Order Augmentation Matching Algorithm ROMA\*

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**ROMA**( $G = (V, E)$ ,  $w: E \rightarrow \mathbb{R}^{\geq 0}$ ,  $\text{int } \ell$ )

1  $M := \emptyset$  (or initialise  $M$  with any matching)

2 **for**  $i := 1$  **to**  $\ell$  **do**

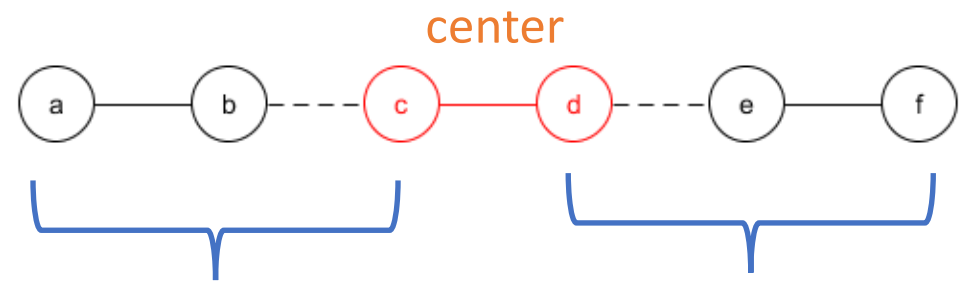
3     **for each** node  $v \in V$  in random order **do**

4          $M := M \oplus \text{aug}(v)$

5 **return**  $M$

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: A 2-augmentation centered at  $v$ .



Can also be implemented in a shared memory parallel machine.\*\*

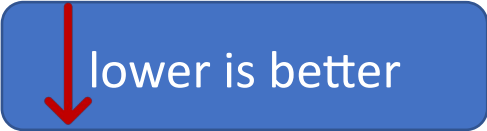
\*J. Maue and P. Sanders, "Engineering Algorithms for Approximate Weighted Matching," in *Experimental Algorithms*, 2007

\*\*A. Berge, "A parallel version of the Random Order Augmentation Matching Algorithm," Master's thesis, University of Bergen, 2020

# Experiments

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- Metrics
  - Operator Complexity
    - Measure of the memory footprint of the multigrid and estimated cost of a V-cycle
  - Setup time
    - Build time the preconditioner using bi-objective matching
  - Number of iterations
    - Number of iterations of the preconditioned CG solver
  - Solving time
    - Total solution time of preconditioned CG



↓ lower is better

# Problems

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- Poisson Benchmark with Axial anisotropy in 2D and 3D
- The boundary value problem

Here,

# Machine

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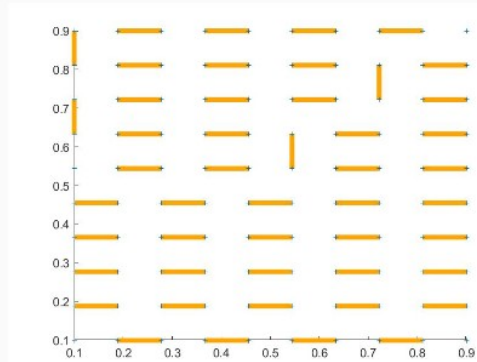
- CINECA Marconi 100
  - **Nodes:** 980
  - **Processors:** 2x16 cores IBM POWER9 AC922 at 3.1 GHz
  - **Accelerators:** 4 x NVIDIA Volta V100 GPUs, Nvlink 2.0, 16GB
  - **RAM:** 256 GB/node

System	Core	Rmax (PFlop/s)
1. Frontiers	8,730,112	1,102.00
2. Fugaku	7,630,848	442.01
3. Lumi	2,220,288	309.10
24. Marconi	347,776	21.64

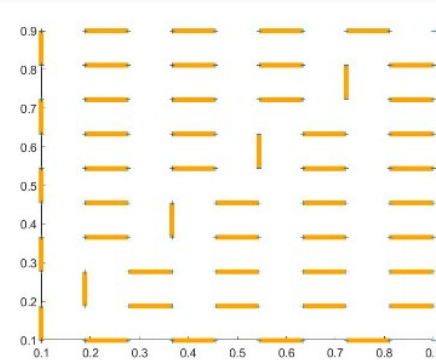
# Effect of

$k_1 \gg k_2$

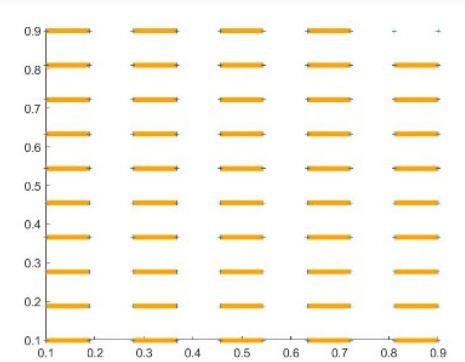
$\lambda$	opc	cr	nit
0	1.552	1.923	9
1.25	1.559	1.961	10
1.75	1.489	1.961	7



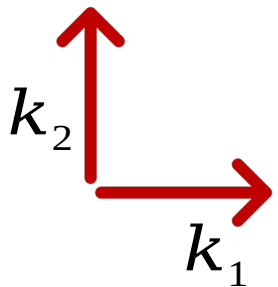
$\lambda = 0, k_1 \gg k_2$



$\lambda = 1.25, k_1 \gg k_2$



$\lambda = 1.75, k_1 \gg k_2$



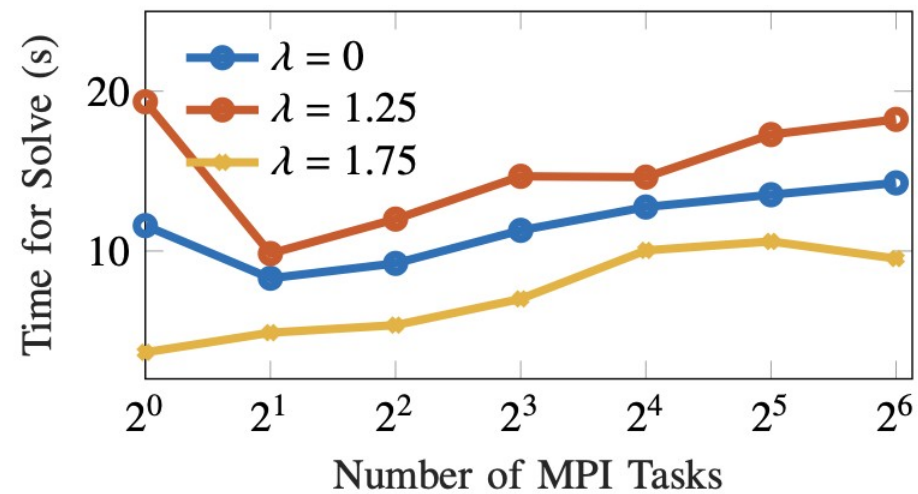
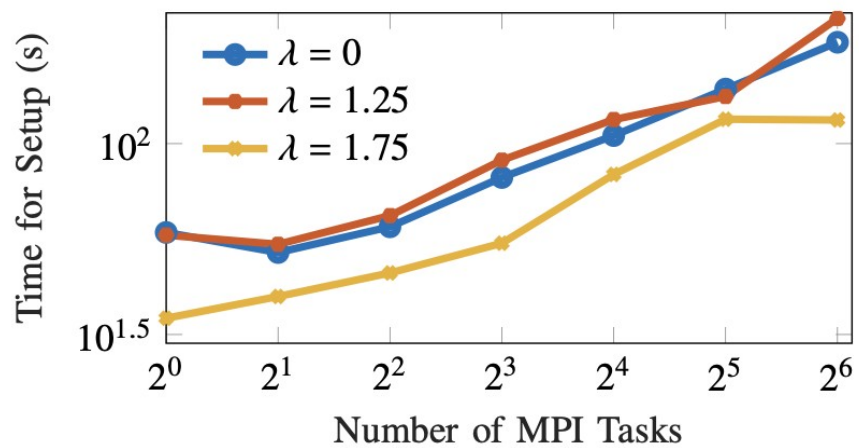
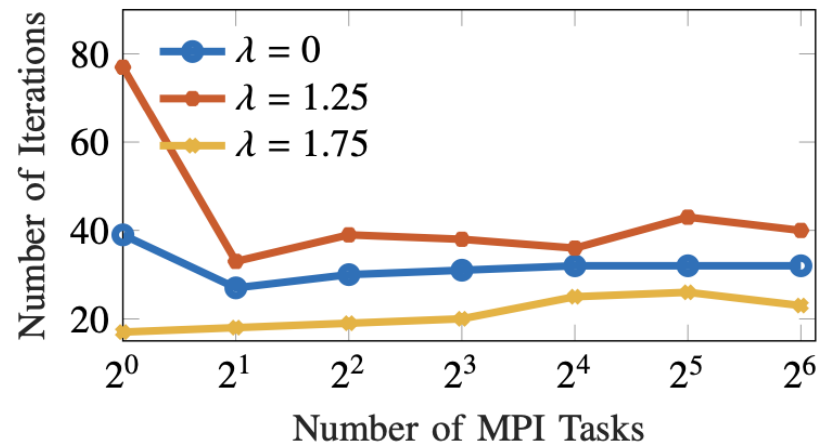
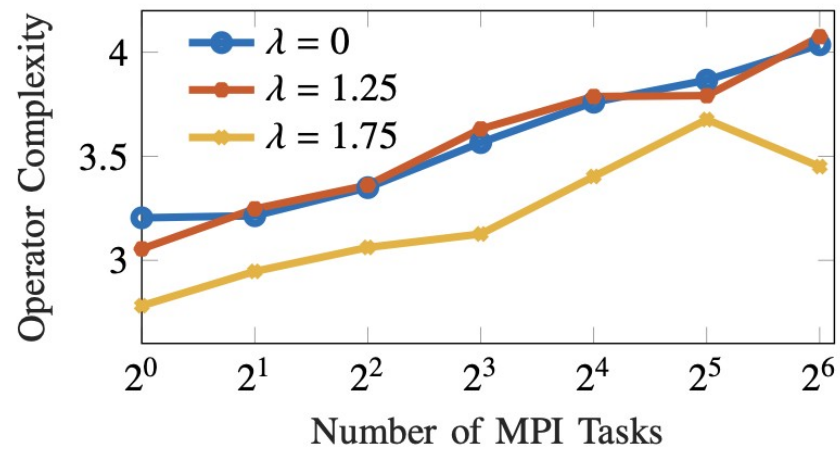


# Weak Scaling Analysis

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## Weak Scaling Setup

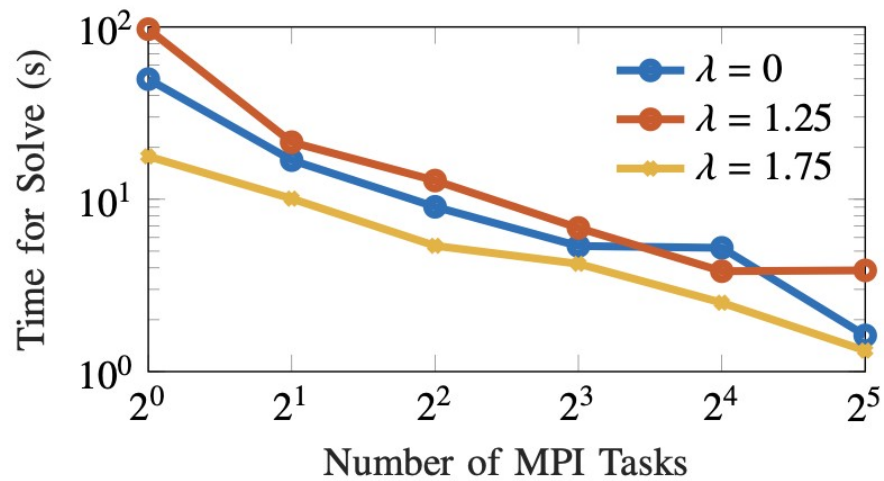
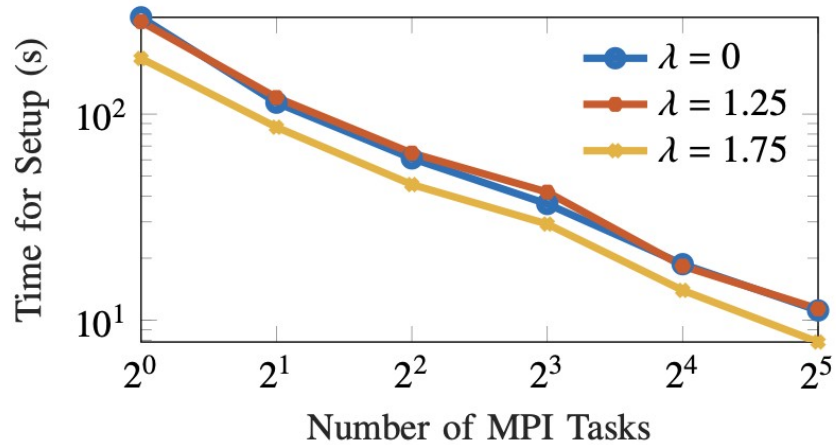
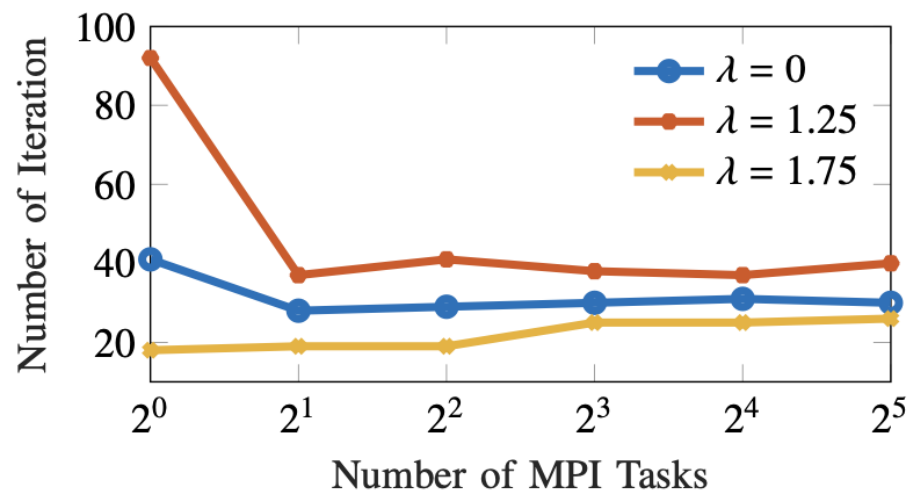
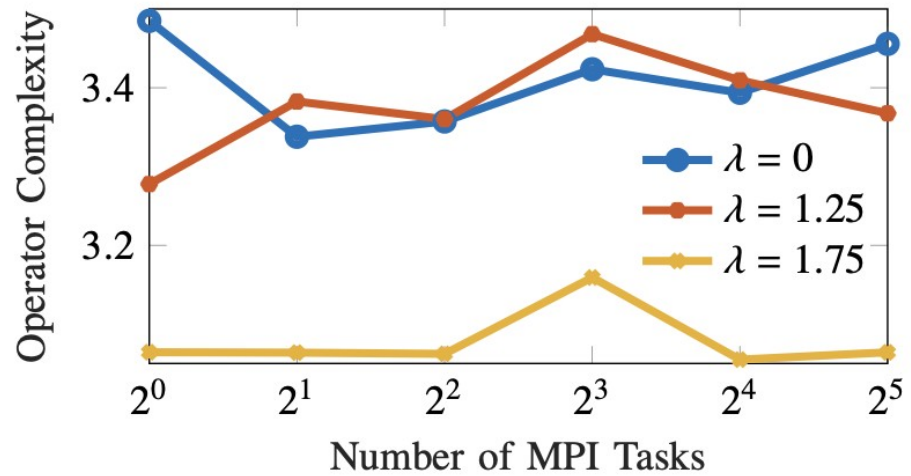
- MPI tasks
- 16 threads per task
- dofs per task
- cores with dofs for the largest problem



# Strong Scaling Analysis

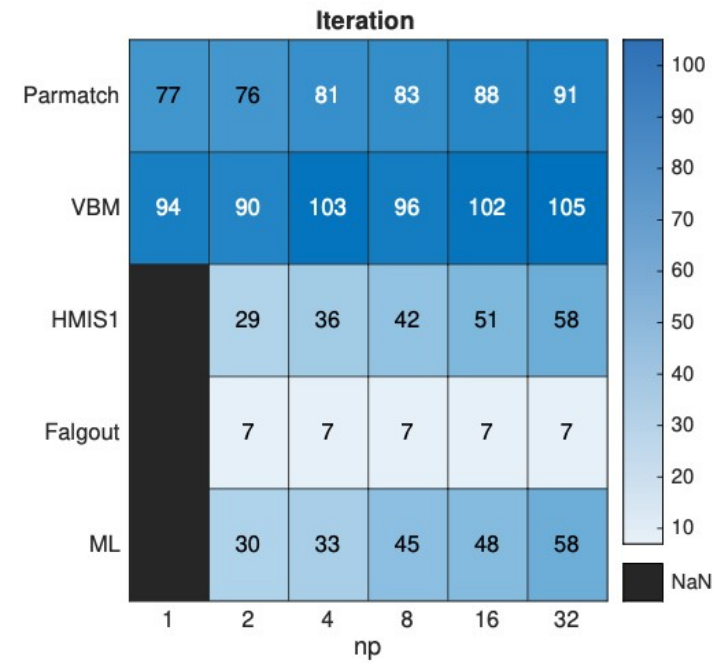
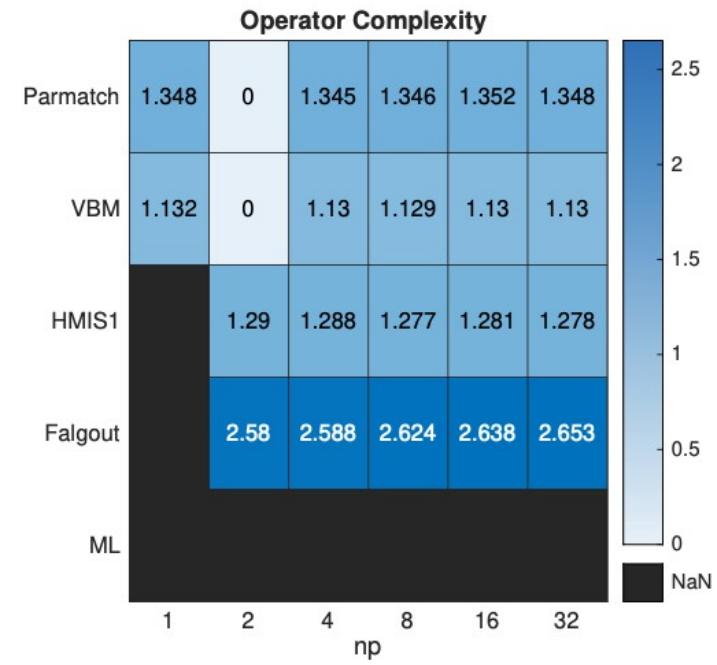
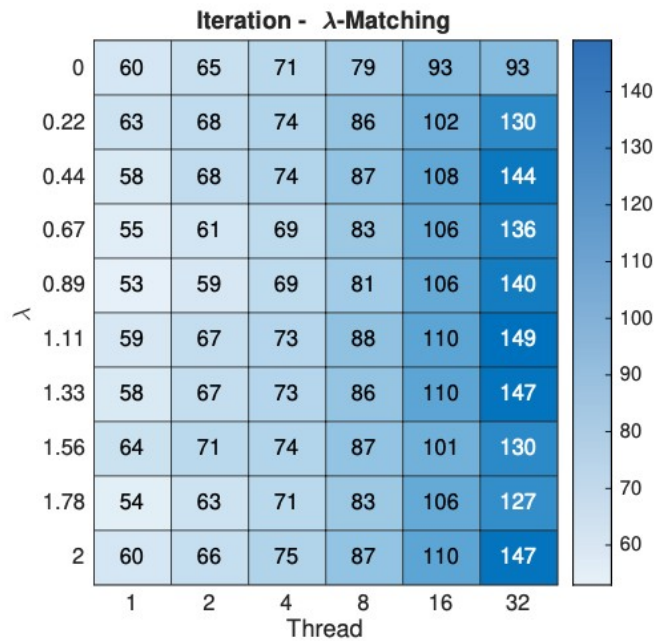
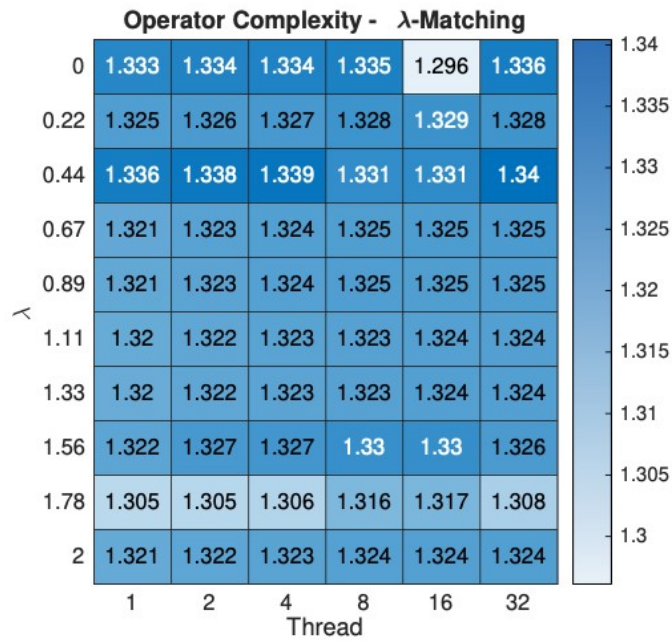
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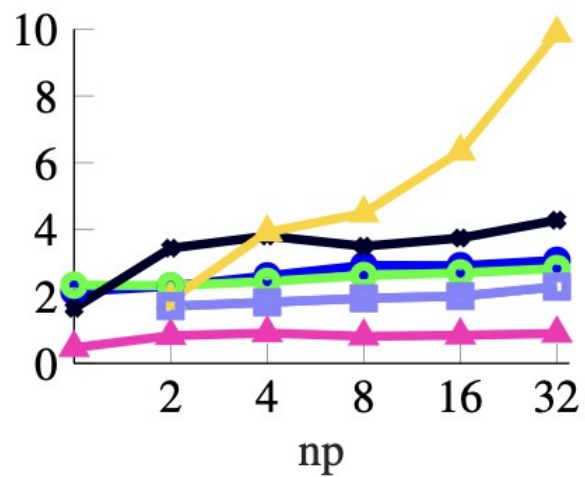
- Setup
  - Number of dofs
  - MPI processes
  - Each task with 16 threads



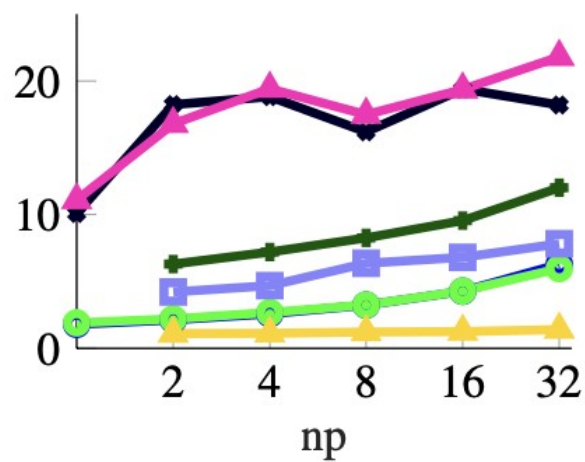
# Comparison with other algorithms

- Benchmark Algorithms
  - Bi-objective Matching (our)
  - Parmatch (greedy matching as aggregator)
  - VBM
  - Falgout
  - HMIS1
  - ML

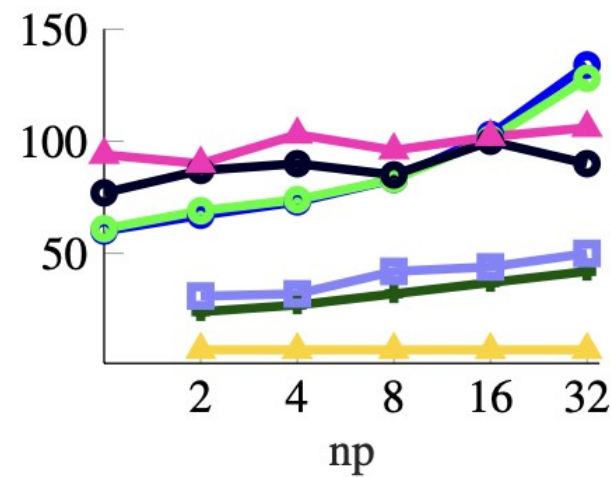




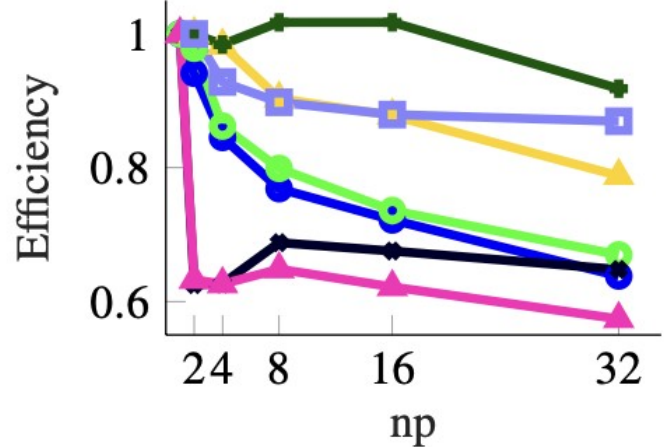
(a) Build time (s).



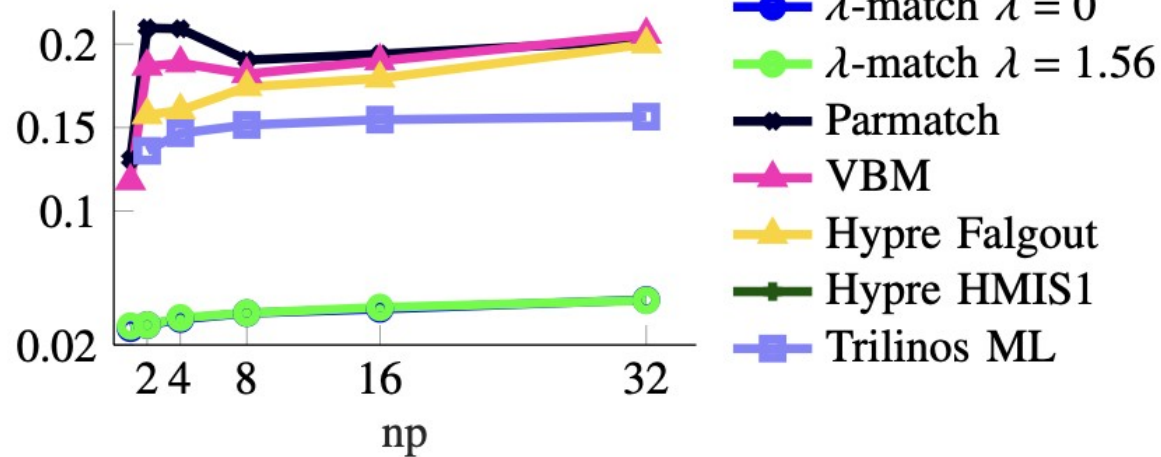
(b) Solve time (s).



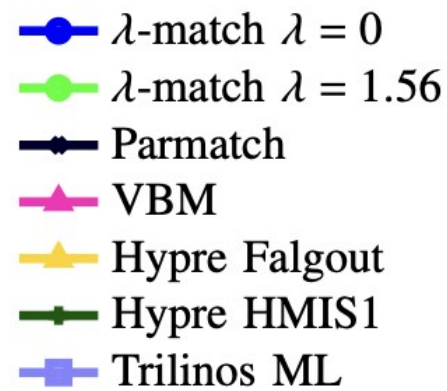
(c) Iteration number.



(d) Time per iteration (Efficiency)



(e) Time per iteration (s).



# Thank You!



<https://smferdous1.github.io/>



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