Enumeration of Molecular Clusters

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Introduction

- Terminologies
 - Floppy cluster
 - Rigid cluster
- Why enumerate?
 - Scientific importance
- How to enumerate?
 - Start by breaking a bond

Interesting question

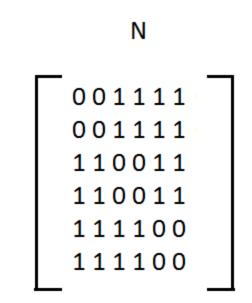
- How many rigid clusters are there for a particular n (n: number of particles)?
- The number of rigid clusters increases dramatically

up to n=5, there is one

- n=6, there are 2 n=7, there are 5
- n=8, there are 13 n=9, there are 52
- n=10, there are 263 n=11, there are 1657, etc

Representation

• Rigid cluster, N = 6



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Algorithms

- Algorithm 1
 - Floppy to rigid
- Algorithm 2
 - One rigid cluster to all possible enumerations

Algorithm 1(Floppy to Rigid)

- Start with a particle and a matrix M of bond constraints
- Compute the Null space of M
- Take a step in the direction of the vector in the tangent space
- Get back to manifold by solving the nonlinear system of equations by Newton's method
- Repeat until new bond is formed

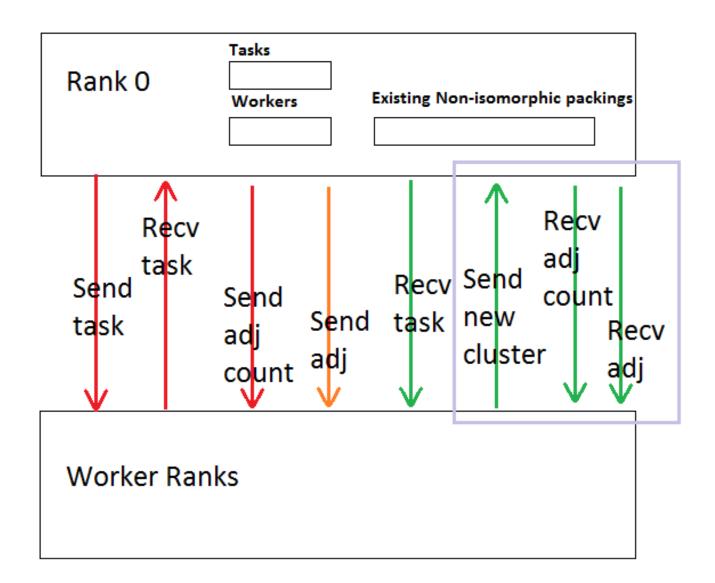
Algorithm 2(Permute and find all packings)

- Break a bond, then you get a floppy particle(s)
- Use Algorithm 1 to find a rigid cluster
- The new rigid cluster might be different
- You have to do this for all bonds from all rigid clusters

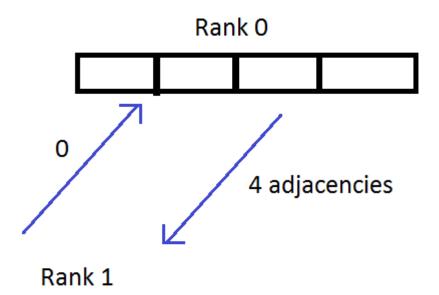
Parallelization

- You have to break all the bonds from all rigid clusters
- You can do it independently
- This operation is parallelizable!

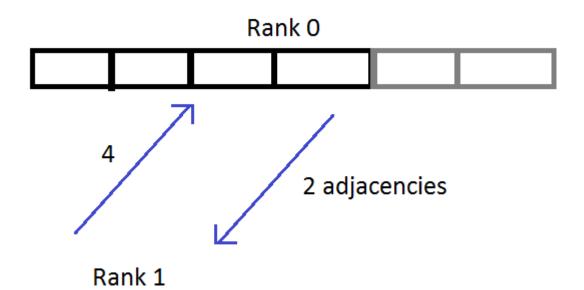
Process Communication



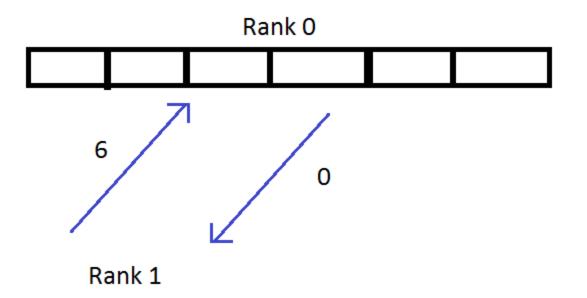
Isomorphism Check



Isomorphism Check



Isomorphism Check



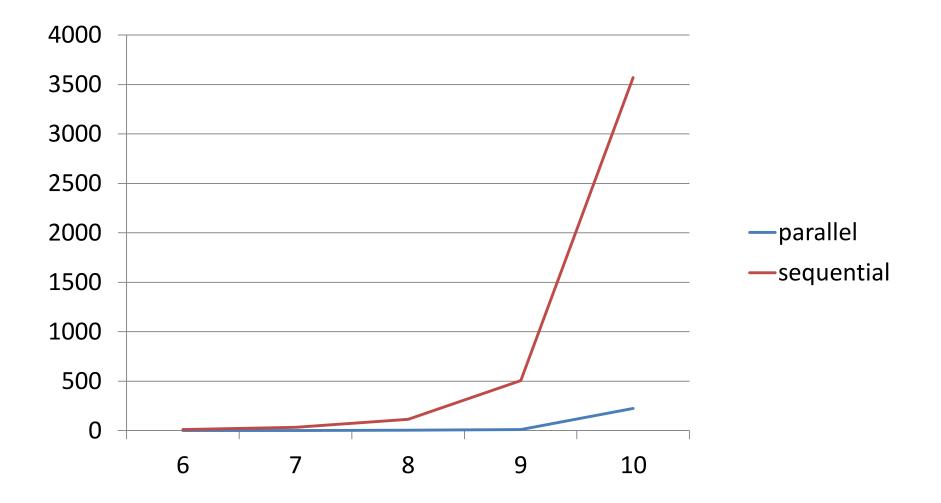
Optimization

- Adjacency check is done in parallel by OpenMP threads
- All send messages are non-blocking

Timing (seconds)

n	parallel	sequential
	(no. ranks)	
6	1.14 (24)	11
7	1.23 (40)	33
8	3.12 (80)	115
9	10.02 (64)	505
10	223.6 (104)	3570

Timing



Thank You