



BioDYNAMO
BIOLOGY DYNAMICS MODELLER

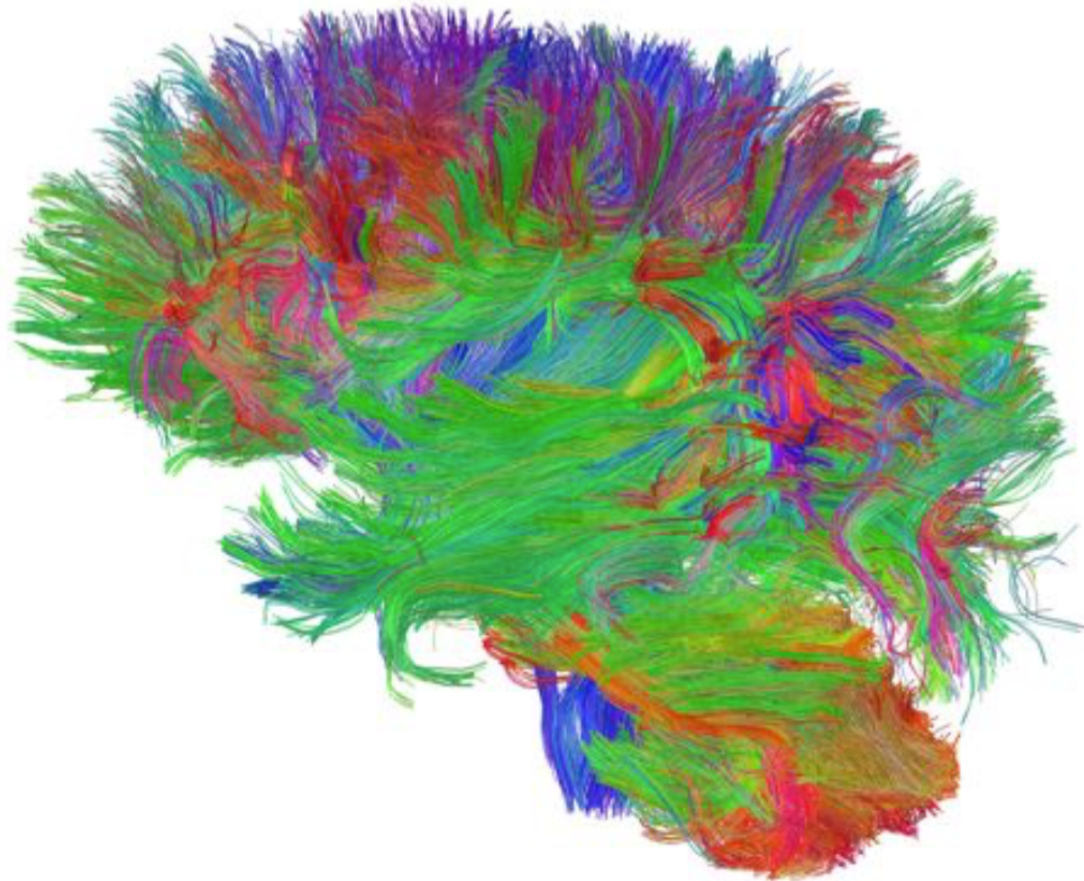
The Anticipated Challenges of Running Biomedical Simulations in the Cloud

Early-Career Researchers in Medical Applications
Short Talks on Computing and Simulation

Lukas Breitwieser



Understanding Developmental Diseases



Executive Summary

- **Motivation:**

Run *tightly-coupled* HPC workloads in the cloud

- Widely accessible, cost-effective

- **Problem:**

Data exchange between servers will be a bottleneck

- **Key Idea:**

 Exploit inherent simulation characteristics to reduce data volume



Outline

1. Biological Simulation Basics

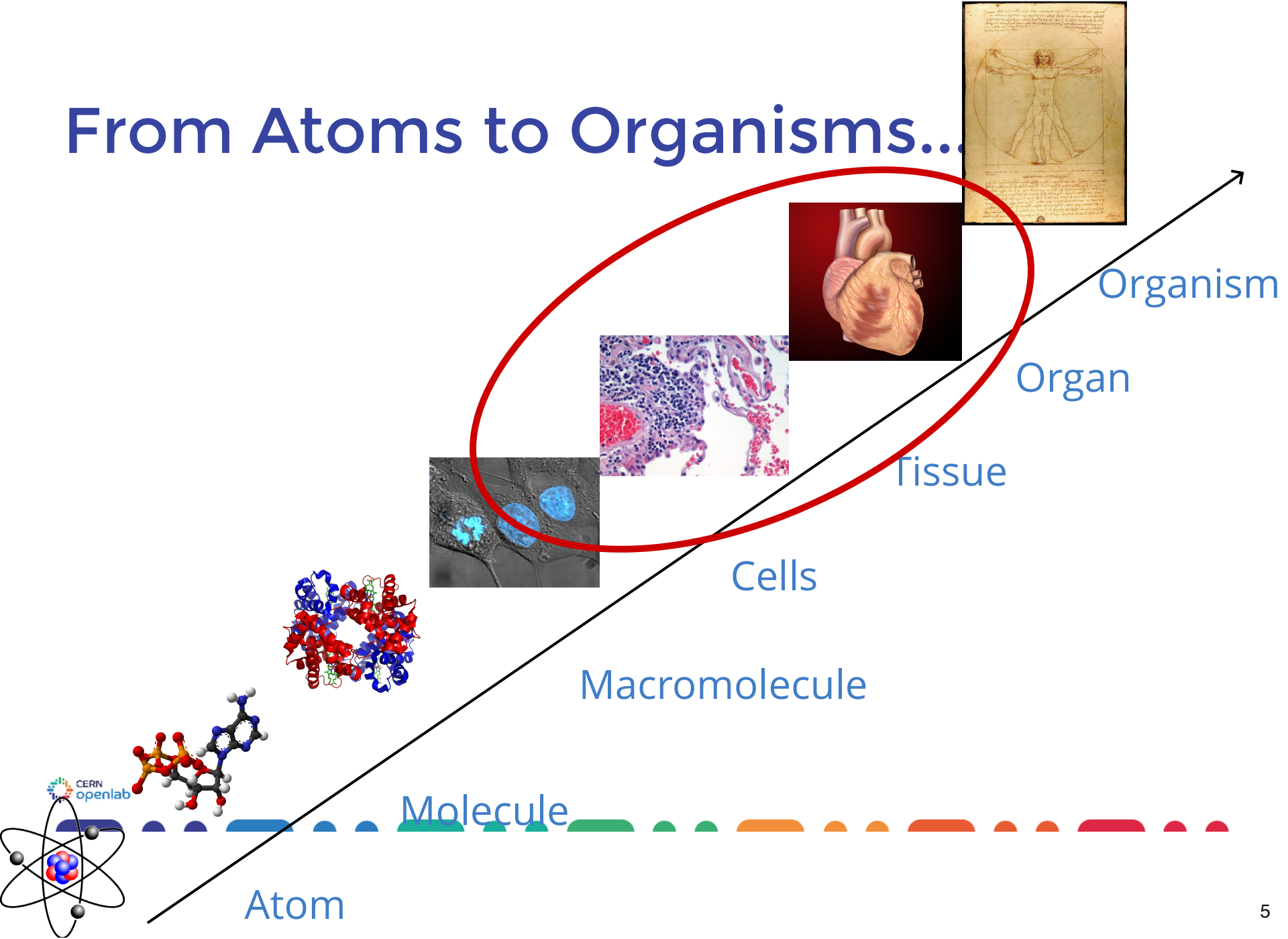
2. Distributed Runtime

3. HPC on Cloud

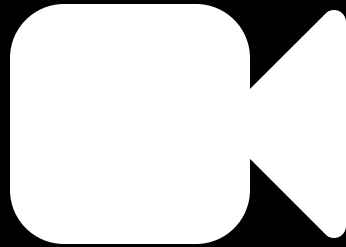
4. Data Movement Minimization



From Atoms to Organisms...

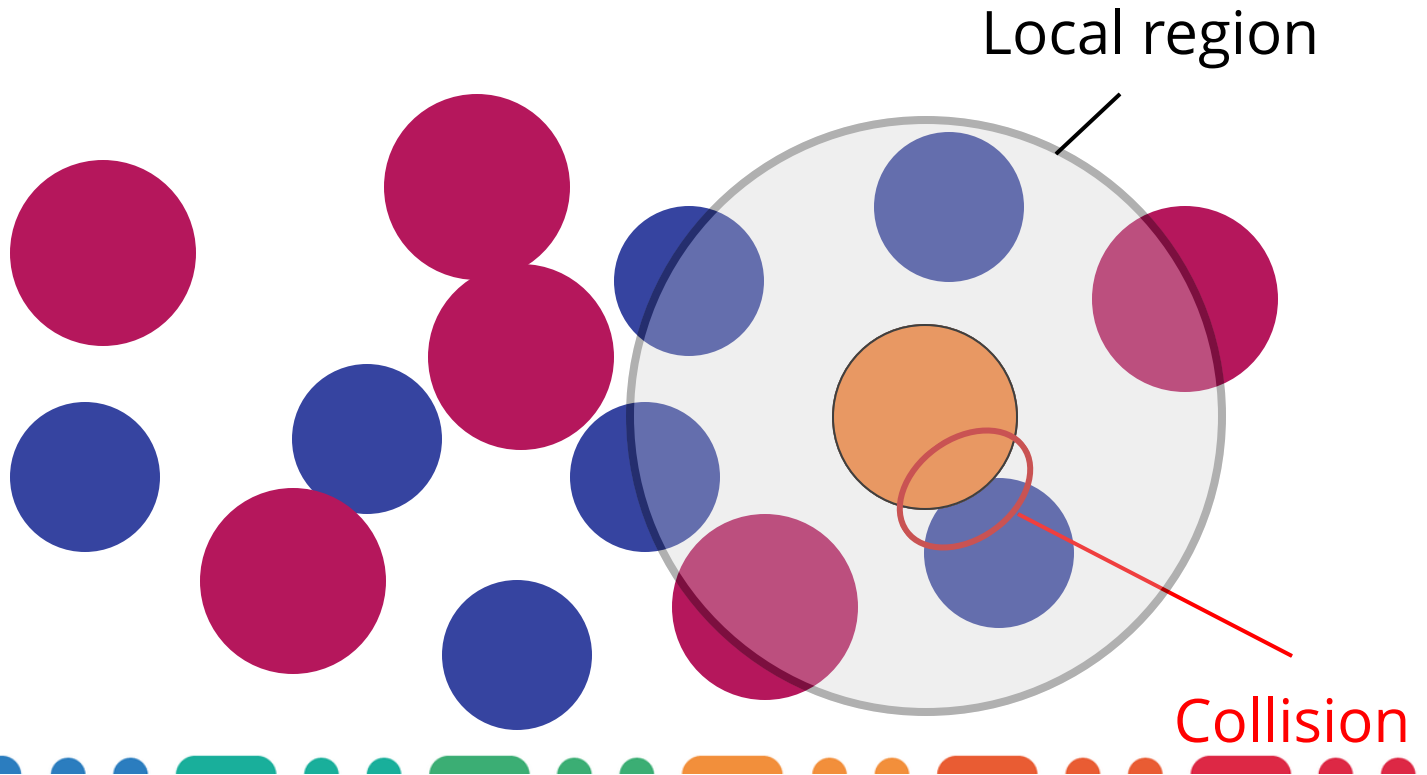


Demo: Tumor Growth



Agent-based simulations

Simulation object = *Agent*



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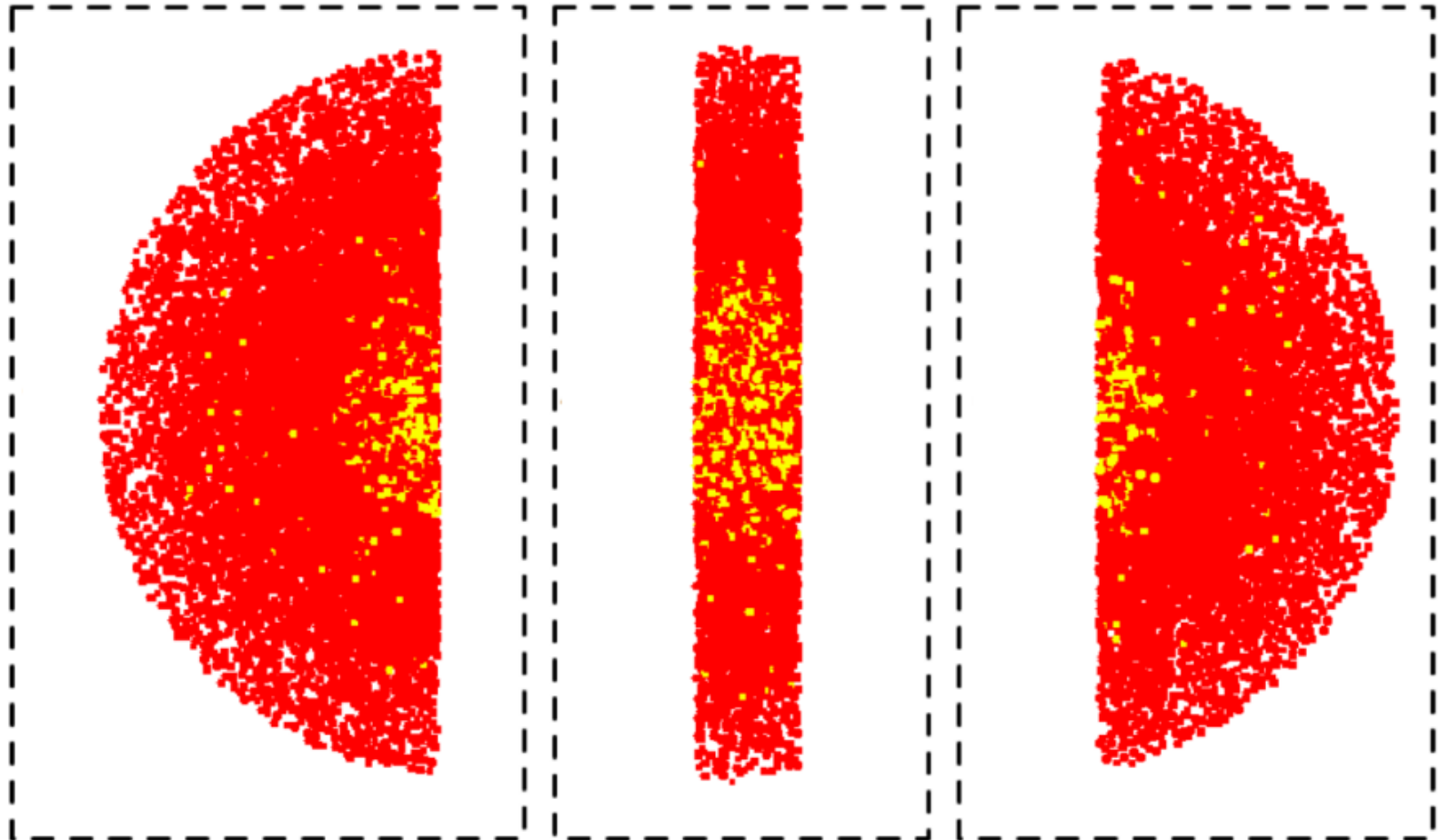
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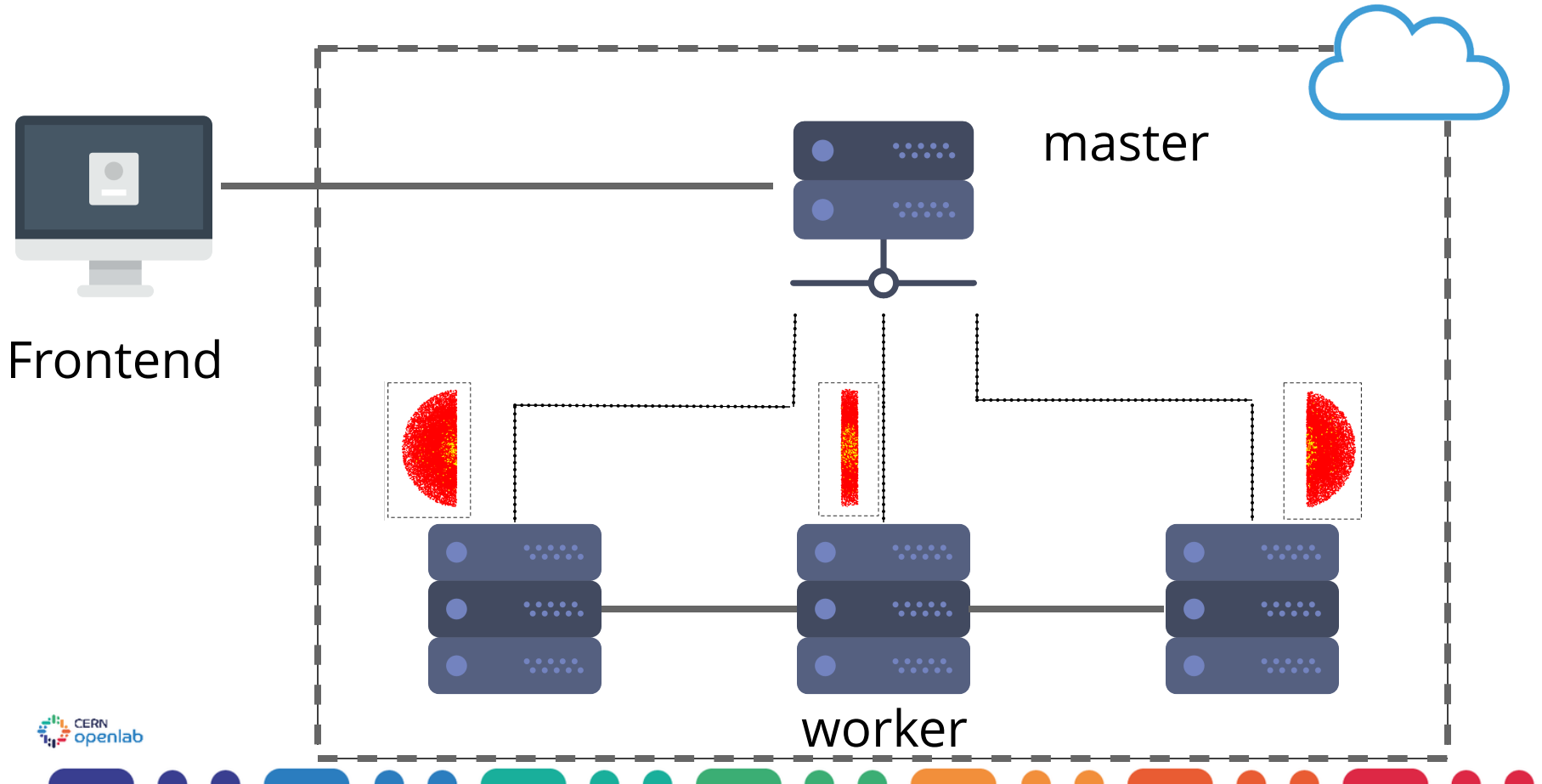
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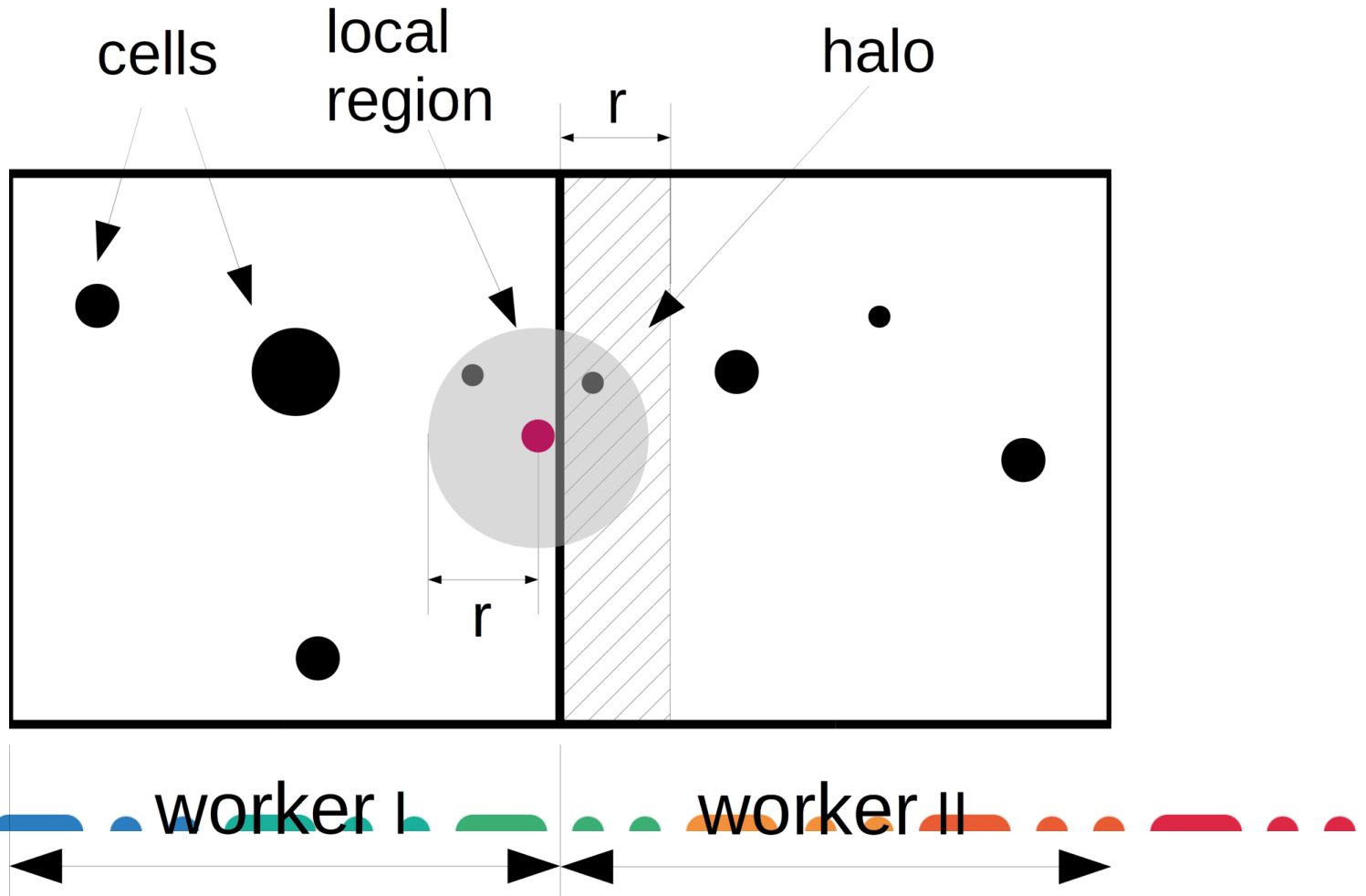
Domain-Decomposition



Distributed Runtime



Border Region



Outline

1. Biological Simulation Basics

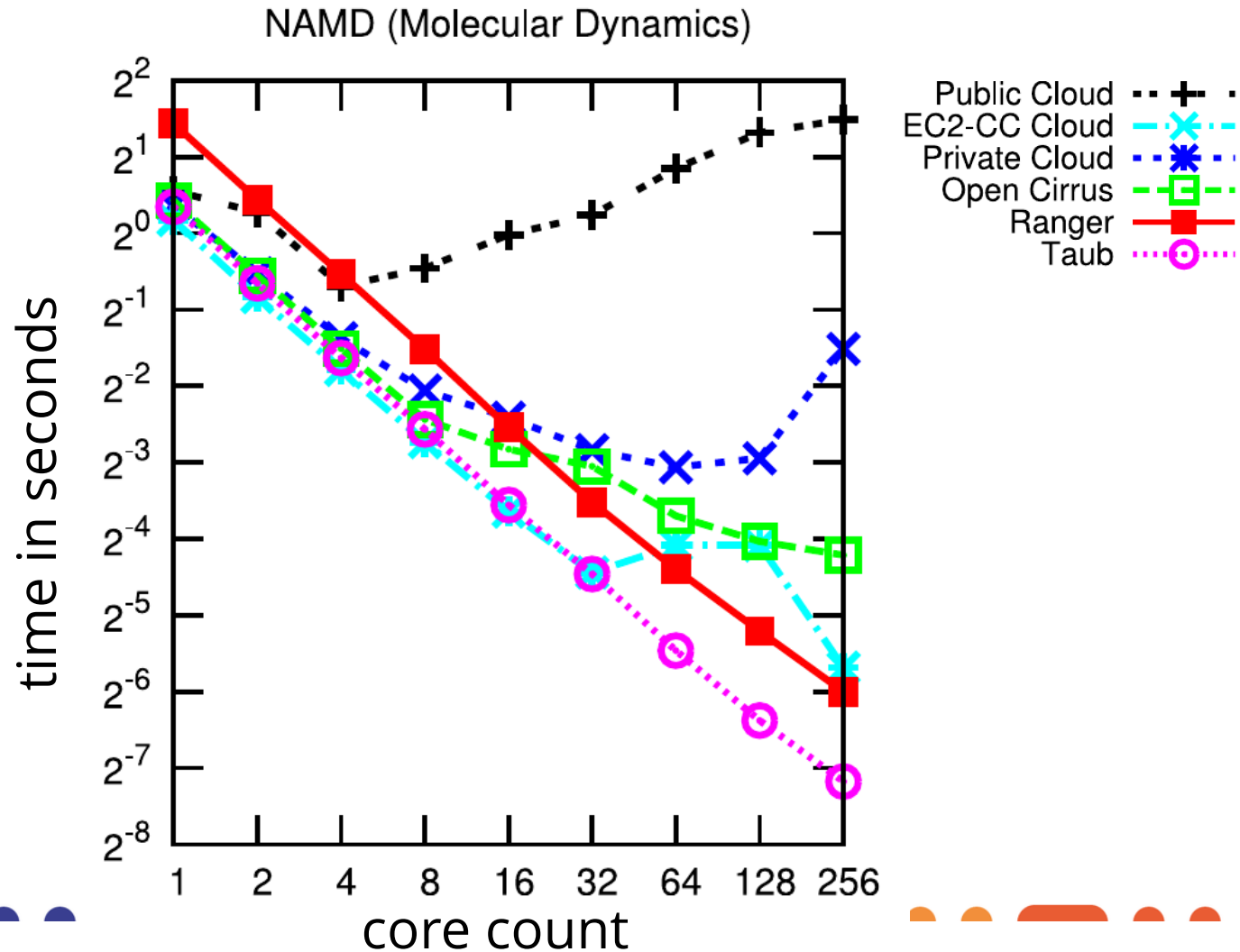
2. Distributed Runtime

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NAMD Scaling



Performance Issues in the Cloud

- Poor network performance compared to supercomputers
- Virtualization
- Resource contention with other tenants

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1. Biological Simulation Basics

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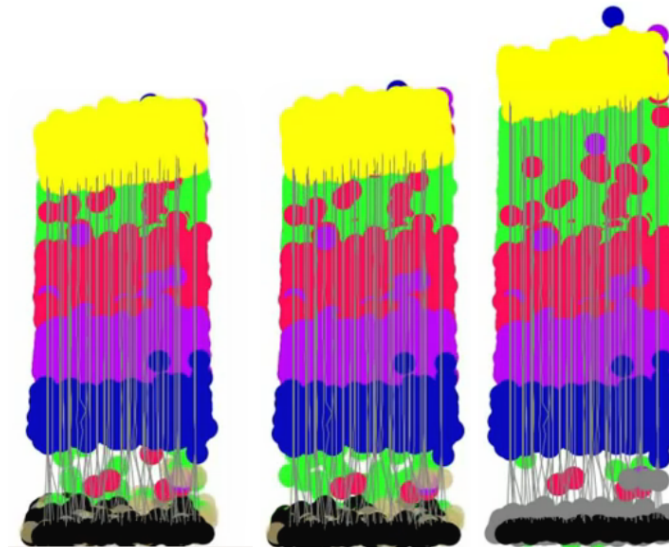
Key Observations

- Some regions and data members are static
- Changes are incremental
- Values might be predicted
- Communication can be replaced with additional computation

Static Regions

- **Definition:** simulation objects whose values do not change along the time dimension

Example: growth of the cerebral cortex



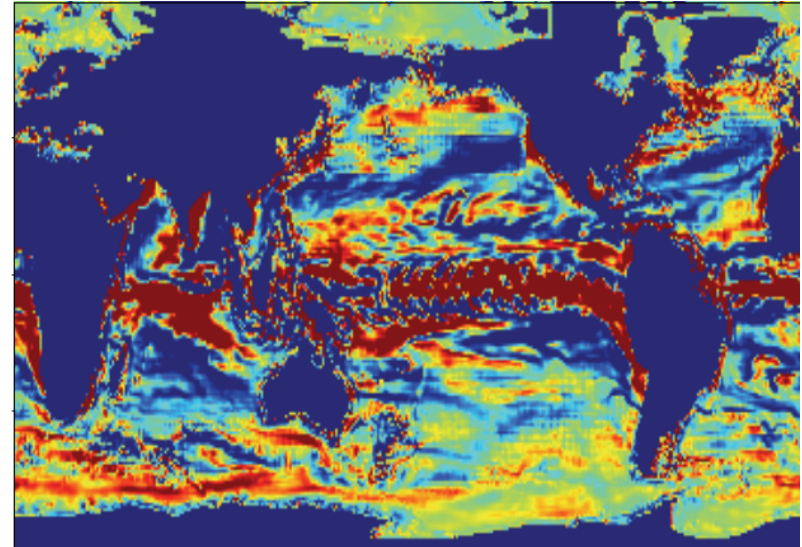
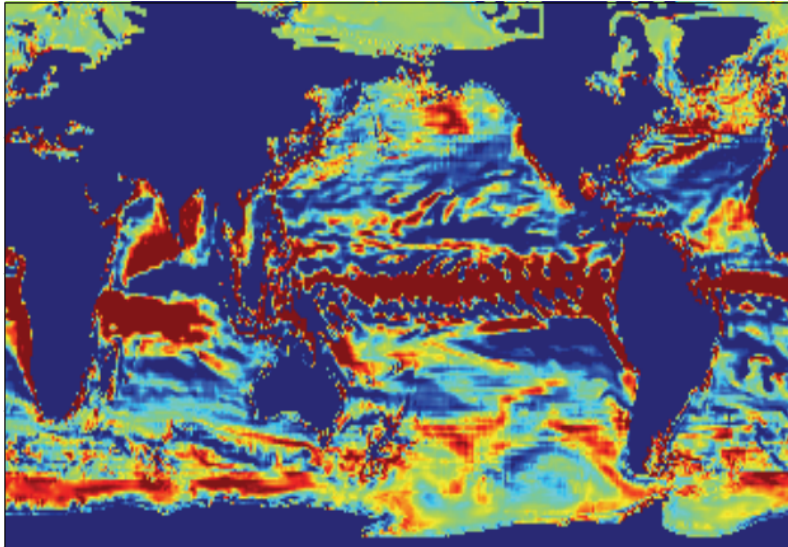
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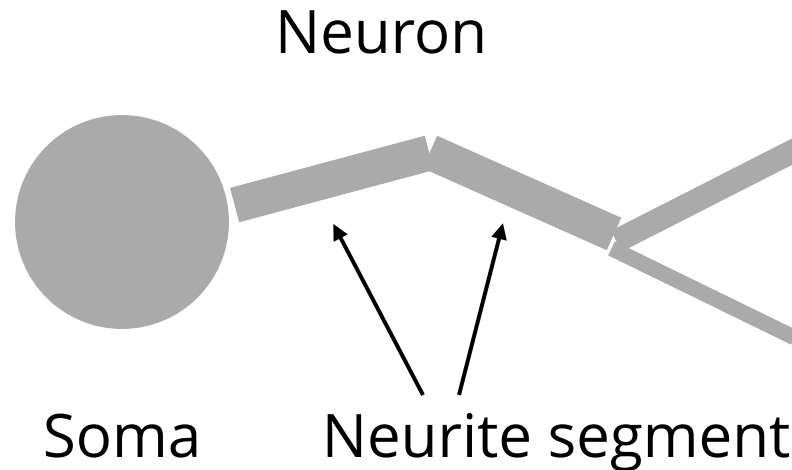
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Static Regions

- Climate Simulation: continental data does not change during simulation of the ocean surface



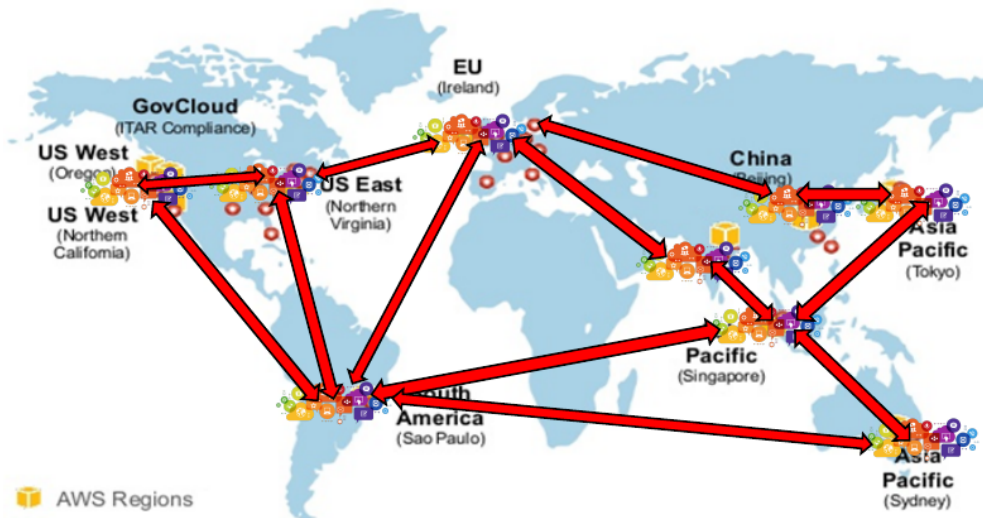
Static Data Members



```
class NeuriteSegment {  
    std::array<double, 3> position_;  
    NeuriteSegment* parent_;  
    ...  
};
```

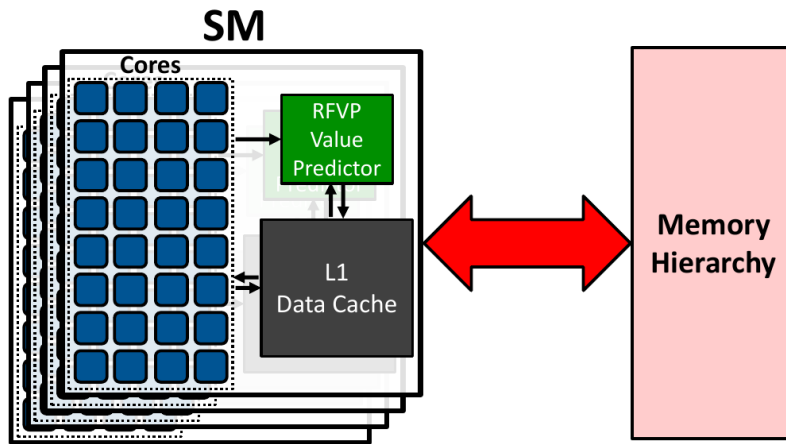
Incremental Changes

- **Observation:** Change between two time steps might be insignificant
- **Idea:** Communicate only significant updates
- **Example:** GAIA: Geo-distributed machine learning approach



Value Prediction

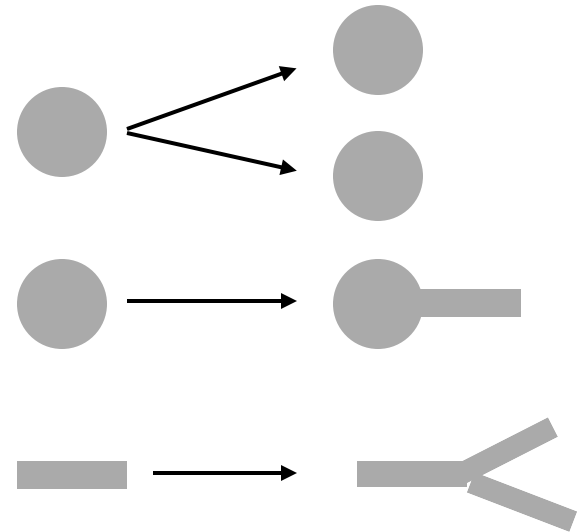
- **Idea:** Predict values for safe-to-approximate variables
- Inspiration from computer architecture:
"Rollback free value prediction"



```
class NeuriteSegment {  
    std::array<double, 3> position_  
    NeuriteSegment* parent_  
    ...  
};
```

Computation vs Communication

- **Idea:** Recompute certain events on the destination server instead of transferring the results
- **Examples:**
 - Cell division
 - Neurite
 - extension
 - bifurcation
 - ...



Send event descriptor instead of whole new simulation object

Next Steps

- Develop the distributed runtime
- Verify that network properties are a bottleneck in the cloud
- Provide detailed analysis for different approaches
- Develop new ideas



QUESTIONS?

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