## P&S Heterogeneous Systems

# Accelerating Agent-Based Simulations with BioDynaMo

#### Lukas Breitwieser

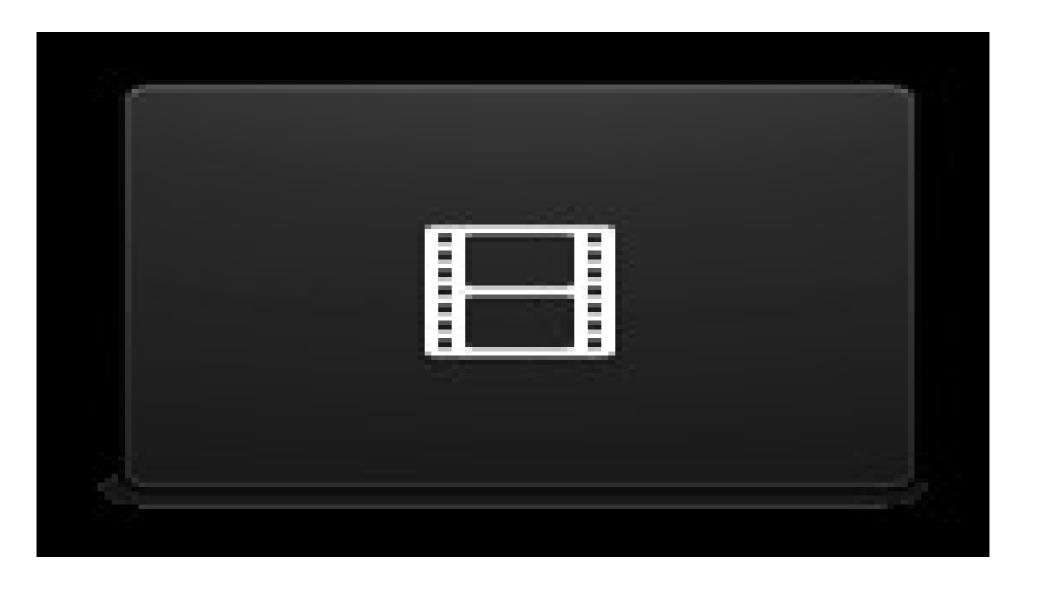
ETH Zürich

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# Introduction to Agent-Based Simulation

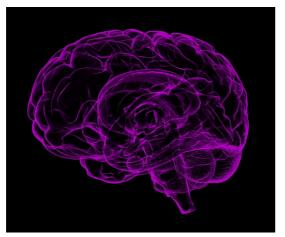
## Modeling complex systems

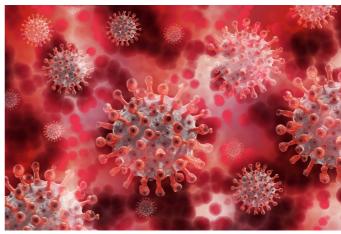


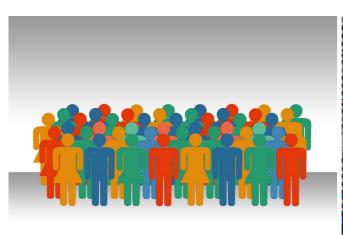
#### What is agent-based simulation?

- Also called individual-based modeling
- Bottom-up approach
  - Modeling the trees not the forest
- Characteristics
  - Local interaction
  - Emergent behavior

### Agent-based simulation is very versatile











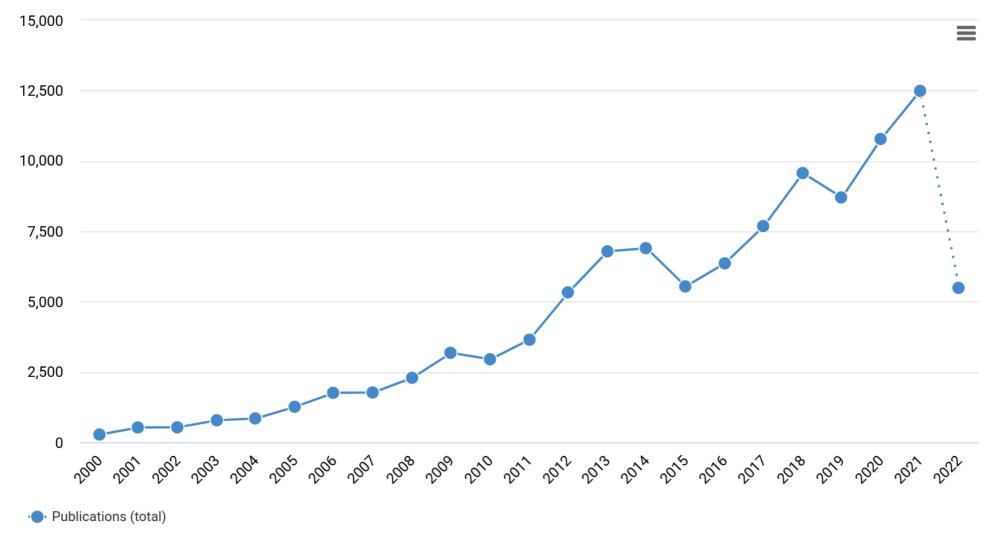
### Agent-based simulation is very versatile (cont.)

Table 1: A sample of recent agent-based applications

	Tuble 1.11 sample of recent agent based approaches
Application Area:	Agent-based Model Focus:
Agriculture	A spatial individual-based model prototype for assessing potential pesticide exposure of farm-workers conducting small-scale agricultural production (Leyk et al. 2009)
Air Traffic Control	Air traffic control to analyze control policies and performance of an air traffic management facility (Conway 2006)
Anthropology	Prehistoric settlement patterns and political consolidation in the Lake Titicaca basin of Peru and Bolivia (Griffin and Stanish 2007)
Biomedical Research	The Basic Immune Simulator, to study the interactions between innate and adaptive immunity (Folcik et al. 2007)
Crime Analysis	A realistic virtual urban environment, populated with virtual burglar agents (Malleson 2010)
Ecology	Investigate the trade-off between road avoidance and salt pool spatial memory in the movement behavior of moose (Grosman et al. 2011)
	Predator-prey relationships between transient killer whales and other marine mammals (Mock and Testa 2007)
Energy Analysis	A building occupant network energy consumption decision-making model (Chen et al. 2011)
	Application for the Smart Grid (Jackson 2010)
	Energy investment decision making (Tobias 2008)
	Oil refinery supply chain (Van Dam et al. 2008)
Epidemiology	Pandemic disease model accounting for individual behavior and demographics (Aleman et al. 2010)
	Global-scale agent model of disease transmission (Parker and Epstein 2011)
Evacuation	Tsunami evacuation using a modified form of Helbing's social-force model applied to agents (Puckett 2009)
Market Analysis	Consumer marketing model developed in collaboration with a Fortune 50 firm (North et al. 2009)
	Consumer airline market share (Kuhn et al. 2010)
	Simulation that models the possibilities for a future market in sub-orbital space tourism (Charania et al. 2006)
Social Networks	Model of email-based social networks, in which individuals establish, maintain and allow atrophy of links through contact-lists and emails (Menges et al. 2008)

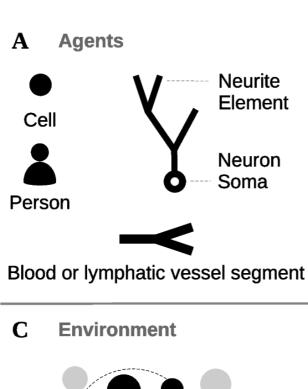
#### Rising Number of Publications in this field

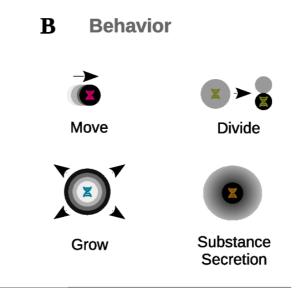
 Keywords used: "agent-based modeling" or "agents-based simulation"

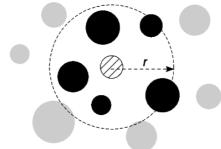


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#### Important building blocks







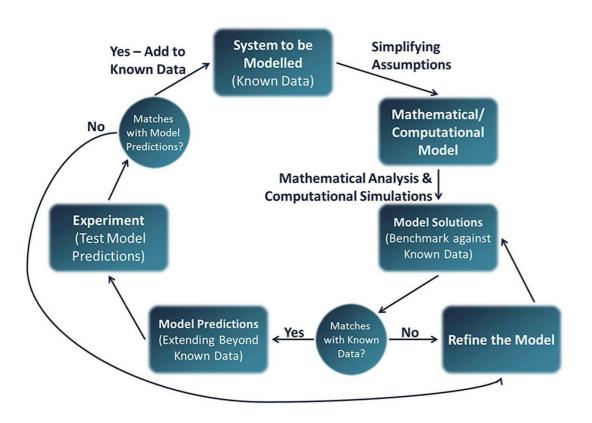
- Observed agent
- Agents inside environment
- Agents outside environment

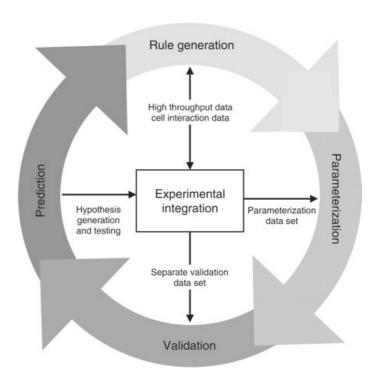
#### **D** Simulation Algorithm

// Define initial model
Place simulation objects in space
Set their attributes
Define behavior

// Run simulation
for each simulation step
 Update environment
 for each agent
 for each agent operation
 Run agent\_operation(agent)
 for each standalone operation
 Run standalone\_operation()

#### The process of developing an ABM

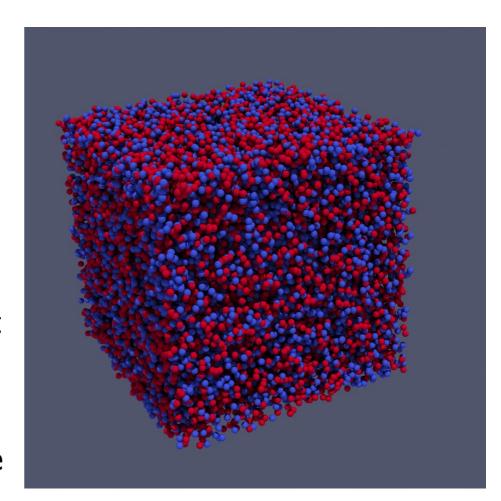




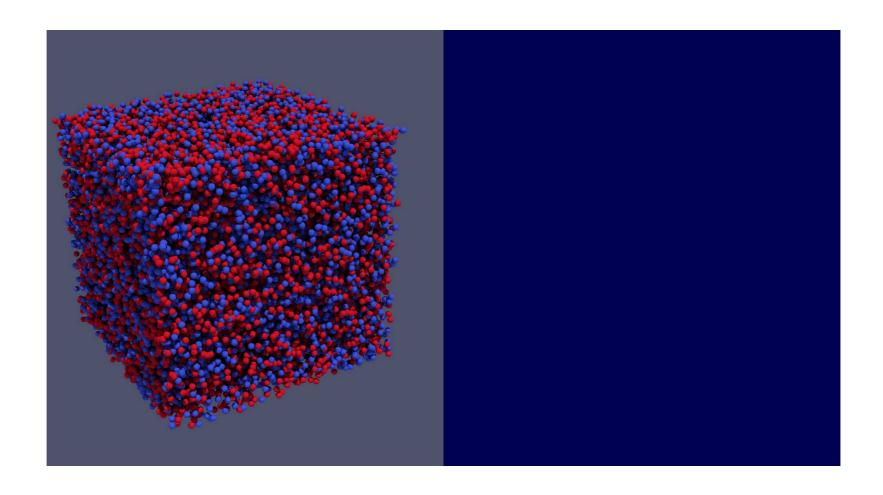
## Demo

#### Cell clustering model

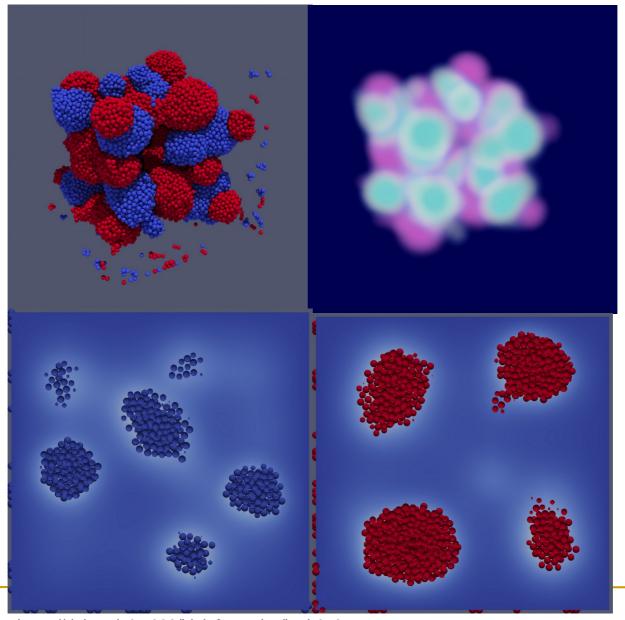
- Agent: Cell
  - Spherical shape
  - cell type
- Behaviors
  - Secrete a substance into the extracellular matrix
  - Follow the concentration gradient (chemotaxis)
- Initial condition
  - Randomly distributed in 3D space



## Cell clustering video



## Cell clustering result



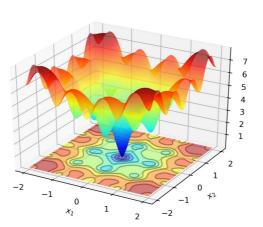
# Performance considerations

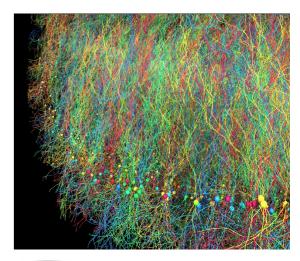
#### The problem

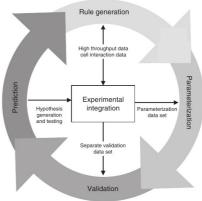
Existing simulation platforms do not always take full advantage of modern hardware.

#### Impact of low performance

- Limitation of the size and complexity of models
- Longer development time
- Limited capability to explore parameter space
  - → less optimal solution
- Increased cost







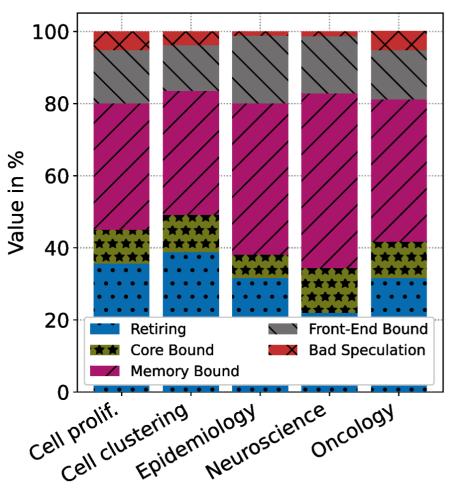
#### Our solution

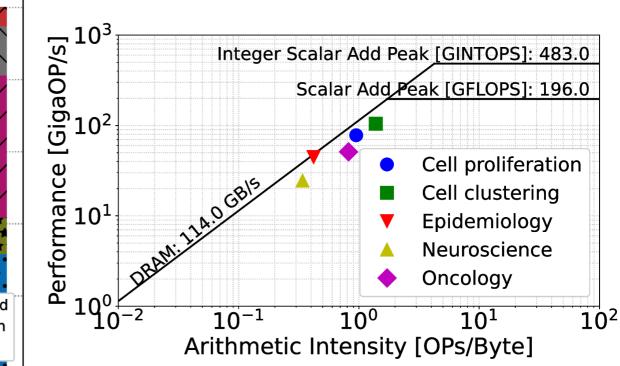
BioDynaMo a high-performance and modular, agent-based simulation platform written in C++.

#### Features and abstraction layers

#### Simulation Simulation Agent geometry: sphere, cylinder Agents: Cell, NeuronSoma, NeuriteElement Behaviors: Secretion, Chemotaxis, Proliferation, BioDynaMo's model GeneRegulation building blocks Extraculluar diffusion Agent interaction force Generation of agent populations Parameter management Agent reproduction & mortality Parameter optimization BioDynaMo's high-level **Environment search** Hierarchical model support Hybrid-modeling Multi-scale simulations features Dynamic scheduling Space boundary conditions Statistical analysis Parallelism & thread-safety Web-based interface BioDynaMo's low-level Performance optimizations Backup & restore of simulations **GPU** support features Quality assurance Visualization infrastructure Libraries ROOT **Others OpenMP ParaView** Linux / MacOS Operating System (Multi-core) CPUs **GPU** Hardware

#### Challenge: Agent-based workload is memory-bound





#### Required efficient agent-based primitives

- Iterate over agents and execute operations
- Add and remove agents to/from the simulation
- Efficient environment implementation

#### Algorithm 1: Agent-based simulation algorithm

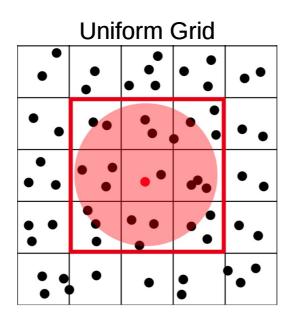
```
1 ModelInitialization()
2 for i \in iterations do
         for op \in pre\_standalone\_operations do
               op():
5
         end
6
         wait()
7
         parallel for a \in agents do
               for op \in agent\_operations do
                     op(a);
10
               end
11
         end
12
         for op \in standalone\_operations do
13
               op();
         end
15
         wait()
         for op \in post\_standalone\_operations do
17
               op();
         end
19 end
```

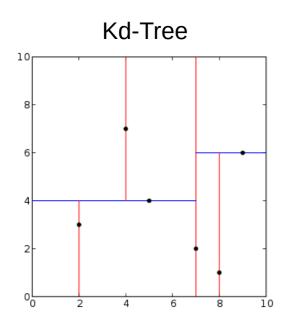
#### Performance Challenges and Improvements

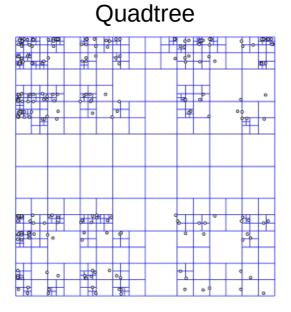
- Maximize parallelization
  - Optimized uniform grid to search for neighbors
  - Parallelize the addition and removal of agents
- Efficient thread synchronization during agent updates
- Minimize memory access latency
  - NUMA-aware iteration
  - Agent Sorting and Balancing
  - Pool-based memory allocator
- Offload computation to the GPU

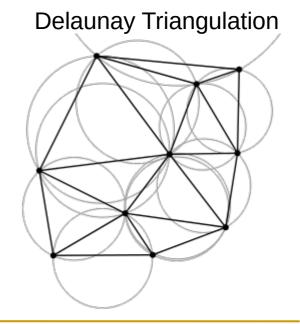
# Maximize parallelization

### Optimized uniform grid to search for neighbors



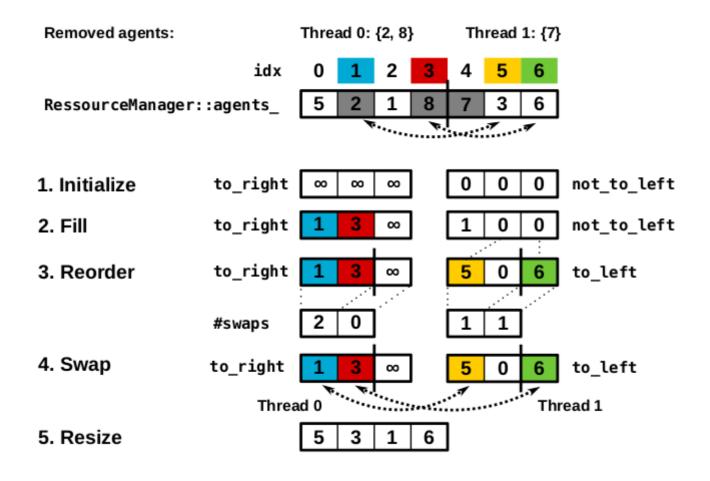






Source: Ahmad Hesam

#### Parallel agent removal mechanism

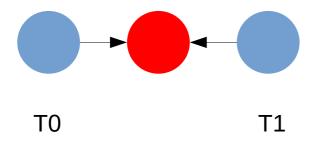


# Optimize Thread-Synchronization

#### Thread-synchronization (TS) during agent-updates

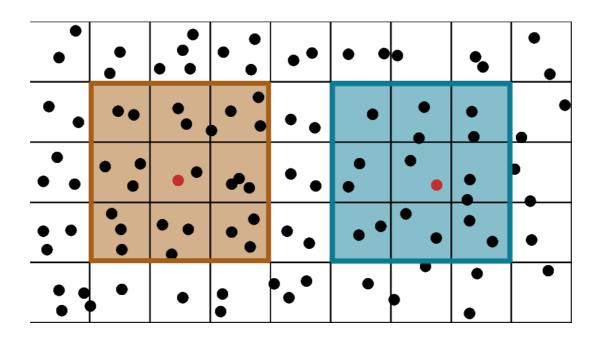
#### **Algorithm 1:** Agent-based simulation algorithm

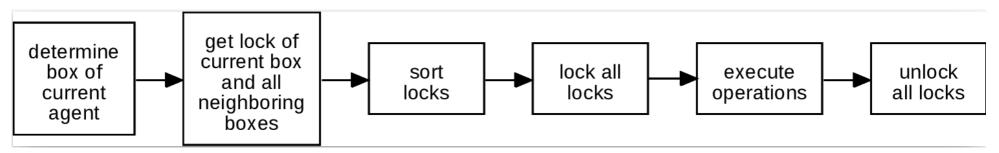
```
ModelInitialization()
   for i \in iterations do
         for op \in pre\_standalone\_operations do
              op();
         end
         parallel for a \in agents do
              for op \in agent\_operations do
                    op(a);
              end
11
         for op \in standalone\_operations do
12
13
14
         end
         for op \in post\_standalone\_operations do
16
17
               op();
19 end
```



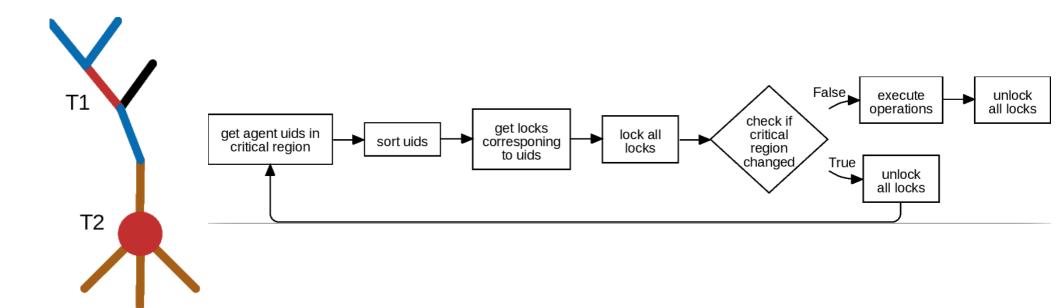
- Only necessary if agents modify their local environment.
  - Two agents (updated by two different threads) could attempt to modify the same neighbor.
- BioDynaMo provides two TS mechanisms
  - Automatic TS
  - User-defined TS

## Automatic thread-synchronization





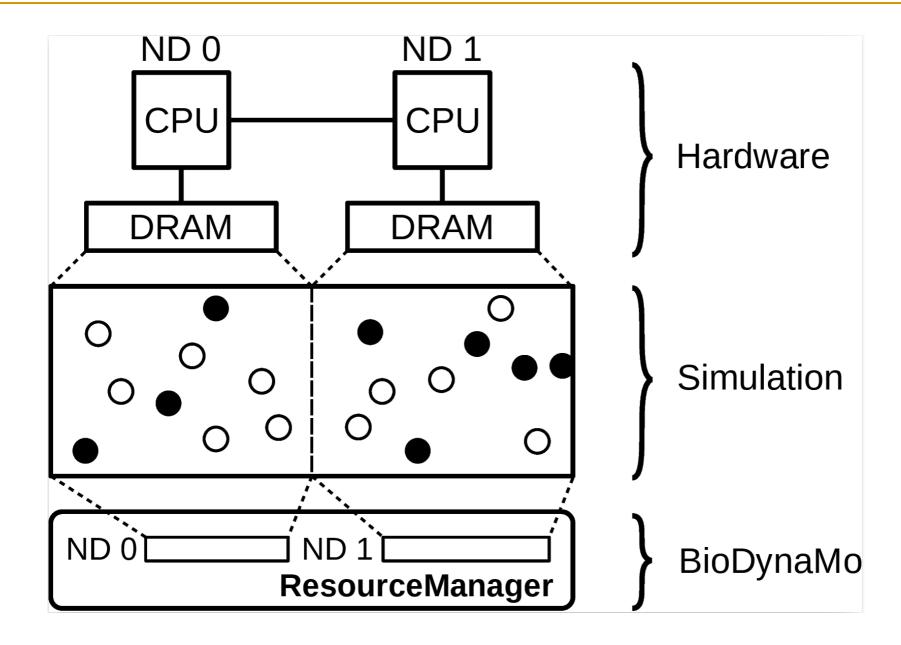
#### User-defined thread-synchronization



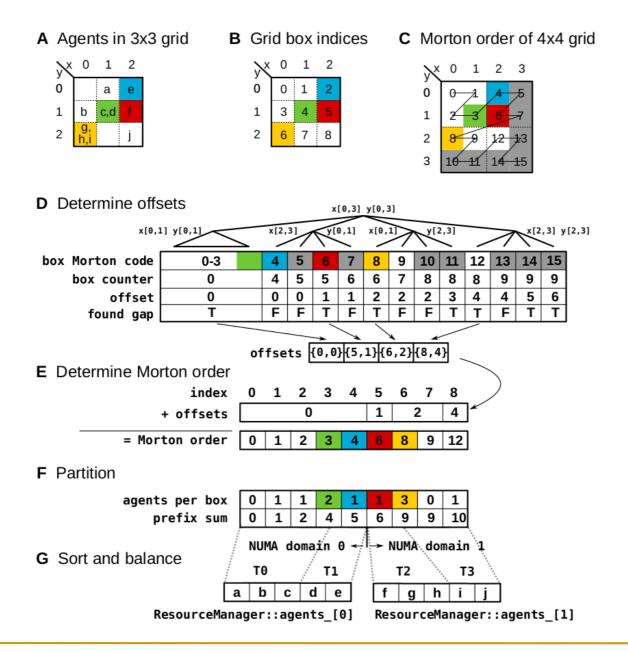
```
void NeuronSoma::CriticalRegion(std::vector<AgentPointer<>>* aptrs->reserve(daughters_.size() + 1);
  aptrs->push_back(Agent::GetAgentPtr<>());
  for (auto& daughter : daughters_) {
    aptrs->push_back(daughter);
  }
}
```

# Minimize Memory Access Latency

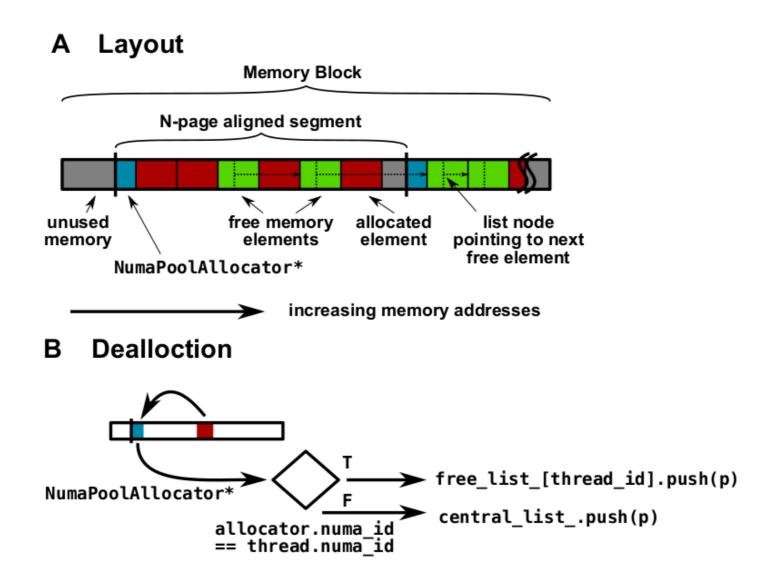
#### NUMA-aware iteration



#### Agent sorting and balancing mechanism



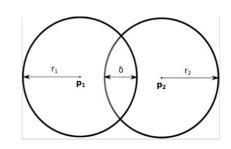
#### BioDynaMo Memory Allocator



# Offload computation to the GPU

#### BioDynaMo's GPU capabilities

- Operations can have implementations for different compute targets (CPU, GPU, and FPGA).
- If an operation has multiple implementations, the scheduler decides which one to use.
- Currently, BioDynaMo provides a GPU operation to calculate mechanical forces between spheres.



$$\delta = r_1 + r_2 - \|\mathbf{p}_1 - \mathbf{p}_2\|$$

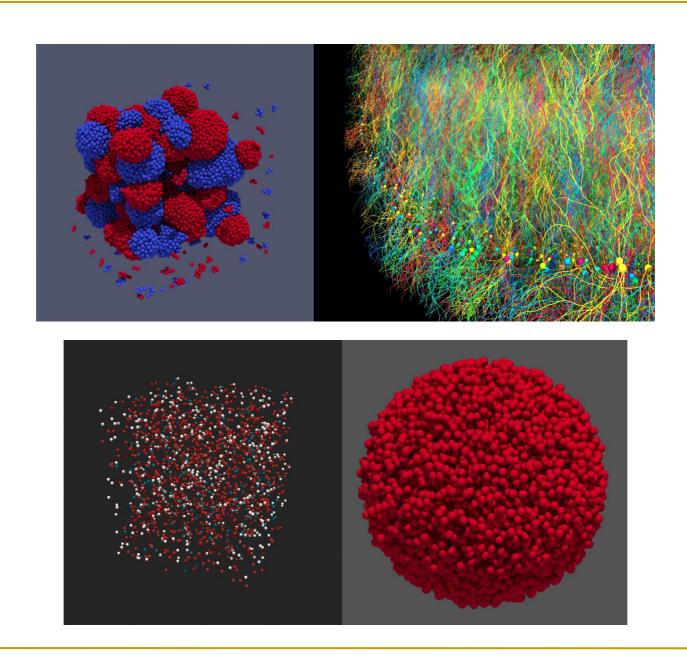
$$r = \frac{r_1 \cdot r_2}{r_1 + r_2}$$

$$\mathbf{F} = (\kappa \cdot \delta - \gamma \cdot \sqrt{r \cdot \delta}) \cdot \frac{\mathbf{p}_1 - \mathbf{p}_2}{\|\mathbf{p}_1 - \mathbf{p}_2\|}$$

Collision force computation

## Performance Evaluation

#### Benchmark simulations



#### Benchmark simulations characteristics

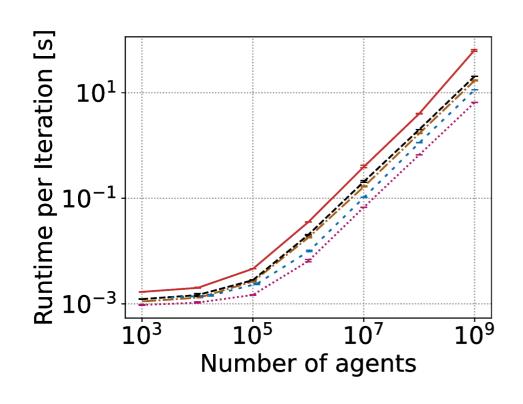
Characteristic	Cell proliferation	Cell clustering	Epidemiology use case	Neuroscience use case	Oncology use case
Create new agents during simulation	X			Х	Х
Delete agents during simulation					Х
Agents modify neighbors				X	
Load imbalance			X	X	
Agents move randomly			X		X
Simulation uses diffusion		X		X	
Simulation has static regions				X	
Number of iterations	500	1000	1000	500	288
Number of agents (in millions)	12.6	2	10	9	10
Number of diffusion volumes	0	54m	0	65k	0

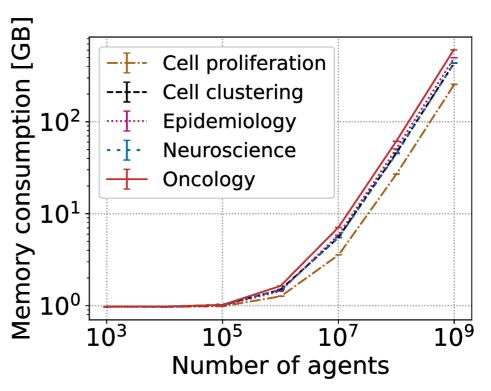
#### Benchmark hardware

#### TABLE II: Benchmark hardware

System	Main memory	CPU	OS
A	504 GB	Server with four Intel(R) Xeon(R) E7-8890 v3 CPUs	CentOS 7.9.2009
В	1008 GB	@ 2.50GHz with a total of 72 physical cores, two threads per core and four NUMA nodes.	
С	62 GB	Server with two Intel(R) Xeon(R) E5-2683 v3 CPUs @ 2.00GHz with a total of 28 physical cores, two threads per core and two NUMA nodes.	CentOS Stream 8

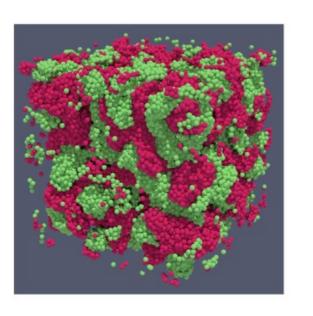
#### Runtime and Memory Complexity

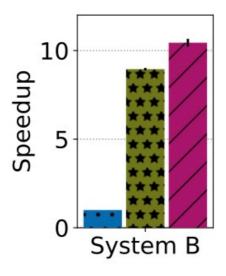


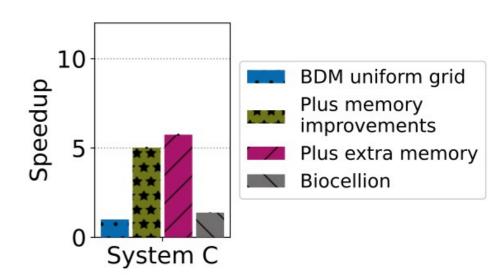


#### Comparison with Biocellion

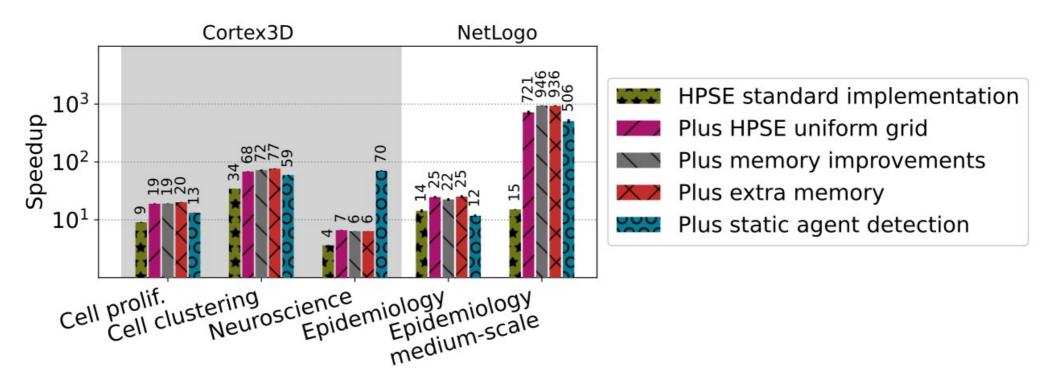
- Single-node 16 CPU cores; 13.4 million cells
  - → BioDynaMo is 4.15x faster
- Biocellion: 21 nodes, 672 CPU cores, 281 million cells BioDynaMo: one node, 72 CPU cores
  - → same runtime, but 9.3x fewer CPU cores used



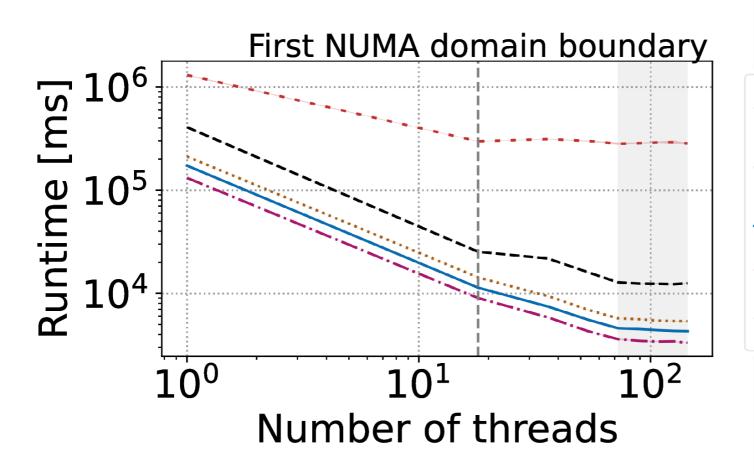


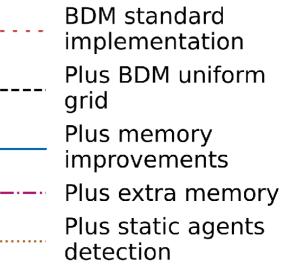


### Comparison with Cortex3D and NetLogo

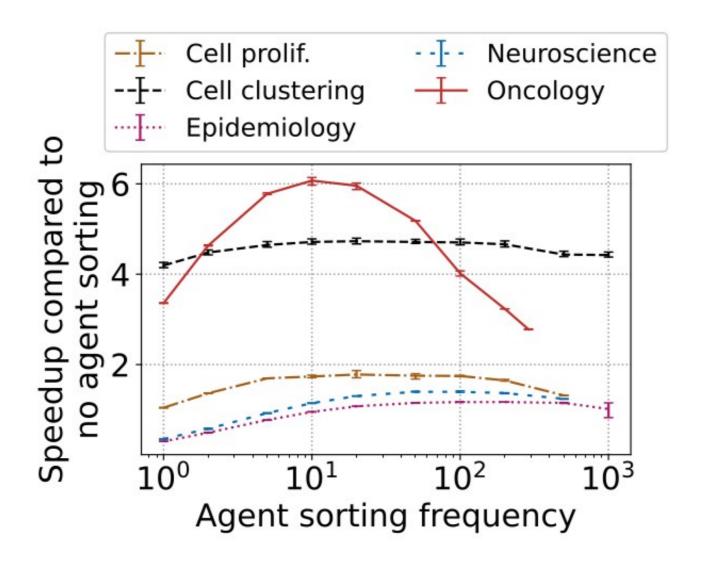


### Scalability





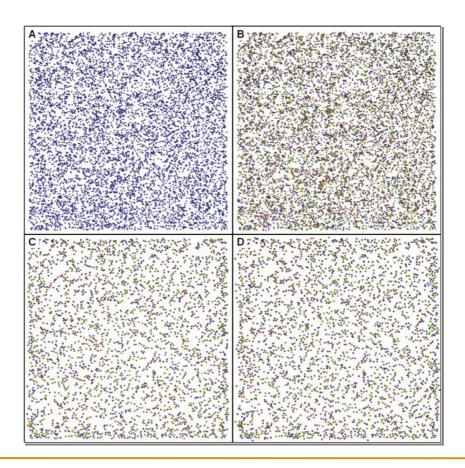
## Agent Sorting and Balancing



# Ongoing Use Cases

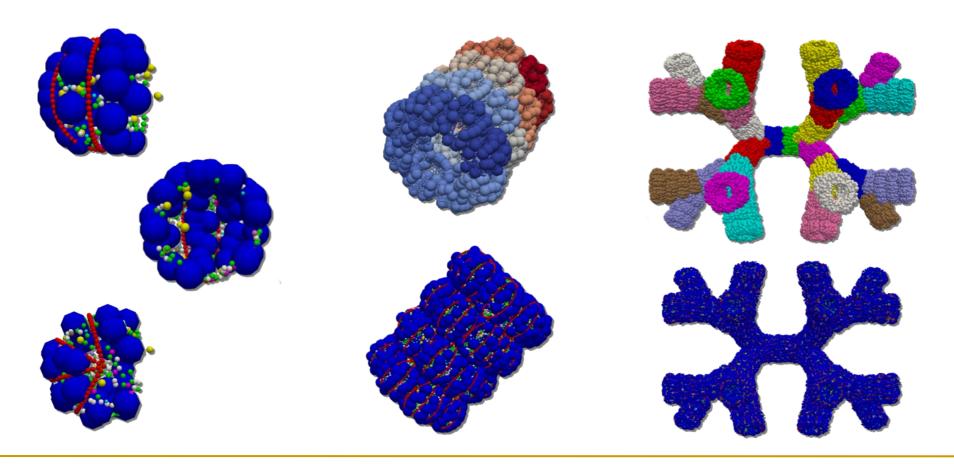
## Retinal self-organization

 Understand the mechanisms of cells self-organization during early development which is pivotal for their function.



#### Radiation-induced lung injury simulation

 Simulate onset of radiation pneumonitis and/or lung fibrosis in normal tissue after exposition to thoracic irradiation.

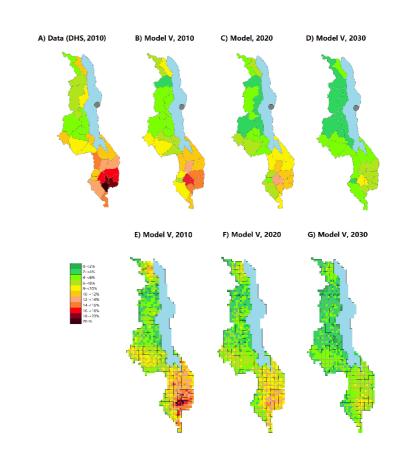


#### Spatial Spread of HIV in Malawi

- Collaboration with UniGE
- Original simulation written in R (Runtime: ~5.5h)
- Goal: speed up execution time
- Preliminary runtime with BioDynaMo: less than 2 minutes
- Further work
   needed to make
   models equivalent

## The spatial spread of HIV in Malawi: An individual-based mathematical model

Janne Estill, Wingston Ng'ambi, Liudmila Rozanova, Olivia Keiser doi: https://doi.org/10.1101/2020.12.23.20248757



#### Summary

- Agent-based simulation can be used to model many systems
- The presented optimizations improve the performance over state-of-the-art
  - Up to three orders of magnitude speedup
- These improvements allow BioDynaMo simulating billions of agents on a single server
- BioDyanaMo is currently being used in:
  - neurosience
  - oncology
  - epidemiology
  - cryobiology
  - socioeconomics
  - finance

- ...

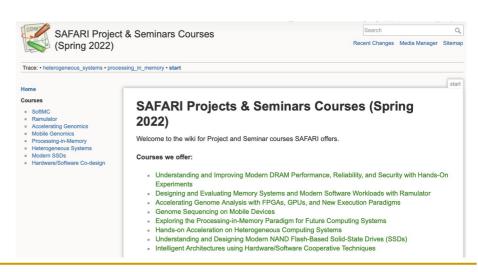
## Thank you for your attention!

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#### More P&S Courses: SSDs, Memory, Bioinformatics...

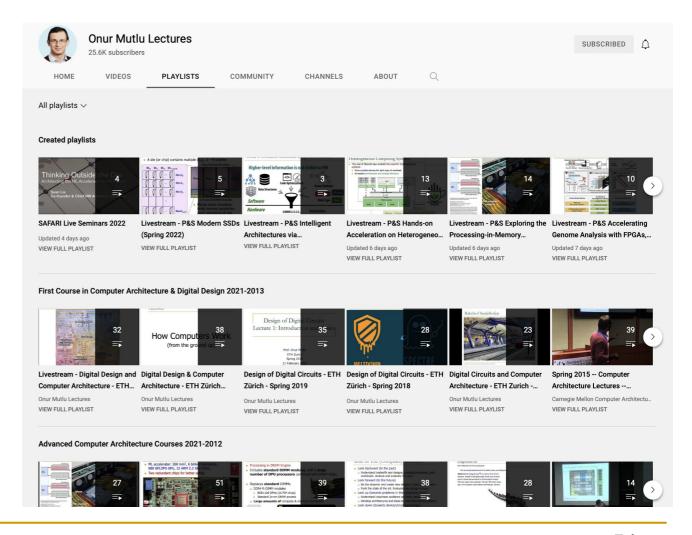
- Understanding and Improving Modern DRAM Performance, Reliability, and Security with Hands-On Experiments
- Designing and Evaluating Memory Systems and Modern Software Workloads with Ramulator
- Accelerating Genome Analysis with FPGAs, GPUs, and New Execution Paradigms
- Genome Sequencing on Mobile Devices
- Understanding and Designing Modern NAND Flash-Based Solid-State Drives (SSDs)
- Intelligent Architectures using Hardware/Software Cooperative Techniques

https://safari.ethz.ch/projects\_and\_seminars/spring2022/doku.php?id=start



#### More Resources: Onur Mutlu Lectures

- All P&S courses
- Digital Design and CompArch course
- Advanced CompArch course
- Seminar in CompArch



## P&S Heterogeneous Systems

## Accelerating Agent-Based Simulations

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11 July 2022