

From Molecular Dynamics towards a Node-Level Auto-Tuning Library for N-Body Simulations



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Molecular Dynamics (MD) Simulations

Applications:

Chemical Engineering:
cavitation, surface tension,
gas separation, etc..

Goals:

- High node-level performance in arbitrary scenarios.
- Minimize time to solution.

Main Challenges:

- Drastic impact of simulation variables on time to solution.
- Performance depends on many variables that can change during runtime.

Is1-mardyn

Highly parallel MD Code

- Language: C++.
- Small rigid, multi-centered molecules
- Double, single and mixed precision
- Implemented Interactions:
 - 12-6 Lennard-Jones
 - Coulomb
 - Charge
 - Dipole
 - Quadrupole
- Highly efficient reduced memory mode enabling largest known MD simulation: $2 \cdot 10^{13}$ molecules.

Node-Level Challenges

Factors affecting performance

- Number of particles
- Particle density distribution
⇒ Can change over time!
- Number of Lennard Jones centers
- Cutoff radius
- Heterogeneous Hardware

Can change time to solution and performance by orders of magnitude!

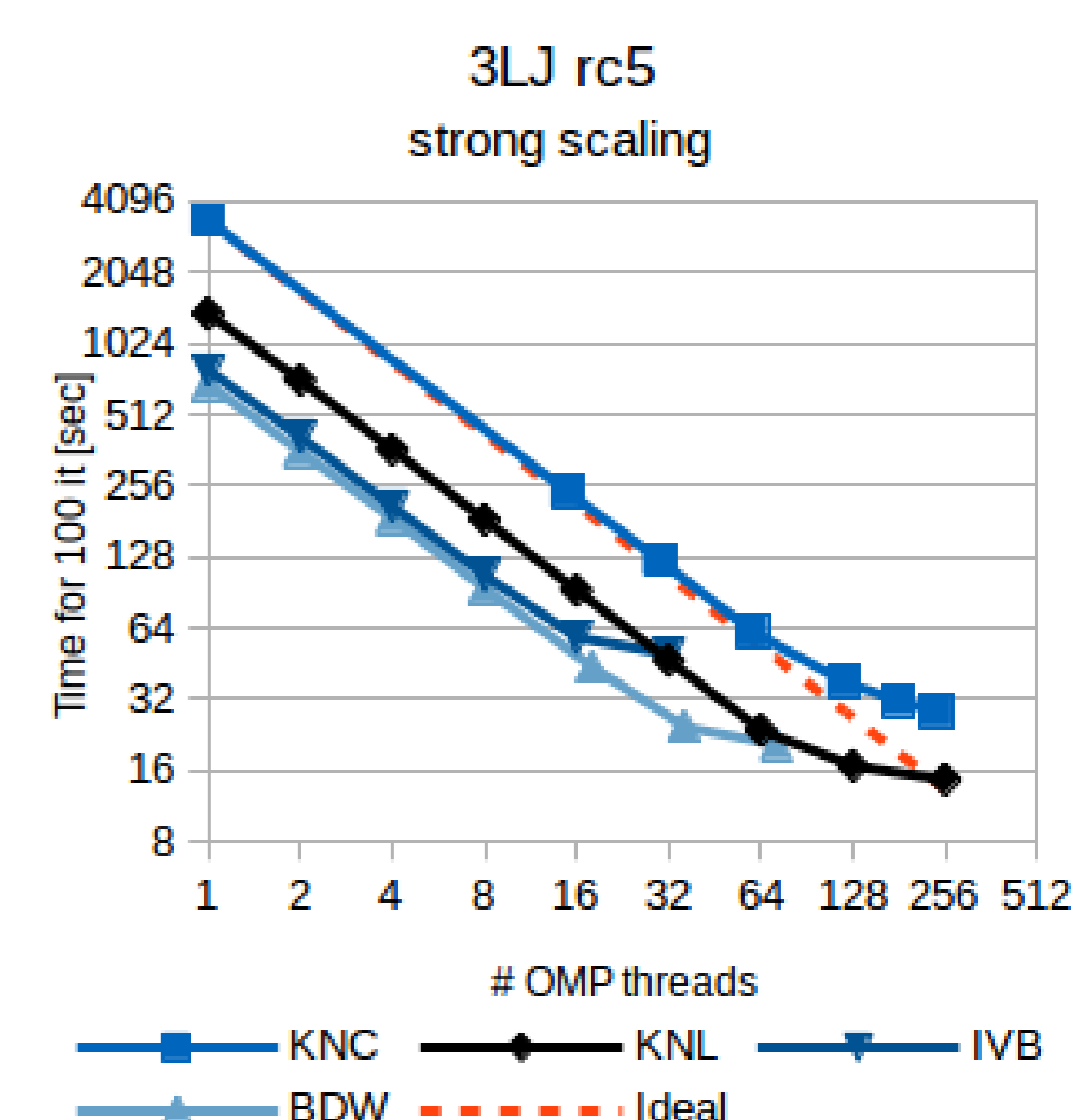
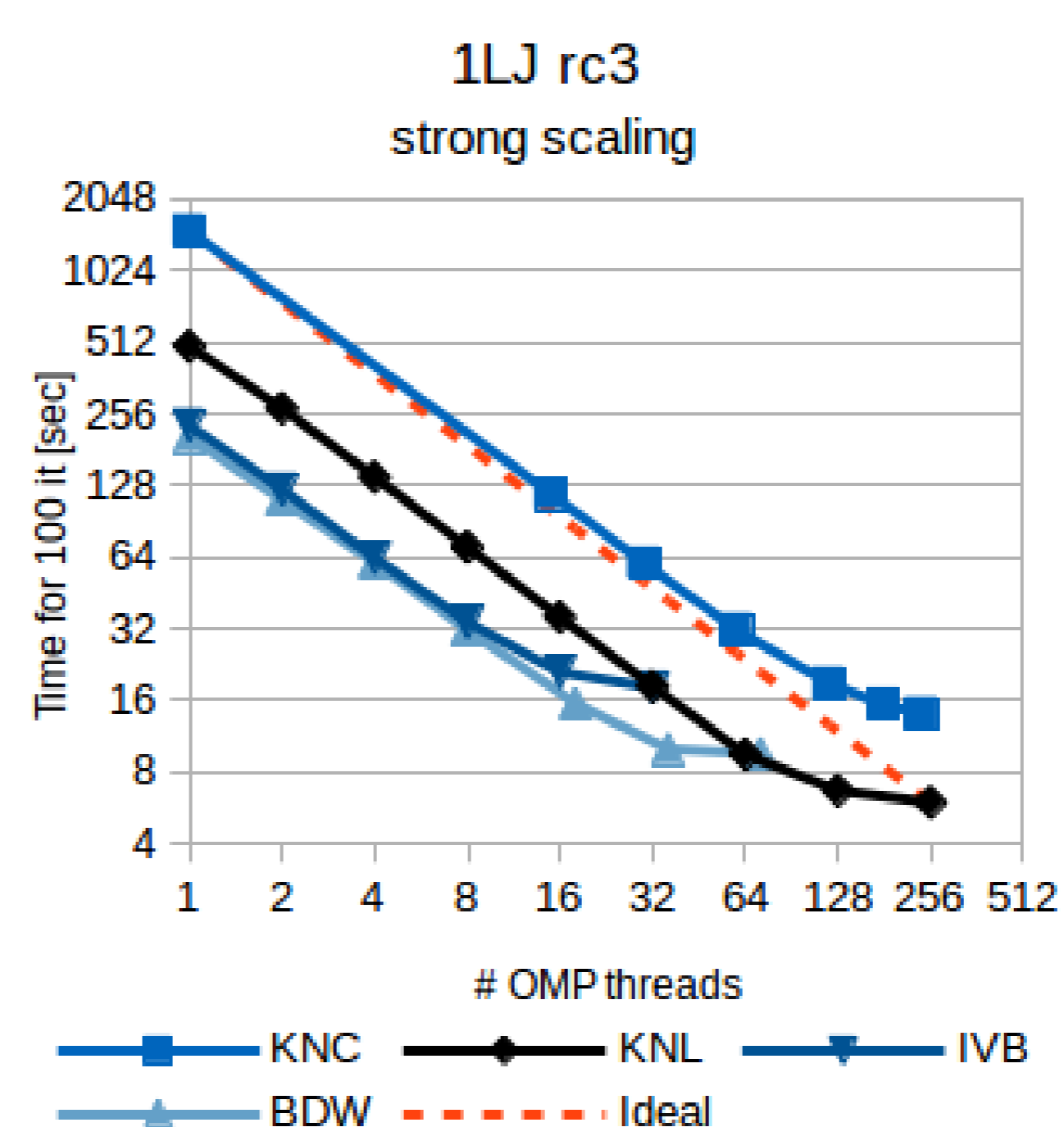
Potential Solutions

- Vectorization of Kernels
- Container (Linked Cells, Verlet Lists)
- Traversal patterns

Results

Homogeneous Scenario

- Large scenario: 1.3 million molecules
- Homogeneously distributed
- OpenMP with 8-way coloring scheme
- Different platforms



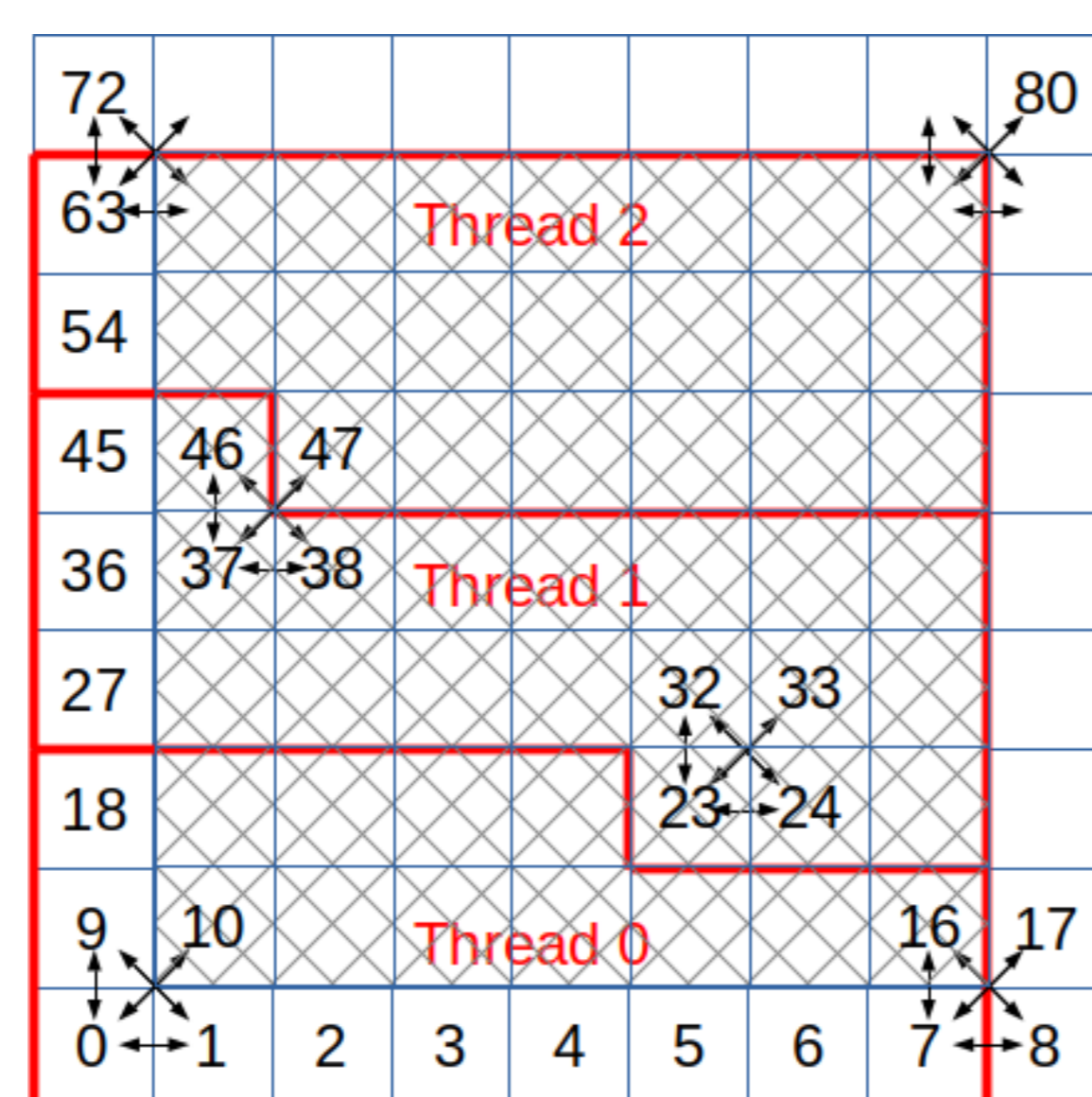
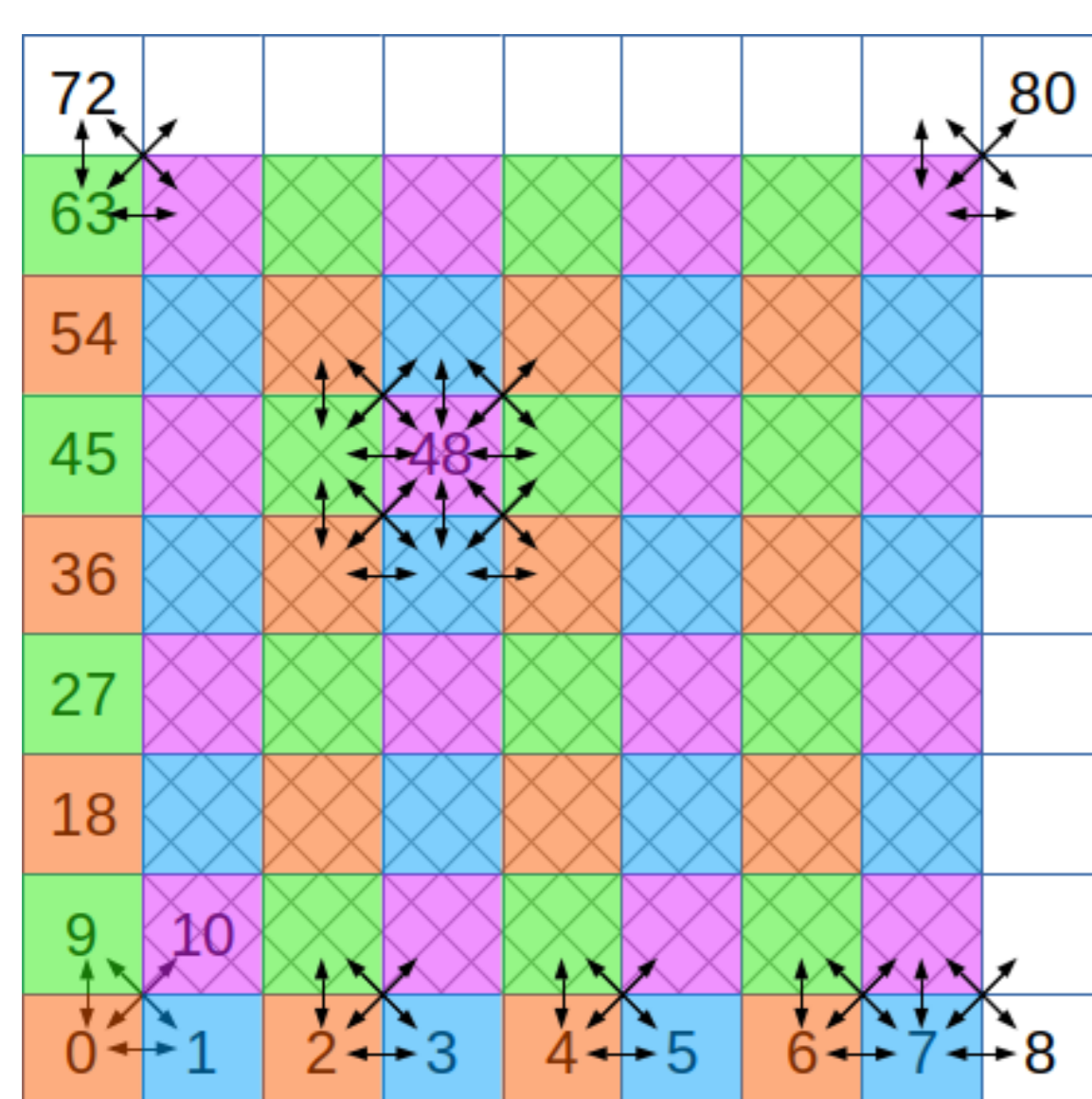
Available Traversal Options

c08

- 8-way coloring scheme (3D).
- ⇒ One barrier per color.
- Load balancing via OpenMP.

slice

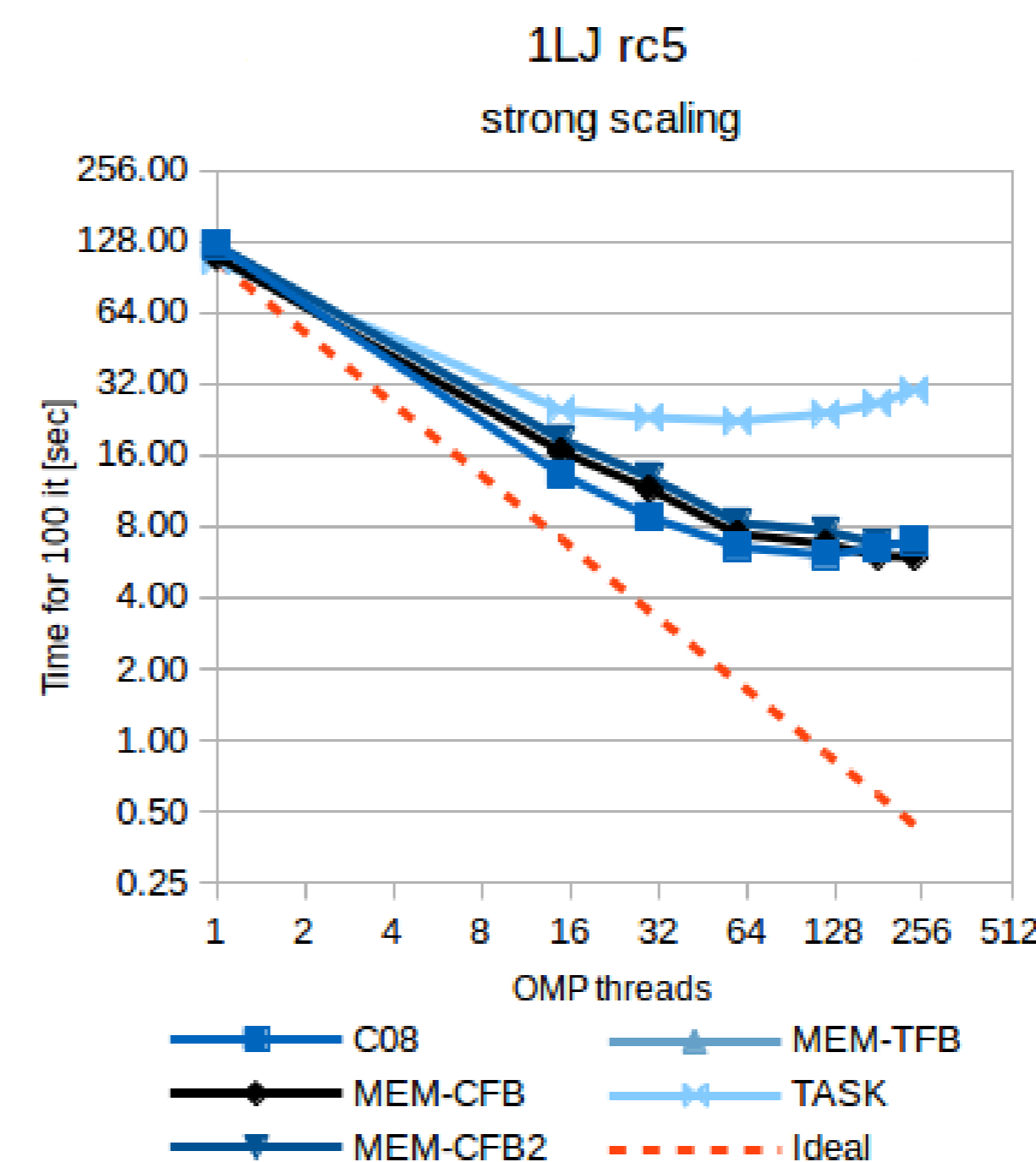
- One slice per thread.
- Only one lock per thread.
- No load balancing.



Comparison of OpenMP schemes: Towards Auto-Tuning

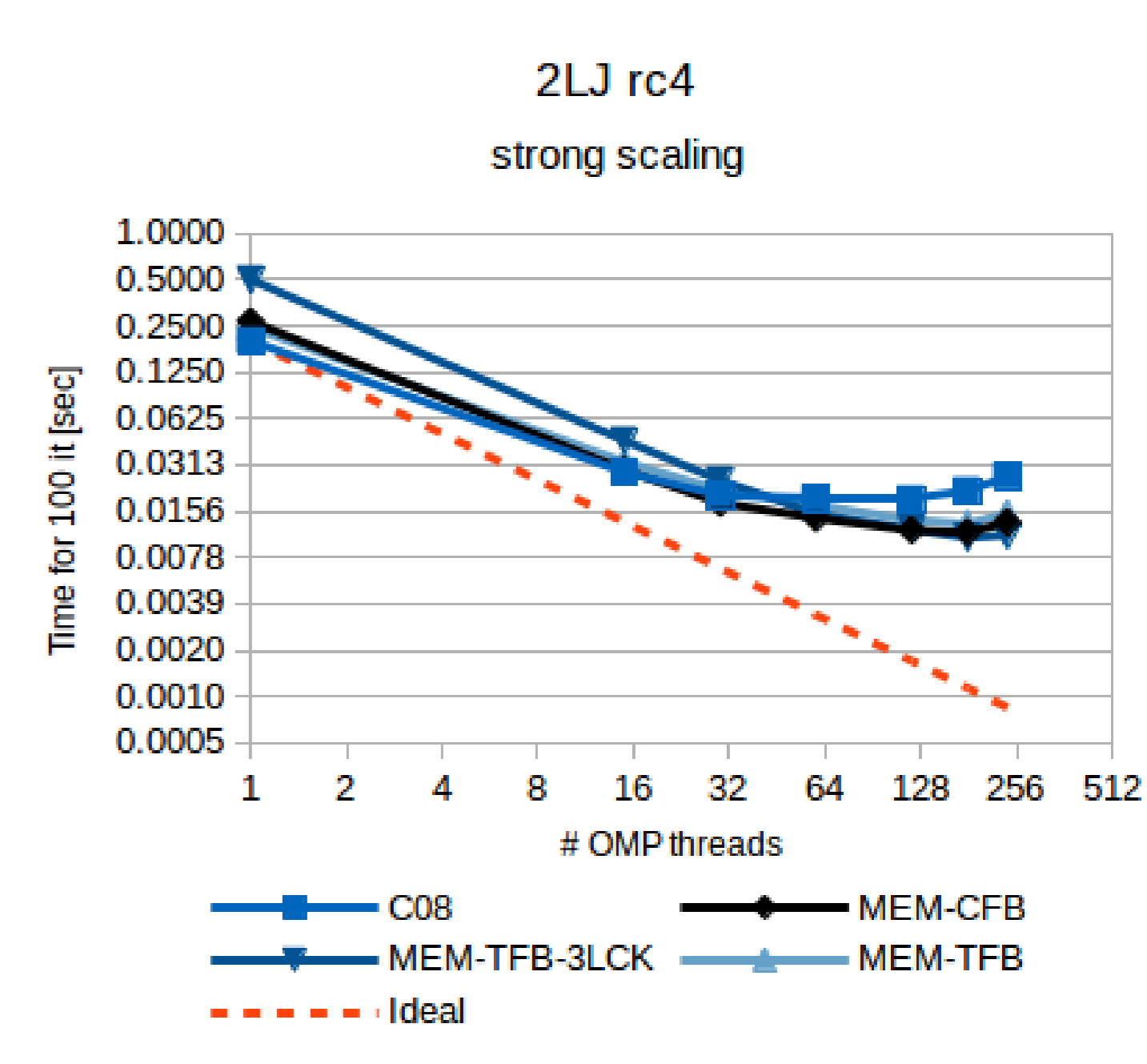
Inhomogeneous Scenarios

- Medium size: 40.000 molecules



Small Scenarios

- Small size: 1000 molecules



Library Goals

- **Optimal performance on arbitrary scenarios**
 - Simulation needs to be able to also handle domains with low particle count or inhomogeneous particle distributions.
 - ⇒ Specialized approaches needed.
- **More modular code structure**
 - Since different scenario settings benefit from dedicated techniques, these need to be easily exchangeable.
 - Example: Cell traversal pattern (see above)
 - ★ Direct Sum, c08, slice, ...

⇒ Auto-Tuning

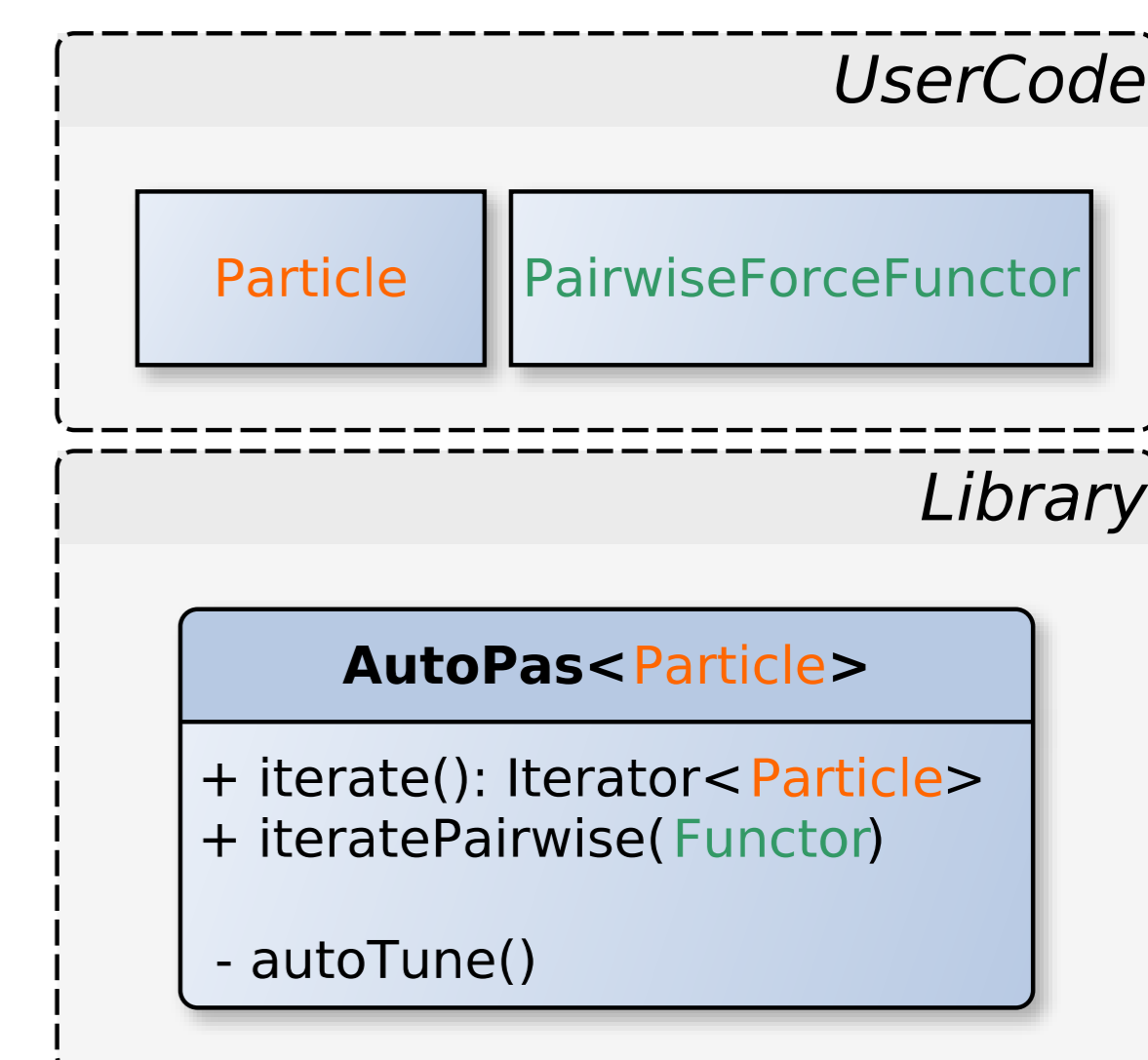
- Instead of the user, the code should find the optimal combination of techniques.
- Too many combinations possible to test all.
 - ⇒ Performance Modeling (e.g. automated empirical with ExtraP).
- Reevaluate combination during runtime and adapt appropriately.
- Provide flexibility through "strategy" software pattern.

There is no silver bullet
⇒ We want to export containers, traversals and kernels to make them available as a library.

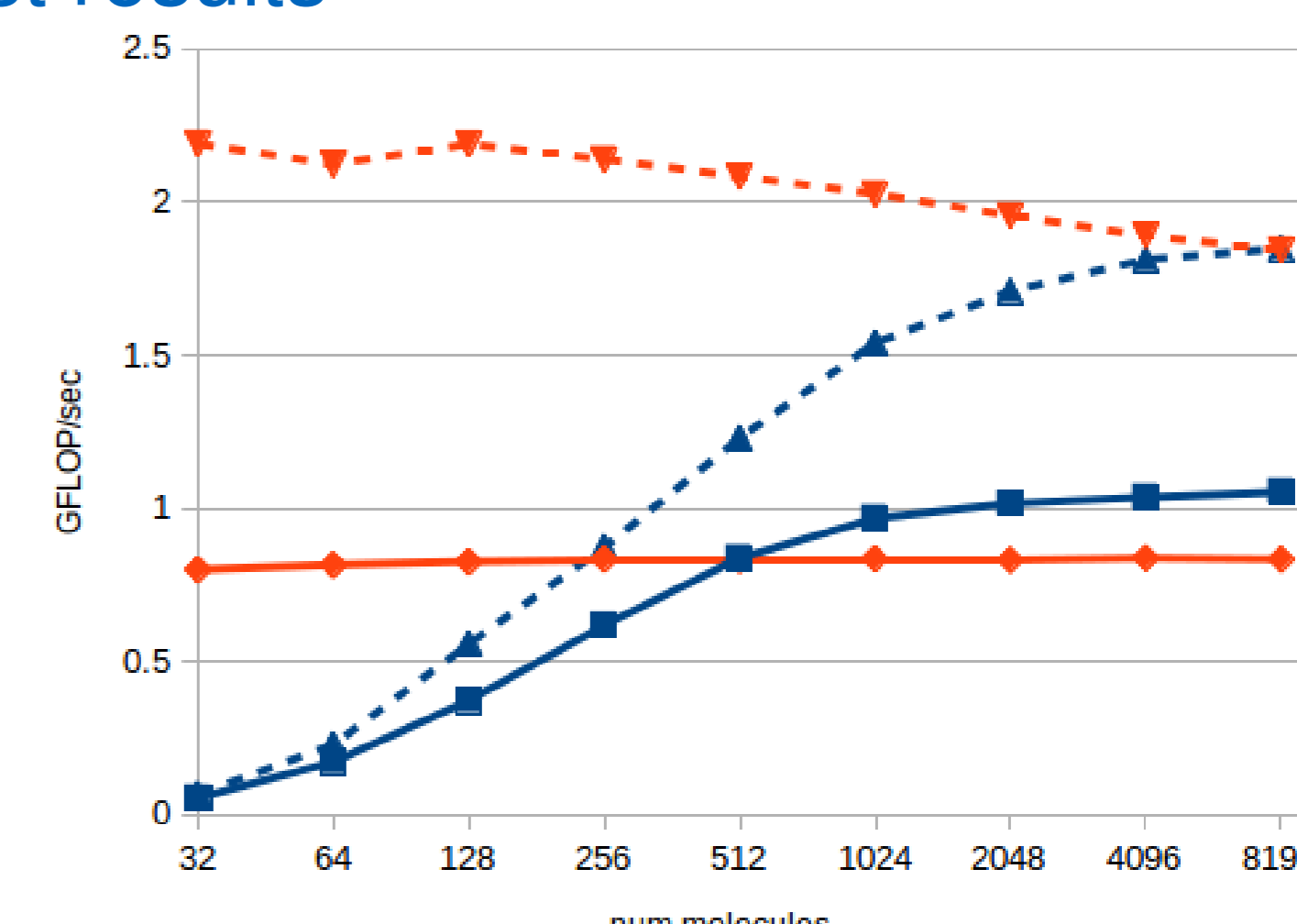
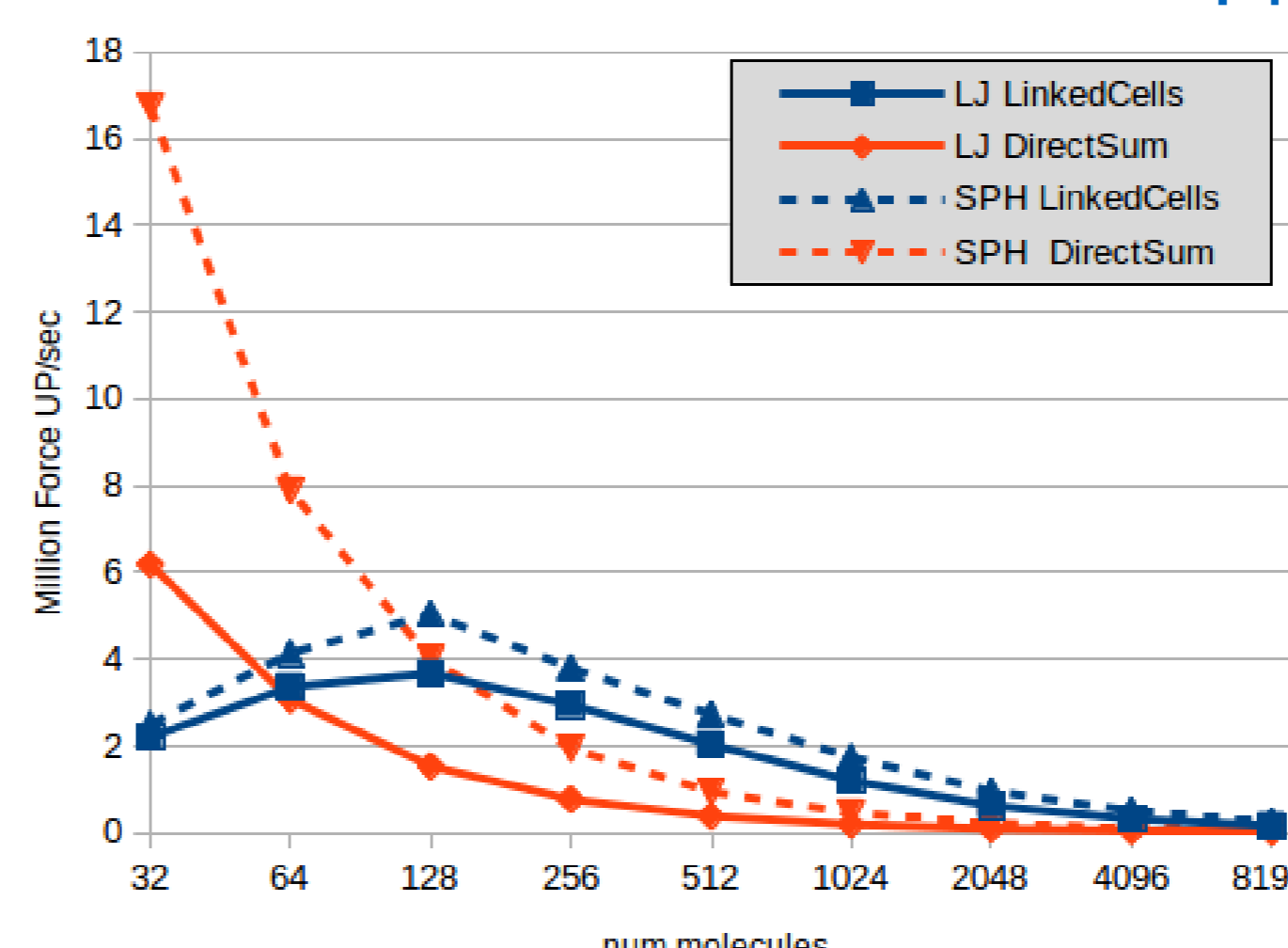
Vision

- Base to build full N-Body Simulations on top of.
- Manages node-level performance internally via Auto-Tuning with Performance Modeling.
- Modular C++ Template design to dynamically select optimal SIMD, OpenMP, Data-structures, etc. at runtime.

Outlook: AutoPas



First results



References

- [1] N. Tchipev, A. Wafai, C. W. Glass, W. Eckhardt, A. Heinecke, H.-J. Bungartz, and P. Neumann, "Optimized force calculation in molecular dynamics simulations for the intel xeon phi," in *European Conference on Parallel Processing*, pp. 774–785, Springer, 2015.
- [2] N. Tchipev and et al., "Twetris: Twenty trillion-atom simulation." submitted, 2018.
- [3] N. Tchipev, A. Costinescu, S. Seckler, P. Neumann, and H.-J. Bungartz, "Towards autotuning between openmp schemes for molecular dynamics on intel xeon phi." 2017. SIAM CSE '17.



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