



The nanoHUB: A science Gateway for nanotechnology

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Online simulations and MORE



nanoHUB.org



An Ecosystem

Applications



learning modules



nanoHUB.org



A middleware

letwork for Computational Nanotechnolog

Resources

TERAGRID





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Why Middleware?





nanoHUB Architecture

Remote access Purdue to simulators and R S compute power Web Presence Physical Machine UNIVERSITY OF **FI ORIDA** Virtual Machine Appl Middleware **WISCONSIN Resource Mgmt** Cluster Tue Jun 14 17:03:00 GMT 200 TERAGRII. **Open Science Grid**

letwork for Computational Nanotechnology



- Fundamental goal of Grid computing:
 - "Flexible, secure, coordinated resource sharing among dynamic collections of individuals, institutions, and resources" [Foster et. al]
 - Without forcing the Grid users to modify their applications to make them Grid-aware
- Key challenge to Grid middleware:
 - The provisioning of execution environments that have flexible, customizable configurations and allow for secure execution of untrusted code from Grid users

"The Anatomy of the Grid: Enabling Scalable Virtual Organizations", I. Foster, C. Kesselman, S. Tuecke. International J. Supercomputer Applications, 15(3), 2001





Problems with using physical machines







Our approach: Define once, instantiate ondemand





Accomplishments



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Application Deployment





Summary

• Scripting languages as a tool development philosophy



Build interfaces on scripting foundation

Build tools from component parts

Add scripting language interface to each component

Components coded in C, C++, Fortran



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Wrap Existing Tools







nanoHUB and In-VIGO integration







nanoHUB and In-VIGO integration

Two virtual machine systems:

VMware:

- + Commercial product
- + Mature software
- not open source
- smaller community

User Mode Linux:

- + Open Source (hack the code)
- + Large community of developers
- not as mature

Future with Xen:

- + Open Source
- + Better performance
- +Collaboration with ANL





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NEMO3D on TeraGrid



Accomplishments:

NCN community allocation NEMO3D ported to TeraGrid Scaling study 64 Million atoms simulation ~50,000 SUs used on TeraGrid



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BioMOCA on TeraGrid

Computational challenge:

The "window" of observation is on the order of 100 to 1,000 nanoseconds for porin, and from 1,000 to 10,000 nanoseconds for gramicidin. $\Delta t=1 fs$

The computational requirement for one data point on the current-voltage curve in porin is estimated between 900 to 6,000 hours of single processor time (Xeon 2.4 GHz, 2GB RAM).

Accomplishments:

NCN student on TeraGrid (Gulzar Kathawala)

MOCA and BioMOCA ported to TeraGrid

~75,000 SUs used on TeraGrid

Permeation trajectory of a "lucky" Na⁺ in a gramicidin A ionic channel T. Van der Straaten, G. Kathawala, U. Ravaioli (UIUC)





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Molecular Conduction Learning Module



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Molecular Conduction Learning Module



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Molecular Conduction Learning Module

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Molecular Conduction Learning Module

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Leveraging Educational Standards





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Example of content interoperability





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Users ?





Questions?

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