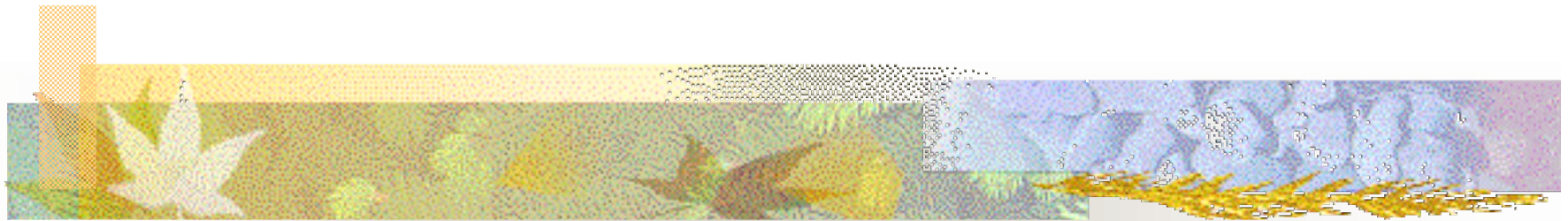


A Stochastic Simulation of Natural Organic Matter and Microbes in the Environment



Xiaorong Xiang

Gregory Madey

Yingping Huang

Steve Cabaniss

(University of New Mexico)

Department of Computer Science and Engineering

University of Notre Dame

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Objectives

- New approach for NOM modeling
 - Agent-based modeling
- E-Science on the Web
- Intelligent interface
- The NOM Collaboratory



Outline

- **Introduction**
- **Modeling**
- **Core simulation engine**
- **Intelligent Web-based interface**
- **The NOM collaboratory**
- **Conclusion**
- **Future work**



Introduction

- What is Natural Organic Matter (NOM)?
- Role of NOM in various science disciplines
 - Mobility and transport of pollutants
 - Availability of nutrients for microorganisms and plant communities
 - Affects quality of drinking water
- Need to understand the evolution and heterogeneous structure of NOM

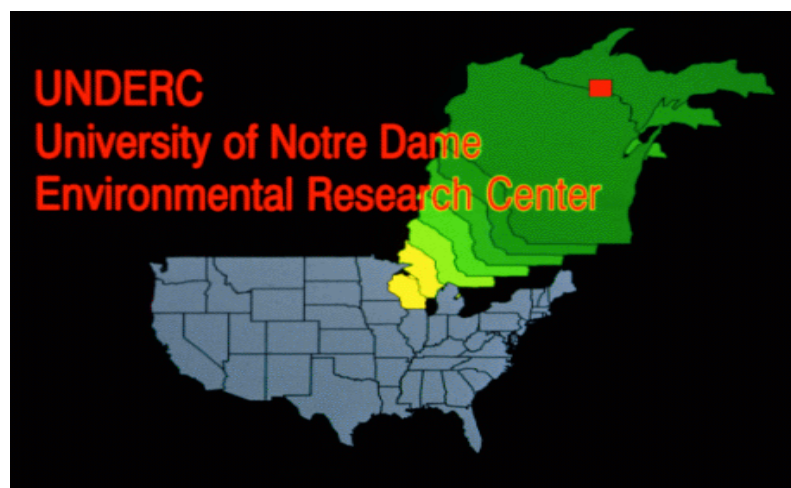
Forest Service Bog [DOC] 7 MW 2200



Twomile Creek [DOC] 17 MW 1500



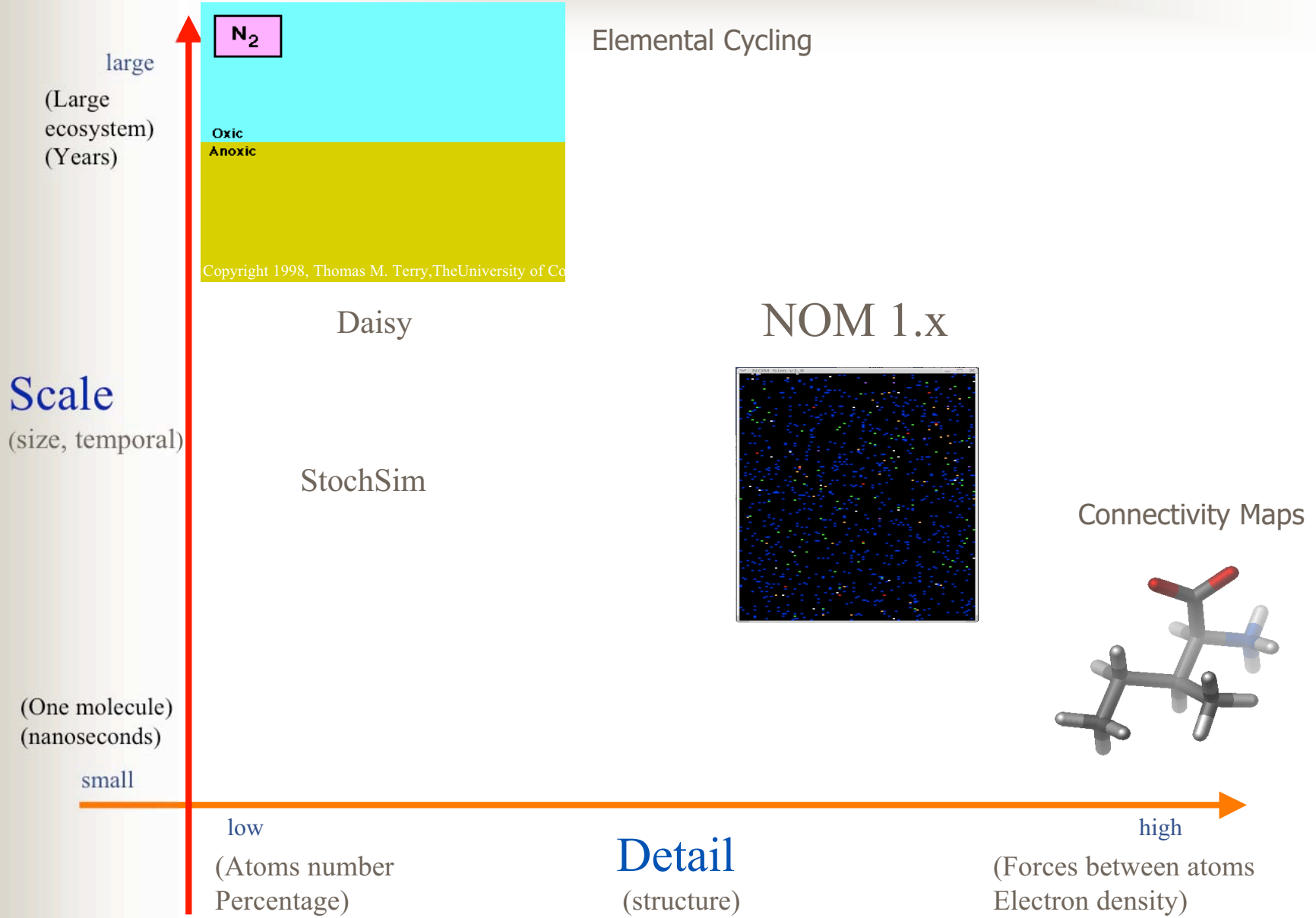
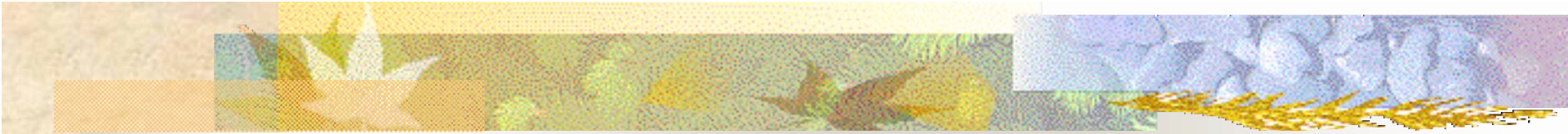
Nelson Creek [DOC] 79 MW 900





Previous models

- Two examples:
 - Daisy (S. Hansen, H. E. Jensen, and N. E. Nielsen 1990-present) : a soil plant atmosphere system model
 - StochSim (C. J. Morton-Firth 1998-present): Stochastic simulation of cell signaling pathways





Our model

- Agent-based modeling (Individual-based modeling)
 - Agent-based model
 - Reynolds (1987): Flocks, herds, and schools: A distributed behavioral model. *Computer Graphics*
 - Each molecule as an individual object with spatial properties
 - Bottom-up approach
 - Stochastic model
 - Trace changes of the system → Database and data mining technologies



Our model (cont.)

- Web-based simulation model
 - Serve as an example for E-Science
 - G. Fox (2002): E-science meets computational science and information technology. *Computing & Engineering*
 - R. M. Jakobovits, J. F. Brinkley, C. Rosse, and E. Weinberger (1998): Enabling clinicians, researchers, and educators to build custom Web-based biomedical information system
 - Support the collaborations, data and information sharing between scientists
 - No installation, expensive computation resources needed by scientists



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Modeling

- A complex system
 - Consists of a large number of objects
 - Molecules, Microbes
 - **Heterogeneous properties**
 - **Individual behaviors**
 - **Interaction between each other**
 - **Objects behavior and interaction are stochastically determined by:**
 - **Attributes** (intrinsic parameters)
 - **Reactions rates**
 - **Environment condition** (extrinsic parameters)
 - No central control
 - Emergent properties



Modeling (cont.)

■ Agent Attributes

- More specific than “percent carbon” but less detailed than a molecular connectivity map
- Elemental composition
 - (C, H, O, N, S, P)
- Functional groups
 - (double bonds, ring structure, alcohols ...)
- The origin of objects
 - spatial position in the system
 - Precursor type of molecule
- Probability table of physical and chemical reactions
- Molecule weight



Modeling (cont.)

- Agent Behaviors (reactions and processes):
 - Transport through soil pores by water (spatial mobility)
 - Adsorb onto or desorbed from mineral surfaces
 - Chemical reactions
 - Total 10 types in current model
 - First order
 - Second order
 - Stochastically determined
- Space:
 - 2-D grid



Modeling(cont.)

- Environmental parameters
 - Temperature
 - pH
 - Light intensity
 - Microbe density
 - Water flow rate
 - ...



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Core simulation engine

- Implementation
 - Swarm toolkit (Santa Fe Institute)
 - Java programming language (Sun JDK 1.4.1_01)
- GUI version
 - View the animation of molecules
 - Easy for debugging and modeling
- Web-based version

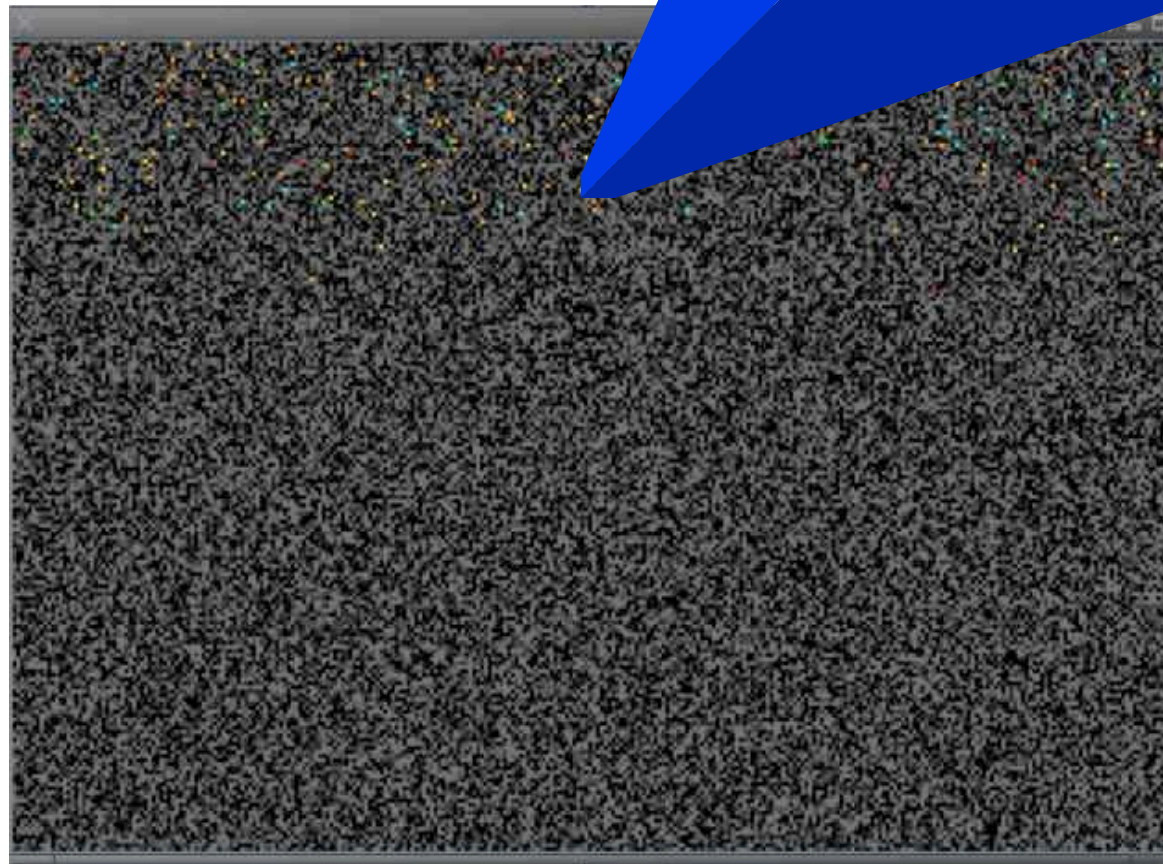


Core simulation engine (cont.)

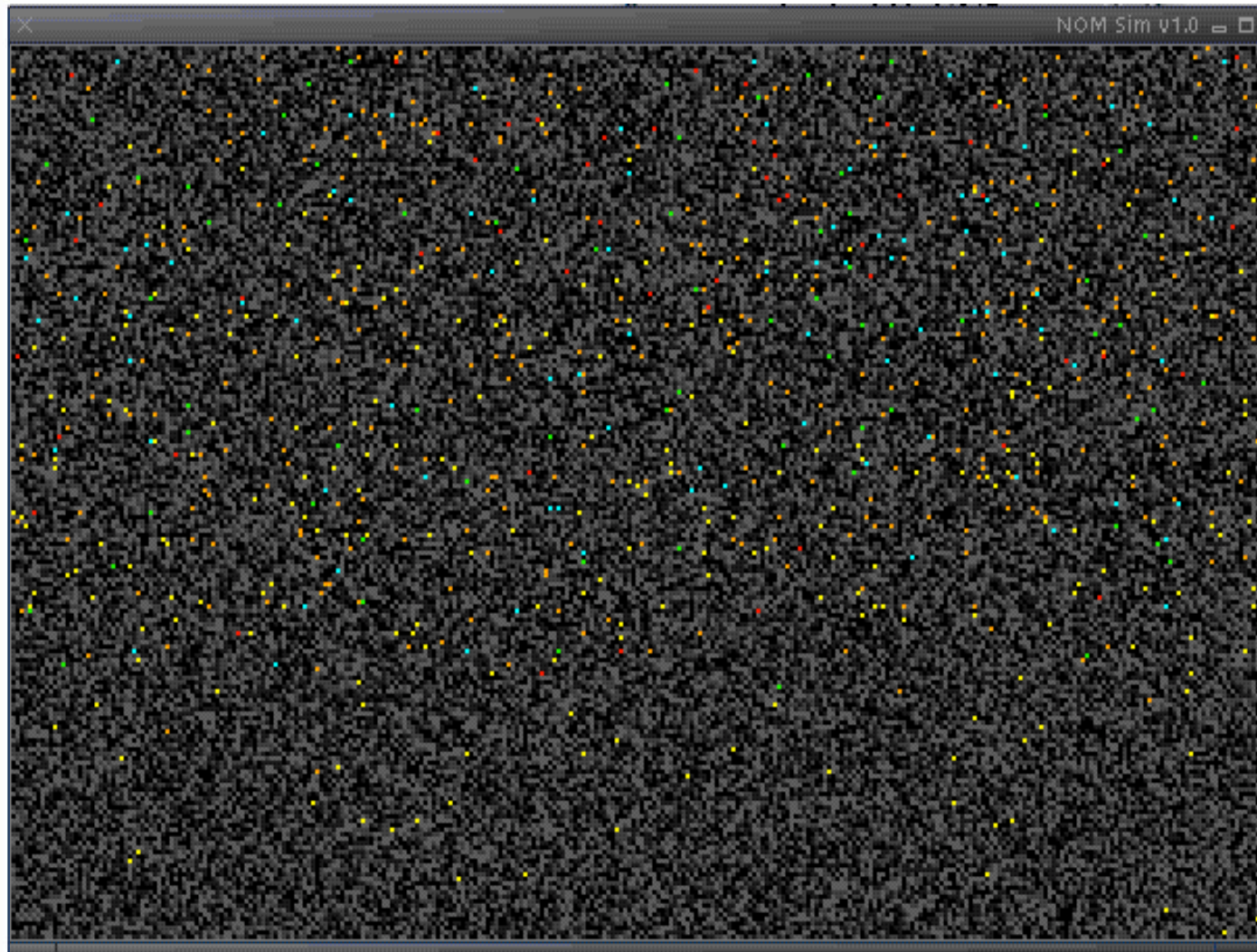
- Read simulation parameter from the database (JDBC)
 - Environmental parameters (pH, temperature, light intensity, and so on)
 - Molecule types and distributions
- User defined time has been separated to a large number of equal size time steps
- Write relevant data into the database every time step (JDBC)
 - Trace the dynamic properties of individuals and the system over time

Visualization

Black - No Adsorption
Grays - Levels of Adsorption
Red - Lignins
Green - Cellulose
Blue - Proteins
Yellow - Reacted
Orange - Adsorbed



Visualizations — NOM molecules in solution and adsorption





Outline

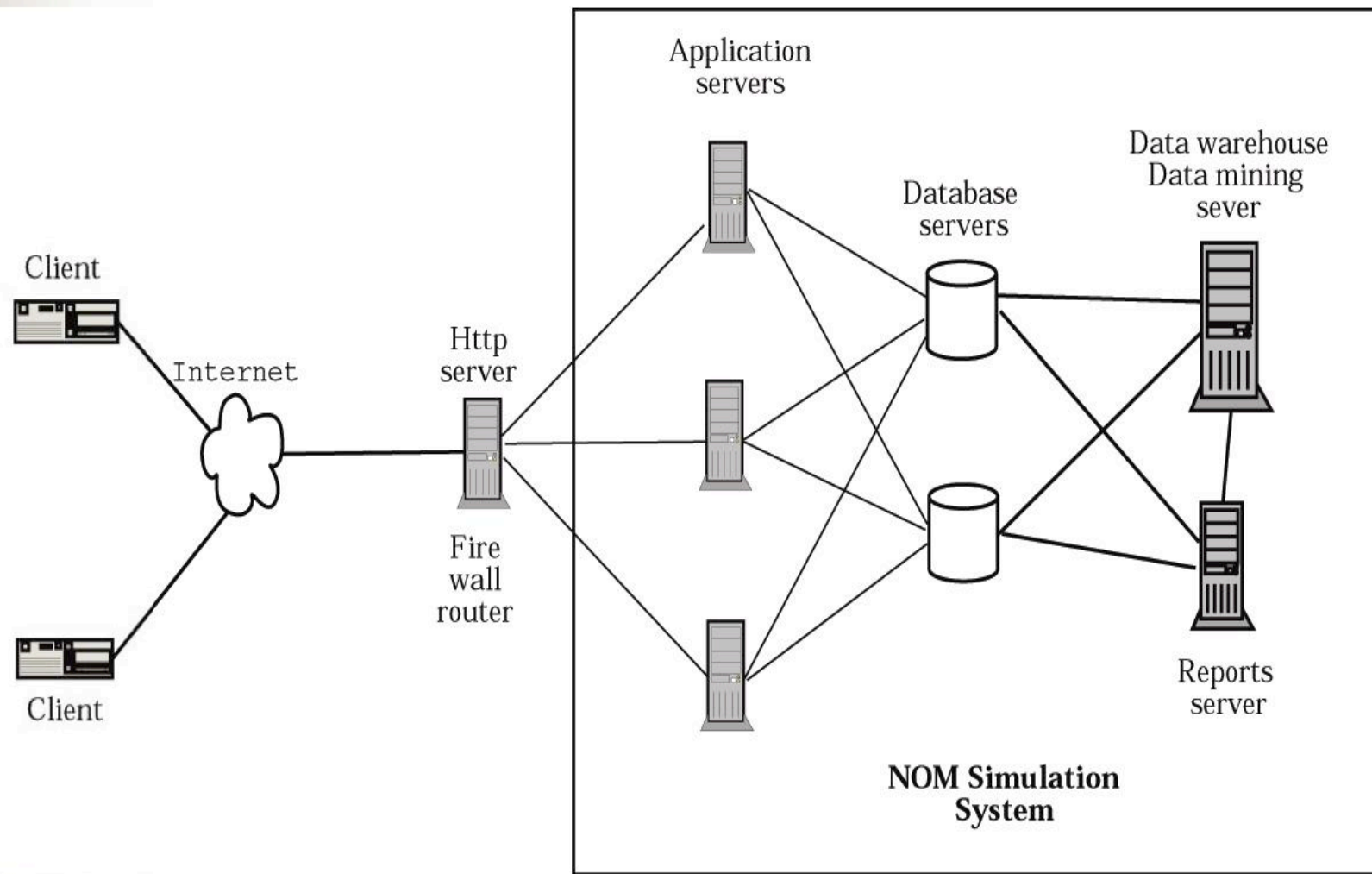
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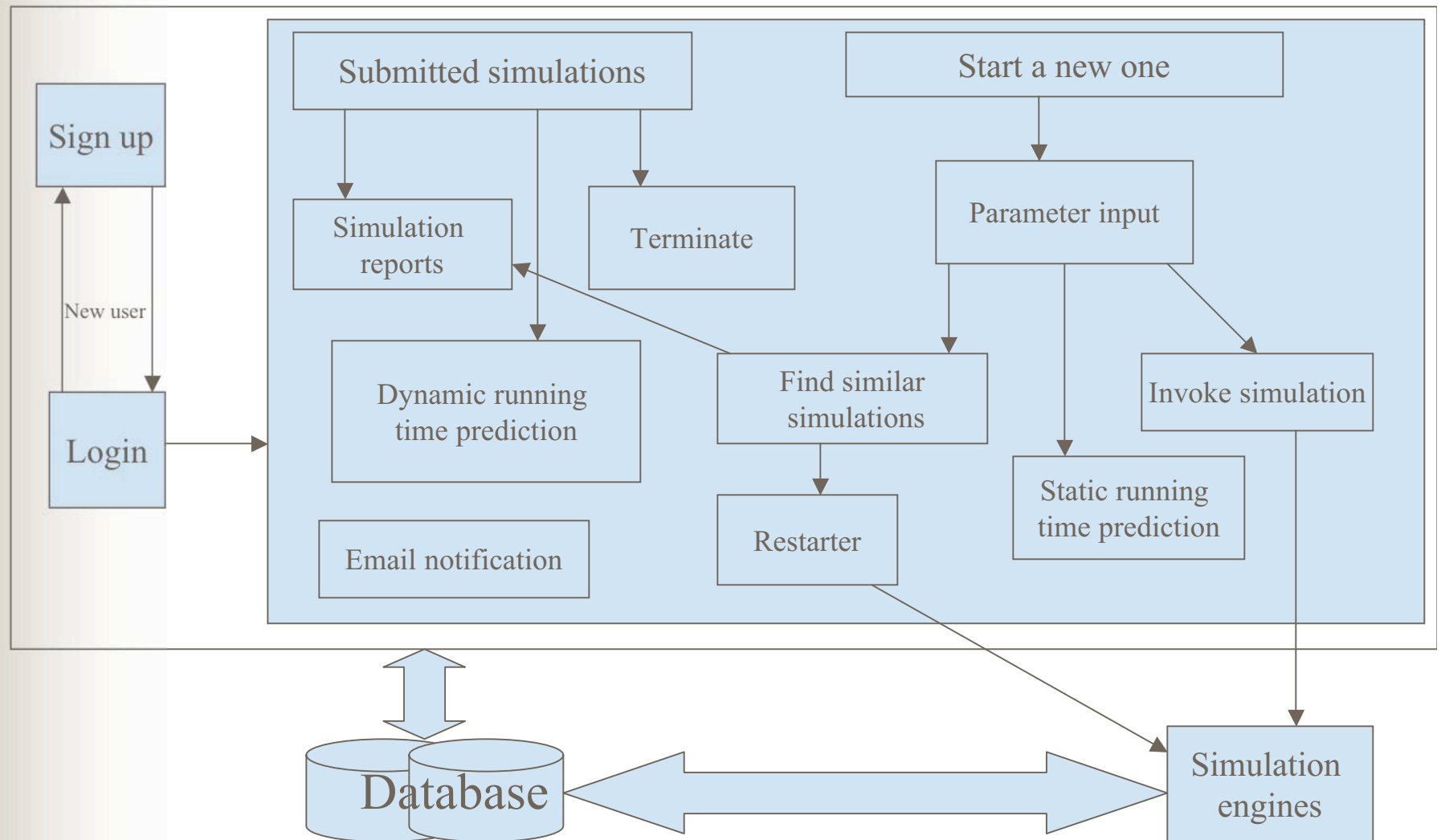
Web-based model

- Distributed, Web-based scientific application model
- Based on Sun Java 2 Enterprise Edition (J2EE)
 - Standard HTML Forms / JSP
 - Java Servlets, Java Beans
 - JDBC - Oracle
 - Oracle Database, Oracle Forms, and Reports
- Three parts:
 - Intelligent Web-based interface
 - Core simulation engine
 - Data analysis packages, Data mining technologies

Access NOM simulation from Web



Web-based interface



Example of Interface



Welcome to NOM Research Group!

Leilani Arthurs

(* Required fields)

Molecule Name* 3ficFA	<input type="checkbox"/> Make it available to public				
Atoms of the molecule Each field must be a non-negative integer. Default value is 0.	(Atom) C 66	(Atom) H 64	(Atom) N 0		
	(Atom) O 38	(Atom) S 0	(Atom) P 0		
Functional groups of the molecule Each field must be a non-negative integer. Default value is 0.	DoubleBond 0	Rings 0	Phenyl 0	Alcohols 0	Phenols 0
	Ethers 0	Esters 0	Ketones 0	Aldehydes 0	Acids 0
	Arylacids 0	Amines 0	RingN 0	Amides 0	Thioethers 0
	Thiols 0	Phosphoesters 0	HPhosphoesters 0	Phosphates 0	

Create New Molecule Type

Report example

Please provide the following information.

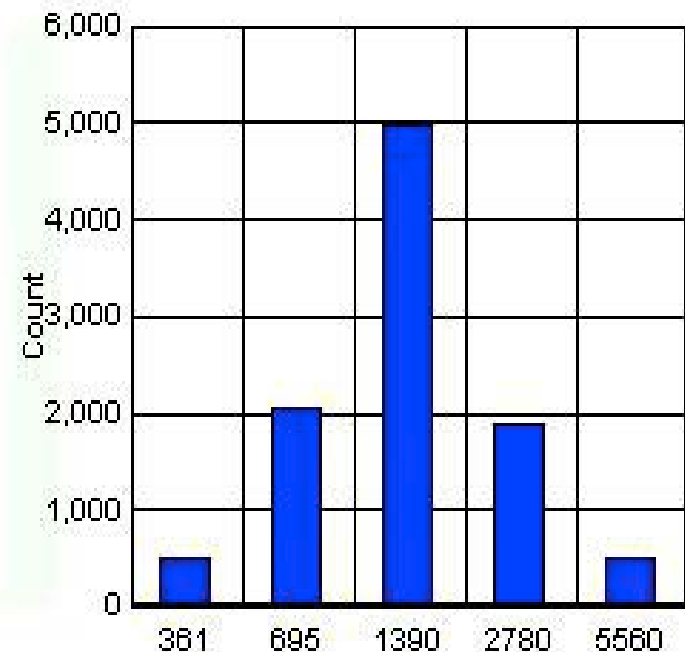
Current Time Step is: 1969 [Refresh](#)

Time Step:
Status:
Start Y:
End Y:

Generate Reports

Generate XML file

Weight Distribution





Intelligent interface components

- Email notification
- Running time prediction
 - Static
 - number of molecules
 - number of time steps
 - Dynamic
 - current time step
 - current wall clock time



Intelligent interface components (cont.)

- Find similar simulations
 - Environment parameters
 - Molecule types and distributions
 - Retrieve the data sets from database
 - Points on a high dimension space
 - Euclidean distance
 - Ordered list
 - Review the simulation results or restart



Intelligent interface components (cont.)

- Automatic restarter
 - Save the state of each objects in the system to database every check point
 - Load the state to the core simulation engine



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Previous work

- Combination of words “collaboration” and “laboratory” first coined by William Wulf (1996):
Richard T. Kouzes, James D. Myers, and William A. Wulf. Collaboratories: Doing science on the internet. *IEEE Computer*, 1996
- Diesel Collaboratory: C. M. Pancerella, L. A. Rahn, and C. L. Yang: The diesel combustion collaboratory: combustion researchers collaborating over the internet. In *Proceedings of the 1999 ACM/IEEE conference on Supercomputing*
- BioCoRE: <http://ks.uiuc.edu/Research/biocore>
- EMSL Collaboratory: <http://www.emsl.pnl.gov:2080/docs/collab>



The NOM Collaboratory









- Interdisciplinary project
 - Chemists
 - Biologists
 - Ecologists
 - Computer Scientists
- Build and integrate software using J2EE
 - NOM modeling & simulation software
 - XML-based standard data format definitions
 - Data querying and manipulation tools
 - Electronic communication tools

NOM Collaboratory - Mozilla

File Edit View Go Bookmarks Tools Window Help

Back Forward Reload Stop http://localhost:8888/interface_COLLABORATION/Collaboratory.html Search Print

NOM Collaboratory

 <p>NOM Simulator</p>	<p>Provide an intelligent interface to facilitate using the NOM simulator</p>	<p>Provide an Interface to define new molecule type</p>	 <p>Molecule Editor</p>
 <p>Search Engine</p>	<p>Provide an interface to search simulation information</p>	<p>Provide an administration role to validate the newly added molecule for public usage</p>	 <p>Molecule Validator</p>
 <p>Discussion Board</p>	<p>Provide a threaded discussion board</p>	<p>Provide a XML-based Markup Language definition</p>	 <p>NOML</p>
 <p>Chat Room</p>	<p>Provide a real time chat box</p>	<p>Provide an Interface to upload publications</p>	 <p>File Upload</p>

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XML-based NOM Markup Language (NOML)

- NOML:
 - Standard data format
 - Environment.dtd, Molecules.dtd, Setup.dtd
- Facilitate communication
 - User ===== User
 - Application ===== Application
- Extension



Other tools

- Molecule editor
 - Define new molecule type
- Molecule validator
 - Authorized persons (Chemists) to validate data
 - Share the molecule type
- Search engine
 - Ad-hoc query
 - View results of the completed simulations
 - Restart some simulations



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Conclusion

- Agent-based stochastic model for simulating the NOM evolution with discrete temporal and spatial properties
- A Web-based simulation architecture (multiple simulation servers, database servers, and data mining technologies)
- Database technologies
- A Web-based intelligent configuration interface
- The NOM collaboratory



Outline

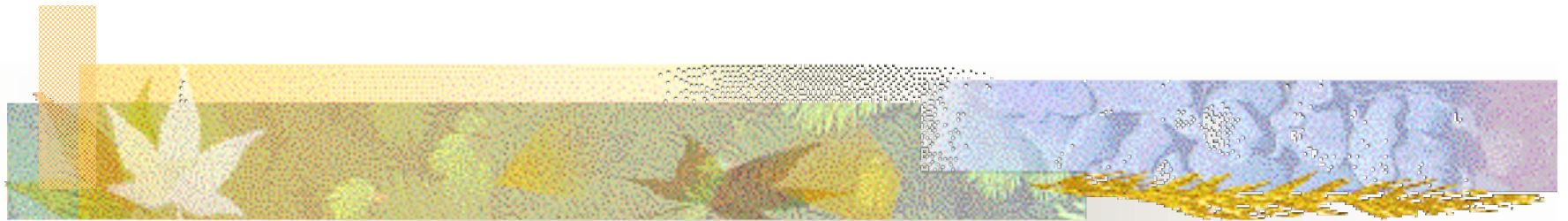
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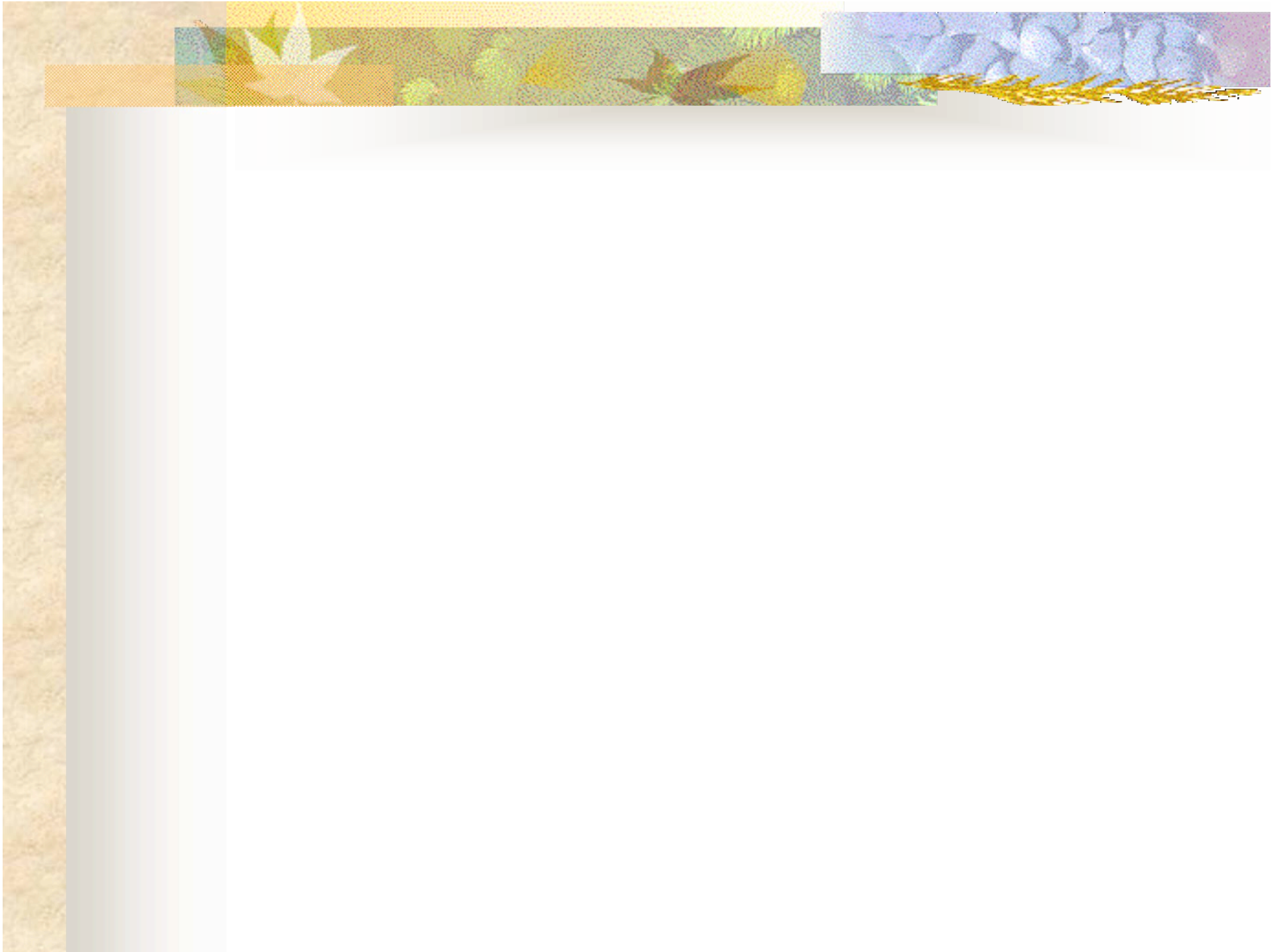
Future work

- Model testing (this summer)
 - Testing of the sorption process
 - More features need to be added into the core simulation engine
- Model validation (this summer)
- Collaboratory:
 - More communication tools
 - More simulation models for NOM study
 - NOML extension

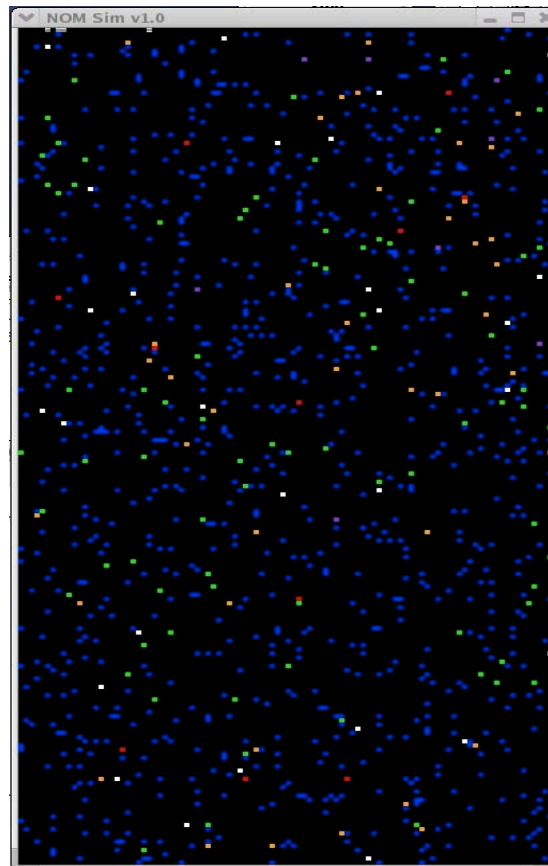
Thank You !



Questions?



GUI version



Work flow

