

VMD+OptiX: Bringing Interactive Molecular Ray Tracing from Remote GPU Clusters to your VR Headset

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<http://www.ks.uiuc.edu/>

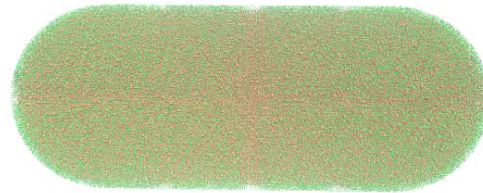
3:30pm, Wednesday Nov 18, 2015

NVIDIA GTC Theater, NVIDIA Booth #1021,
Supercomputing 2015, Austin, TX

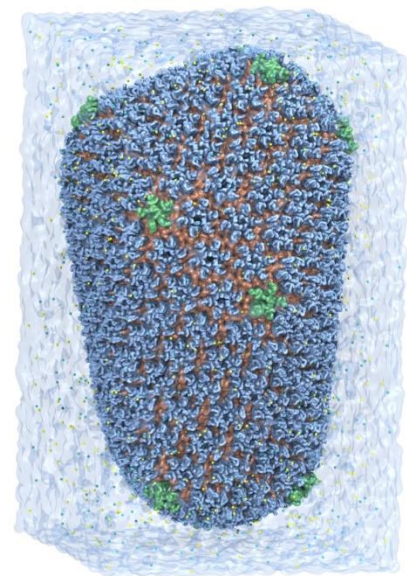


VMD – “Visual Molecular Dynamics”

- Visualization and analysis of:
 - molecular dynamics simulations
 - particle systems and whole cells
 - cryoEM densities, volumetric data
 - quantum chemistry calculations
 - sequence information
- User extensible w/ scripting and plugins
- <http://www.ks.uiuc.edu/Research/vmd/>



Whole Cell Simulation

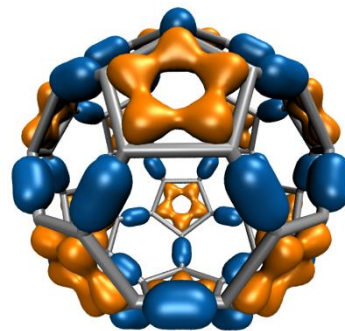


MD Simulations

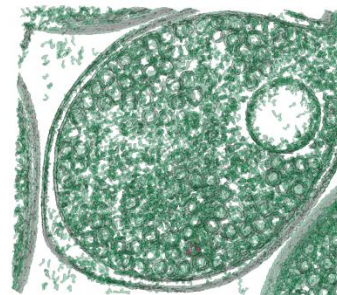
Structural Similarity	
1trp-a	ASFS...EAP...G...D...V...E...K...K...K...I...T...V...O...K...C...A...Q...C...H
1ocr-a	ASFS...EAP...G...D...V...E...K...K...K...I...T...V...O...K...C...A...Q...C...H
1yaa-a	AKESTGFK...P...G...S...A...K...K...G...A...T...L...F...K...T...R...C...Q...Q...C...H
1scya-a	AKESTGFK...P...G...D...V...A...K...G...K...K...T...F...V...O...K...C...A...Q...C...H
1tocy-a	AKESTGFK...P...G...D...V...A...K...G...K...K...T...F...V...O...K...C...A...Q...C...H
1trp-a	AKESTGFK...P...G...D...V...E...K...K...K...I...T...V...O...K...C...A...Q...C...H

Sequence Similarity	
1trp-a	ASFS...EAP...G...D...V...E...K...K...K...I...T...V...O...K...A...Q...C...H
1ocr-a	ASFS...EAP...G...D...V...E...K...K...K...I...T...V...O...K...A...Q...C...H
1yaa-a	AKESTGFK...P...G...S...A...K...K...G...A...T...L...F...K...T...R...Q...Q...C...H
1scya-a	AKESTGFK...P...G...S...A...K...K...G...A...T...L...F...K...T...R...Q...Q...C...H
1tocy-a	AKESTGFK...P...G...S...A...K...K...G...A...T...L...F...K...T...R...Q...Q...C...H

Sequence Data



Quantum Chemistry

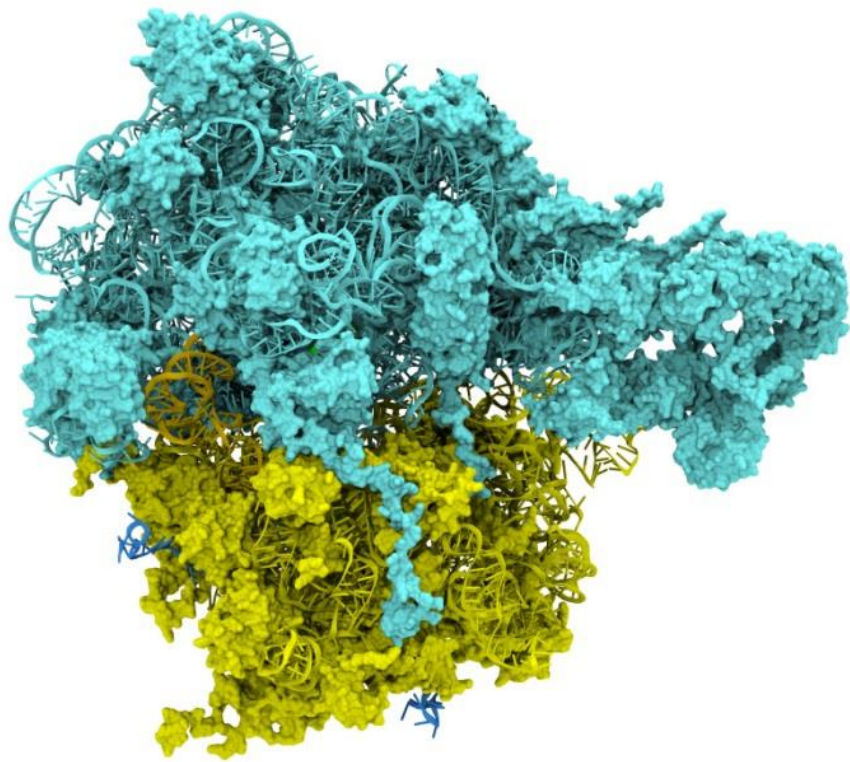


CryoEM, Cellular Tomography

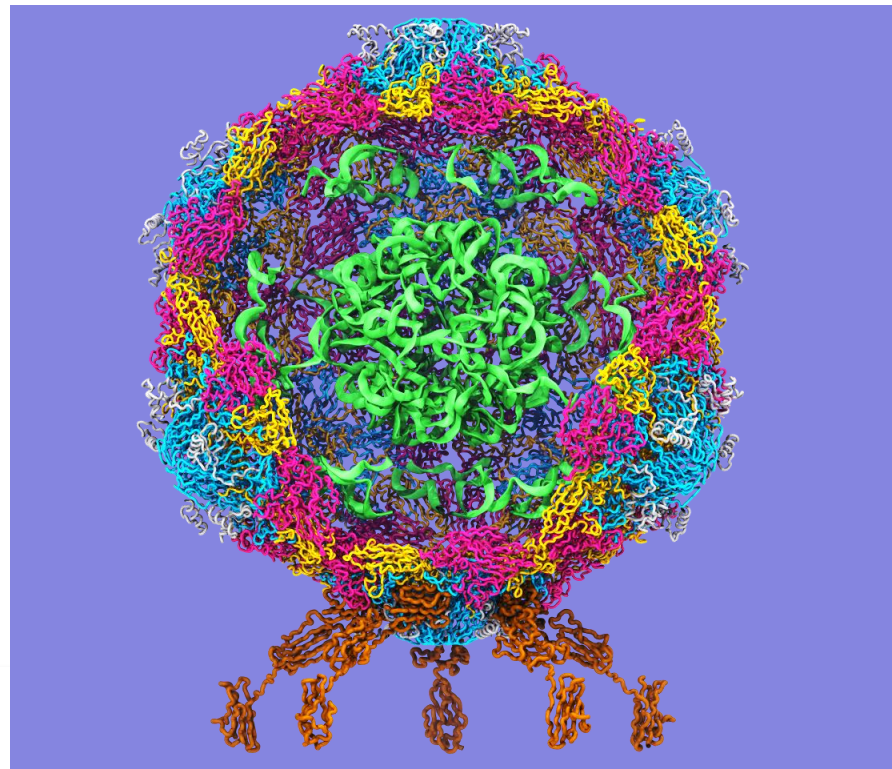
Goal: A Computational Microscope

Study the molecular machines in living cells

Ribosome: target for antibiotics

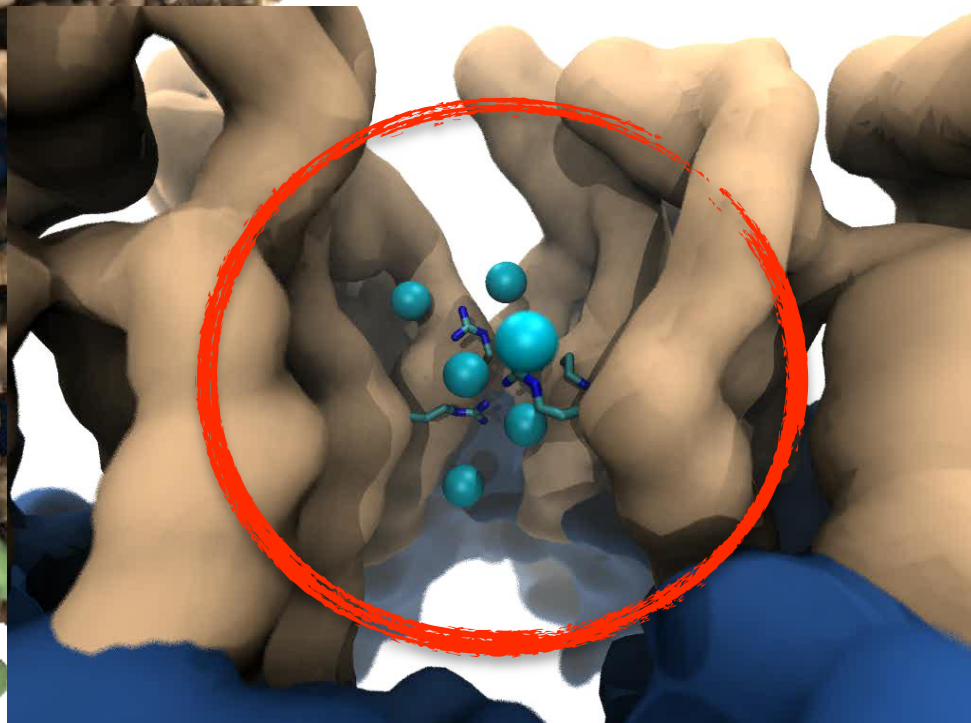
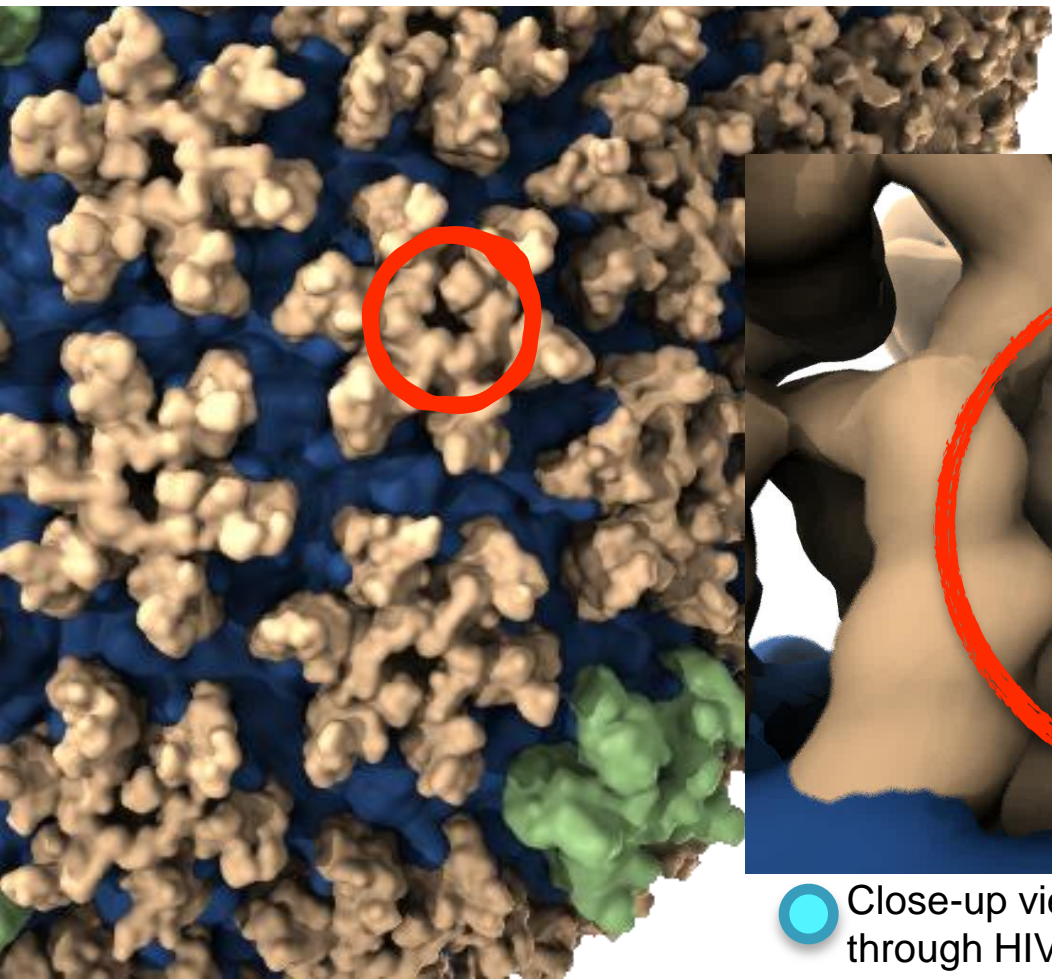



Poliovirus



Goal: Intuitive interactive viz. in crowded molecular complexes

Results from 64 M atom, 1 μ s sim!



 Close-up view of chloride ions permeating through HIV-1 capsid hexameric centers

Immersive Viz. w/ VMD

- VMD began as a CAVE app (1993)
- Use of immersive viz by molecular scientists limited due to cost, complexity, lack of local availability, convenience
- Commoditization of HMDs excellent opportunity to overcome cost/availability
- This leaves many challenges still to solve:
 - Incorporate support for remote visualization
 - UIs, multi-user collaboration/interaction
 - Rendering perf for large molecular systems
 - Accommodating limitations/idiosyncracies of commercial HMDs



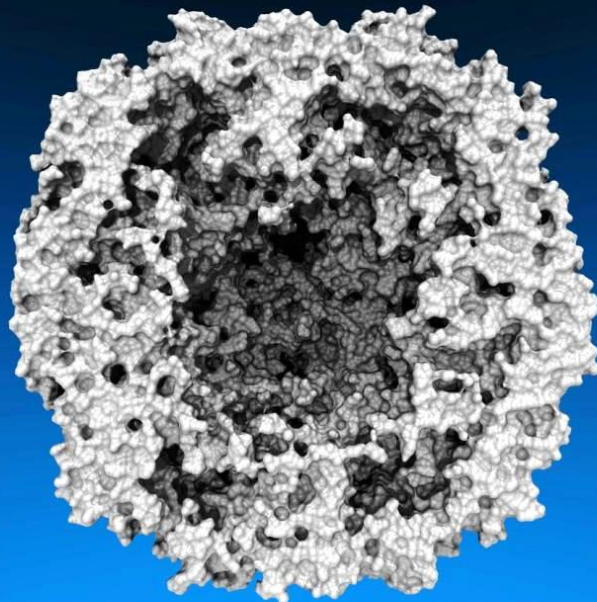
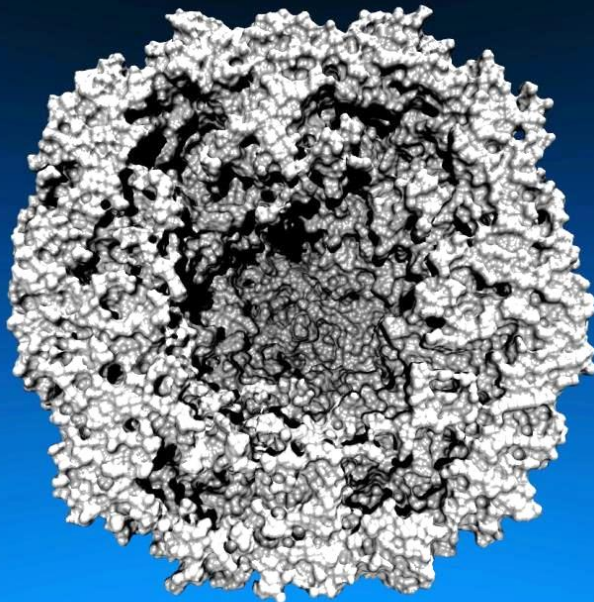
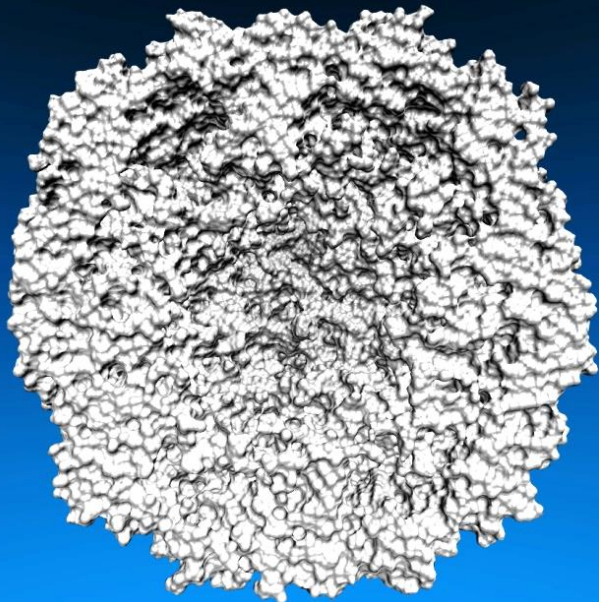
VMD running in a CAVE w/ VR Juggler

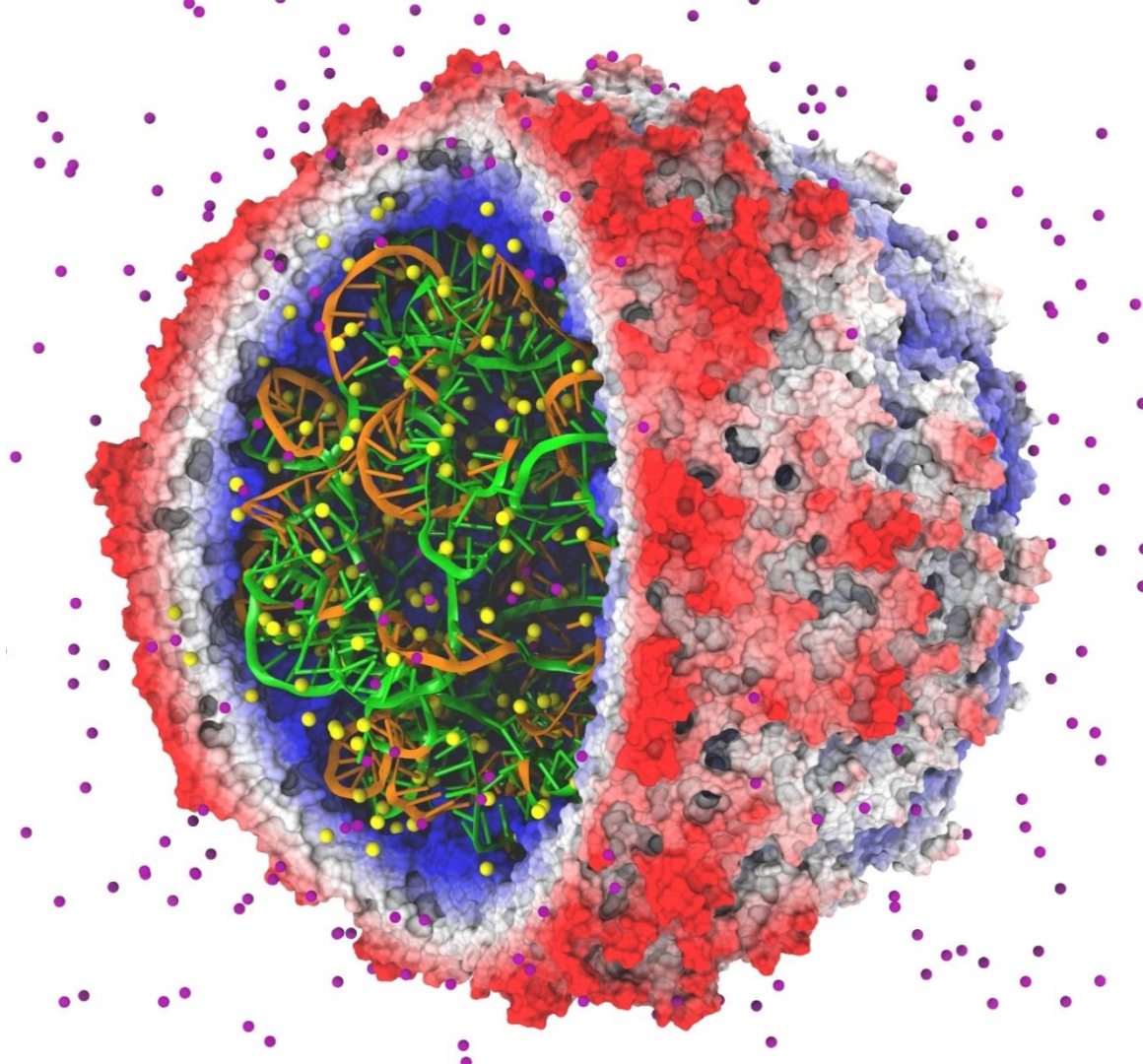
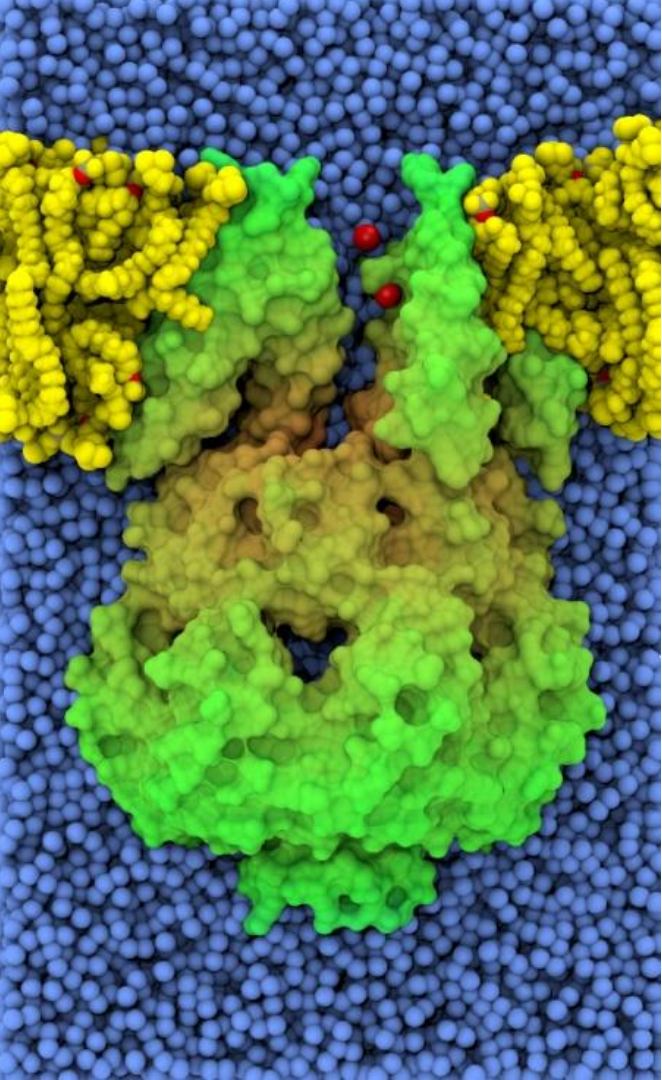
Lighting Comparison

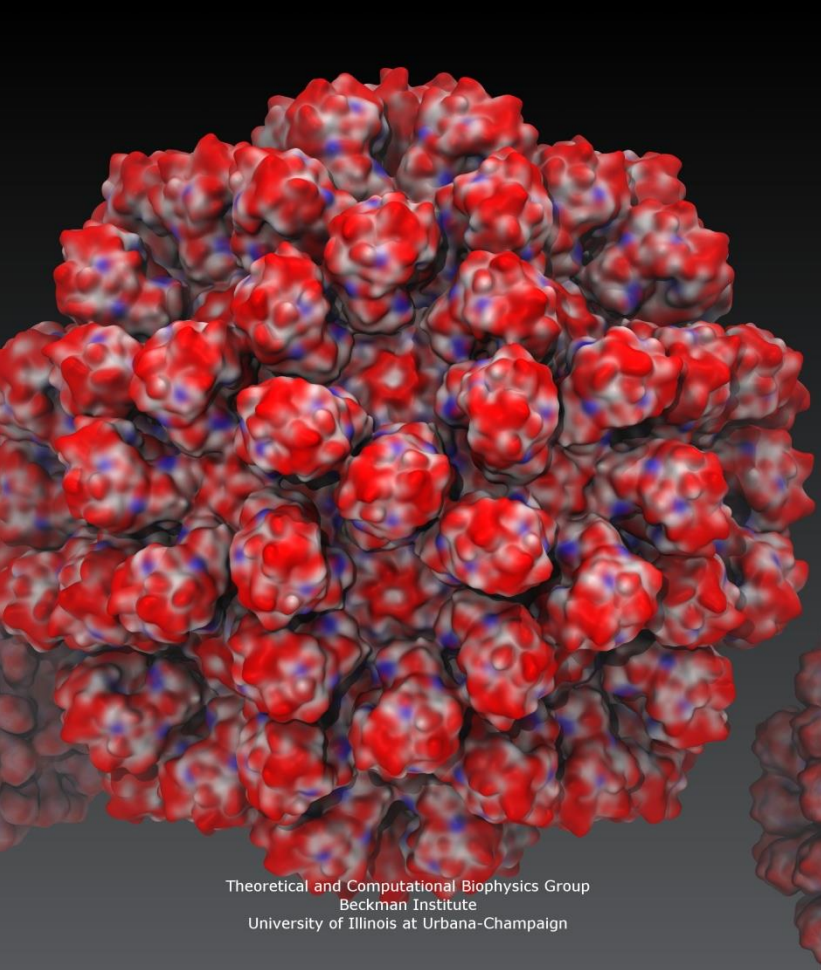
Two lights, no shadows

Two lights, hard shadows, 1 shadow ray per light

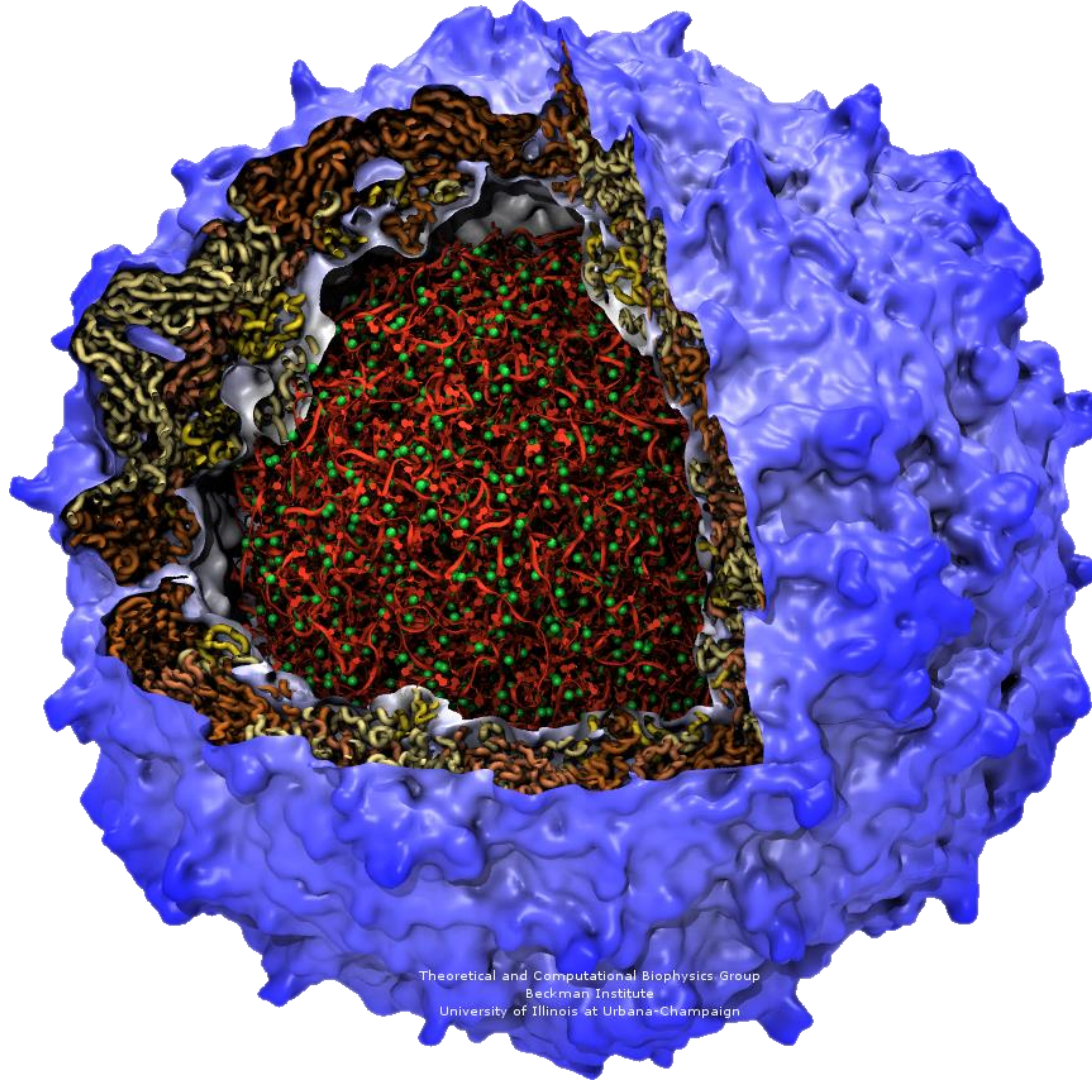
Ambient occlusion + two lights, 144 AO rays/hit





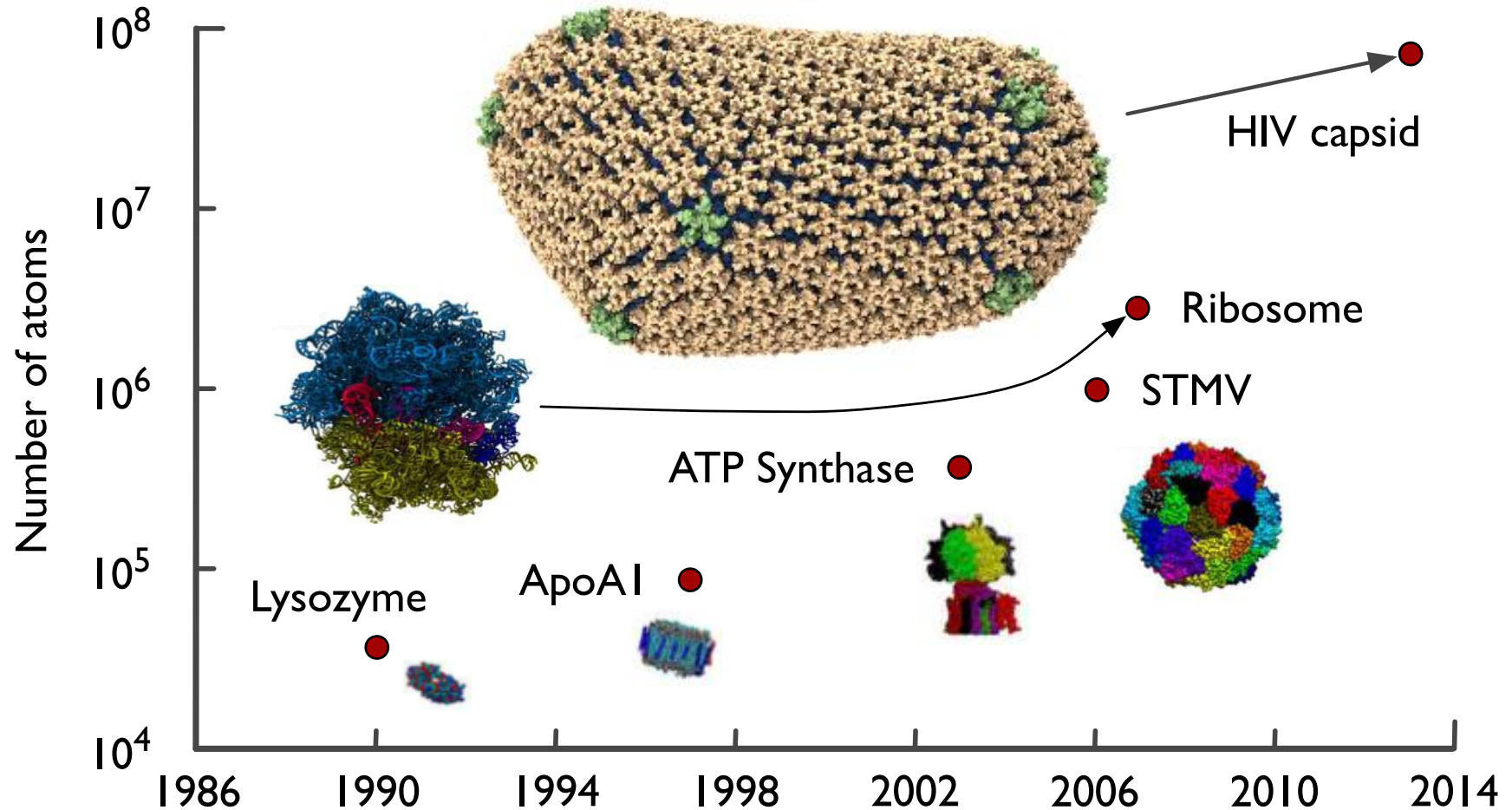


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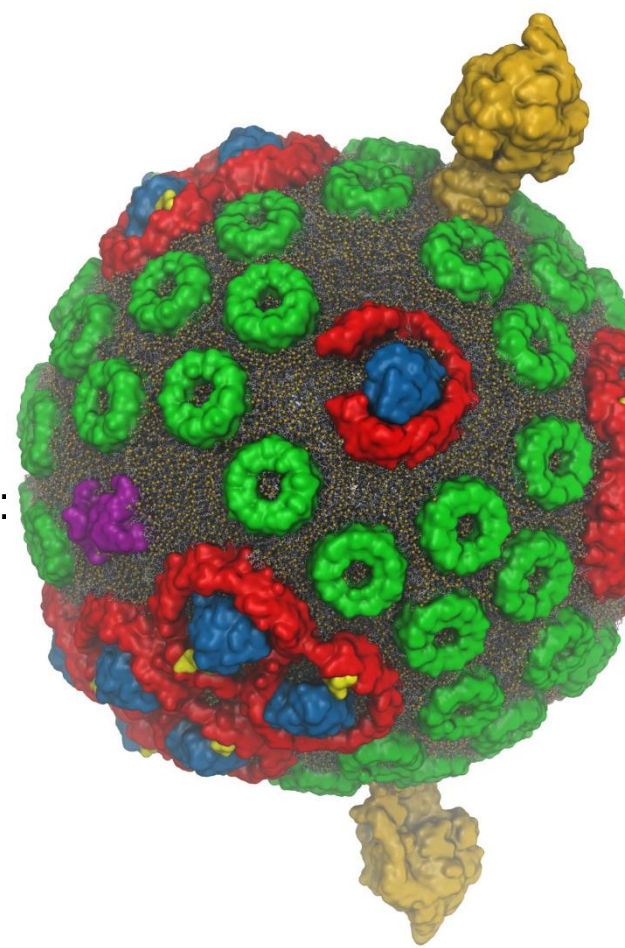
Computational Biology's Insatiable Demand for Processing Power



VMD 1.9.3 + OptiX 3.8 + CUDA 7.0

~1.5x Performance Increase

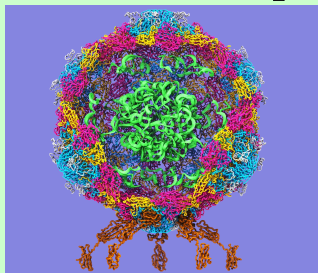
- OptiX GPU-native “**Trbv**h” **acceleration structure builder** yields substantial perf increase vs. CPU builders running on Opteron 6276 CPUs
- New optimizations in VMD TachyonL-OptiX RT engine:
 - **CUDA C++ Template specialization of RT kernels**
 - Combinatorial expansion of ray-gen and shading kernels at compile-time: stereo on/off, AO on/off, depth-of-field on/off, reflections on/off, etc...
 - Optimal kernels selected from expansions at runtime
 - **Streamlined OptiX context and state management**
 - **Optimization of GPU-specific RT intersection routines, memory layout**



VMD/OptiX GPU Ray Tracing of chromatophore w/ lipids.

VMD Molecular Structure Data and Global State

Scene Graph



Graphical Representations

DrawMolecule

Non-Molecular
Geometry

User Interface Subsystem

Tcl/Python Scripting

Mouse + Windows

VR Input "Tools"

Display Subsystem

VMDDisplayList

DisplayDevice

OpenGLDisplayDevice

FileRenderer

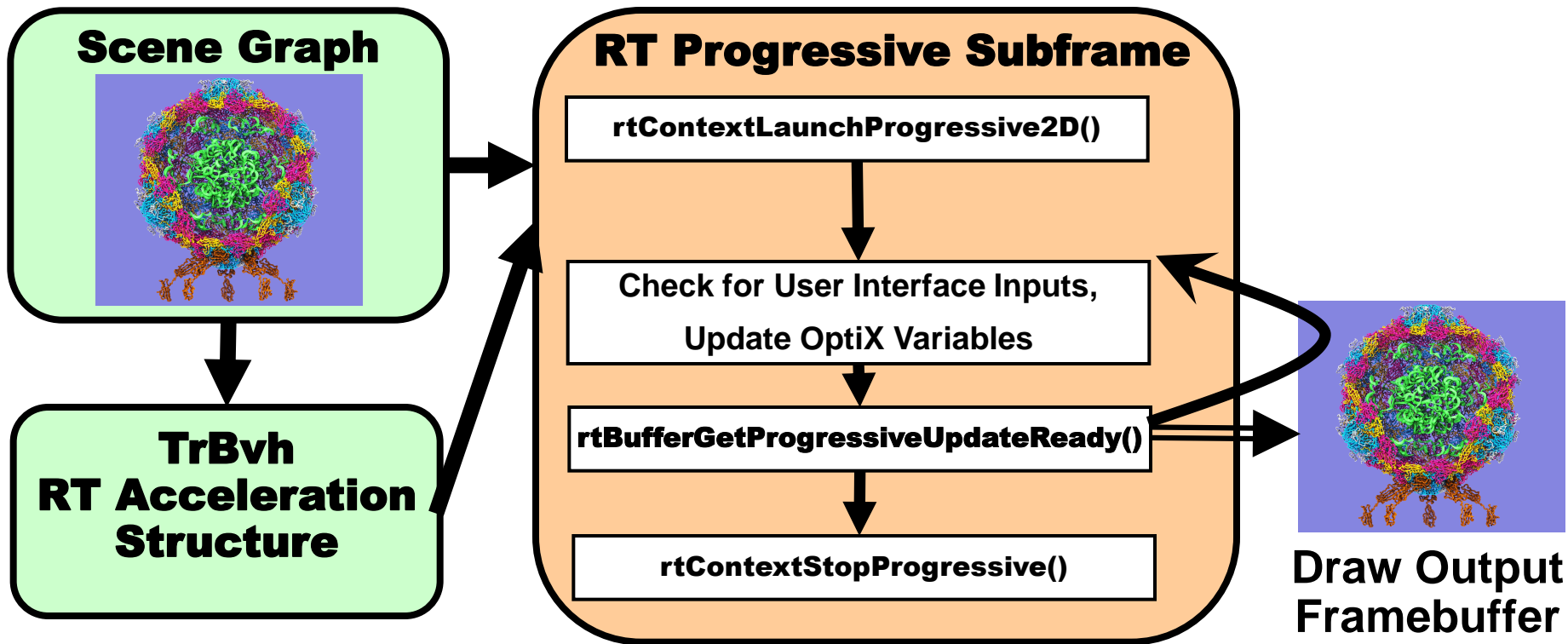
Windowed OpenGL GPU

OpenGL Pbuffer GPU

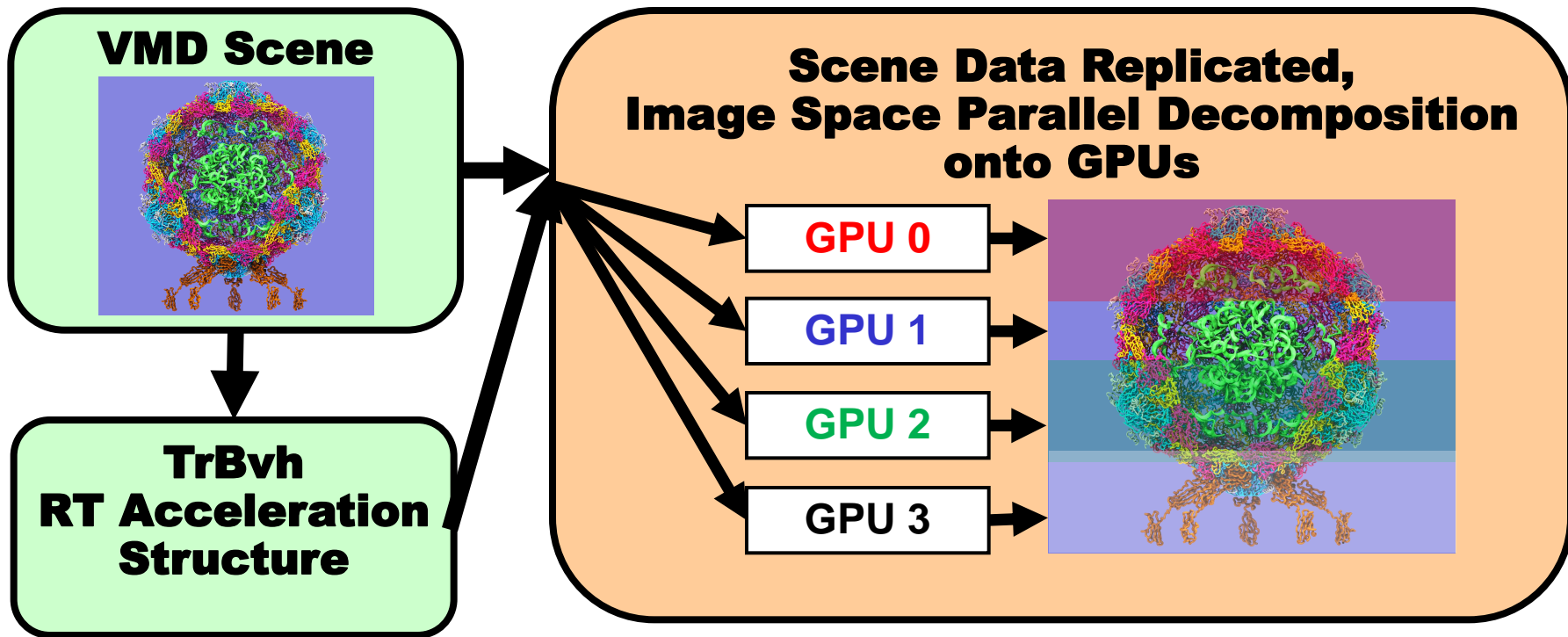
Tachyon CPU RT

TachyonL-OptiX GPU RT
Batch + Interactive

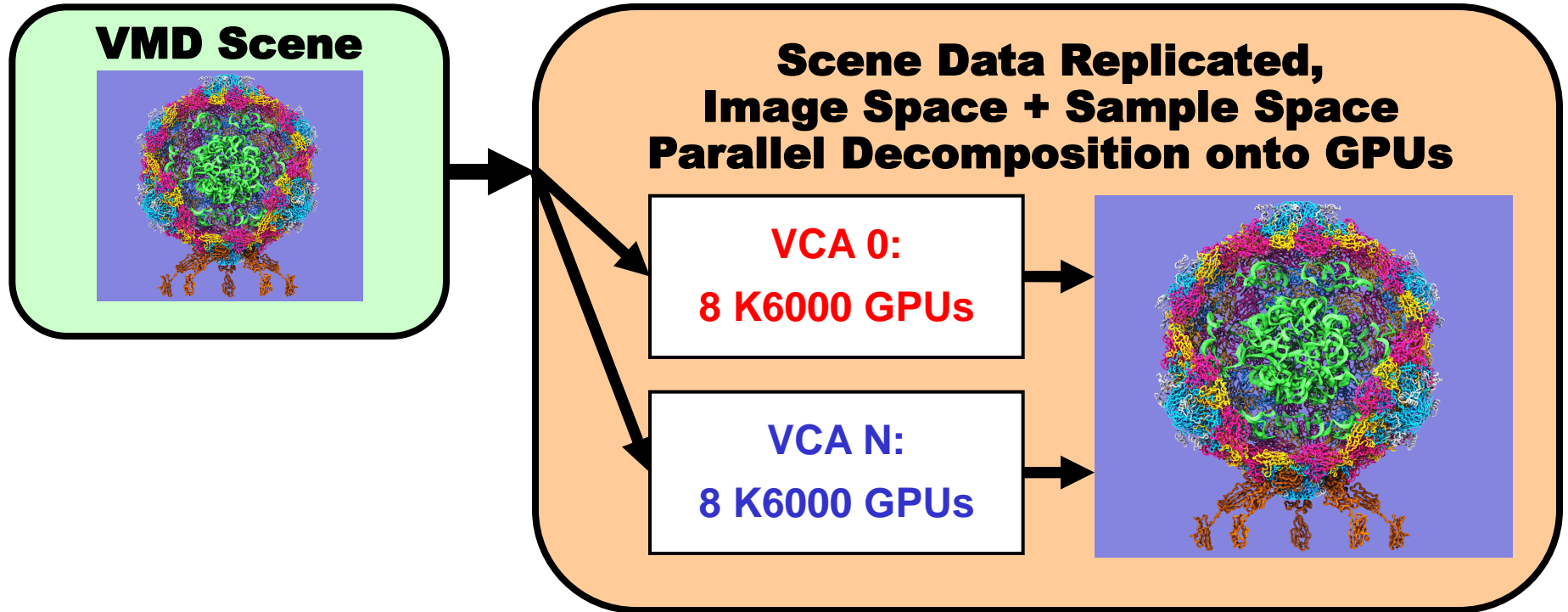
VMD TachyonL-OptiX Interactive RT w/ OptiX 3.8 Progressive API



VMD TachyonL-OptiX: Multi-GPU on a Desktop or Single Node

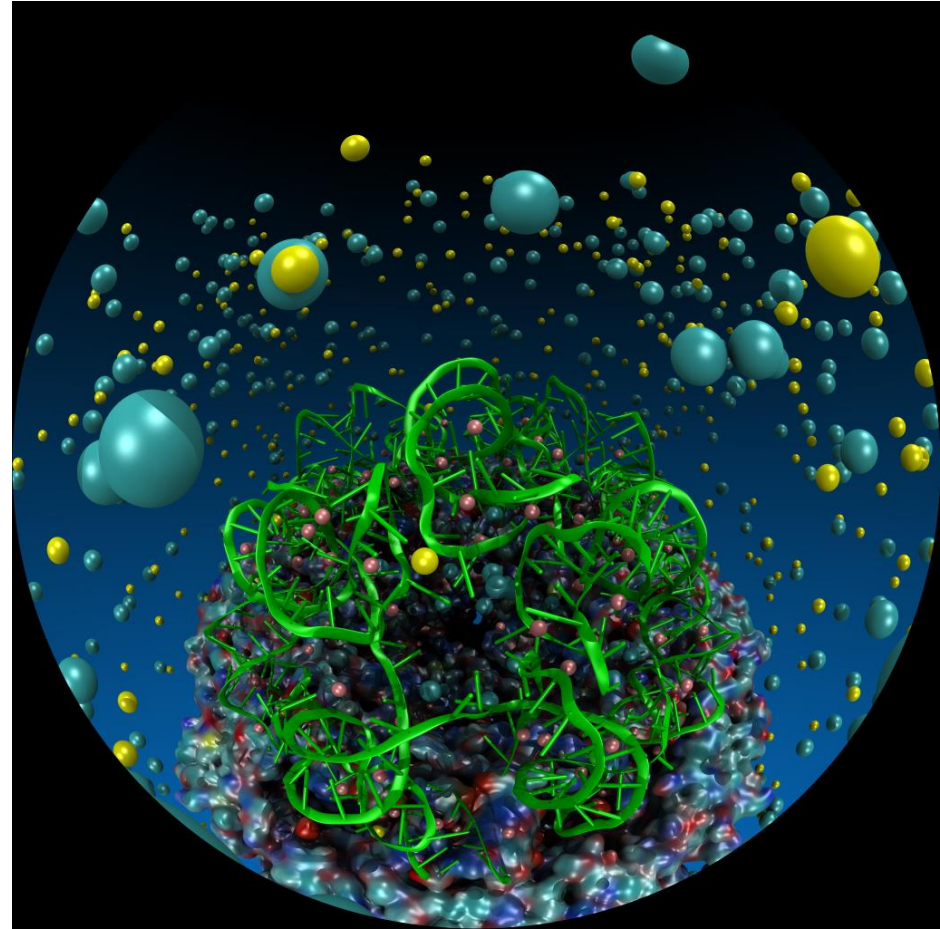


VMD TachyonL-OptiX: Multi-GPU on NVIDIA VCA Cluster



VMD Planetarium Dome Master Camera

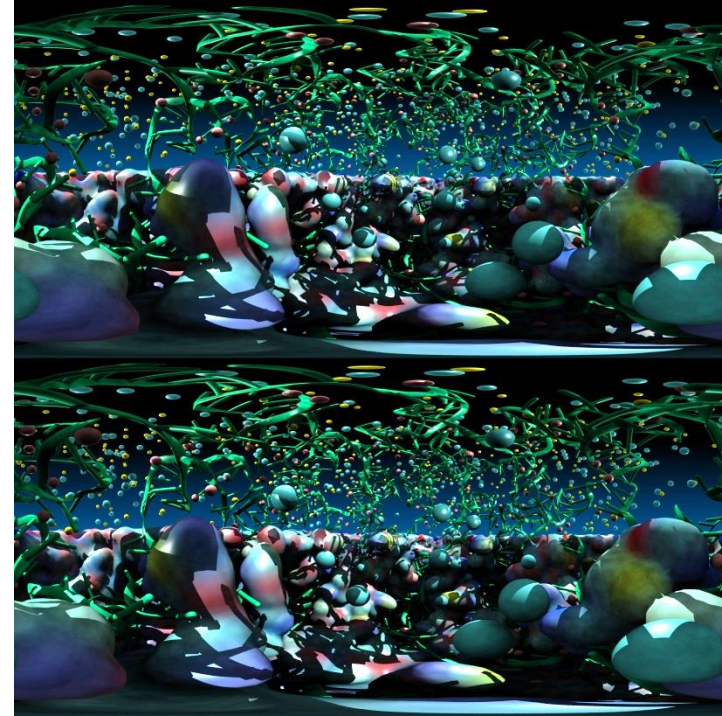
- RT-based dome projection -- rasterization poorly suited to non-planar projections
- Fully interactive RT with ambient occlusion, shadows, depth of field, reflections, and so on
- Both mono and stereoscopic
- No further post-processing required



Stereoscopic Panorama Ray Tracing w/ OptiX



- **Render 360° images and movies for VR headsets such as Oculus Rift, Google Cardboard**
- Ray trace panoramic stereo spheremaps or cubemaps for very high-frame-rate display via OpenGL texturing onto simple geometry
- Stereo requires spherical camera projections **poorly suited to rasterization**
- Benefits from OptiX multi-GPU rendering and load balancing, **remote visualization**



HMD Ray Tracing Challenges

- HMDs require high frame rates (**90Hz or more**) and minimum latency between IMU sensor reads and presentation on the display
- Multi-GPU workstations fast enough to direct-drive HMDs at required frame rates for simple scenes with direct lighting, hard shadows
- Advanced RT effects such as AO lighting, depth of field require much **larger sample counts**, impractical for direct-driving HMDs
- **Remote viz. required** for many HPC problems due to **large data**
- **Remote viz. latencies too high for direct-drive of HMD**
- **Our two-phase approach: moderate-FPS remote RT combined with local high-FPS view-dependent HMD reprojection w/ OpenGL**



**VMD+OptiX
Progressive RT
Engine**

**Omnistereo
Image Stream**

**View-dependent
OpenGL HMD
Reprojection
(up to 150 FPS)**

Scene

H.264 Video

**HMD Quaternion
+ Input Updates**

HDMI Video

HMD Pose

15Mbps Internet Link

**RT @ Remote VCA GPU Cluster
8 to 30 FPS @ 3072x1536**



HMD View-Dependent Reprojection with OpenGL

- Texture map panoramic image onto reprojection geometry that matches the original RT image formation surface
- HMD sees standard perspective frustum view of the textured surface
- Commodity HMD optics require **software lens distortion and chromatic aberration correction** prior to display, implemented with multi-pass FBO rendering
- Low-latency redraw as HMD head pose changes



VMD can support a variety of HMD lens designs, e.g.
<http://research.microsoft.com/en-us/um/redmond/projects/lensfactory/oculus/>



Come See A Live Demo!

- Demo shown by collaborators in Indiana U. booth on and off throughout exhibition
- RT @ NVIDIA VCA cluster in Santa Clara, thousands of miles away
- Work-in-progress:
 - 6DOF controller UI
 - Alternative HMD lens designs

Future Work

- Support for more commodity HMDs as they become generally available
- Support for OSes besides Linux
- Ray tracing engine and optimizations:
 - **Multi-node parallel RT and remote viz. on general clusters and supercomputers, e.g. NCSA Blue Waters, ORNL Titan**
 - Interactive RT stochastic sampling strategies to improve interactivity
 - Improved omnidirectional cubemap/spheremap sampling approaches
- Tons of work to do on VR user interfaces, multi-user collaborative visualization, ...



Acknowledgements

- Theoretical and Computational Biophysics Group, University of Illinois at Urbana-Champaign
- **Bill Sherman at Indiana University**
- NVIDIA OptiX and CUDA teams
- NCSA Blue Waters, ORNL OCLF Titan teams
- NVIDIA CUDA Center of Excellence, University of Illinois at Urbana-Champaign
- Funding:
 - NSF Blue Waters:
NSF OCI 07-25070, PRAC “The Computational Microscope”,
ACI-1238993, ACI-1440026
 - DOE INCITE, ORNL Titan: DE-AC05-00OR22725
 - NIH support: 9P41GM104601, 5R01GM098243-02



Visualization Publications

<http://www.ks.uiuc.edu/Research/vmd/>

- **Chemical Visualization of Human Pathogens: the Retroviral Capsids.** Juan R. Perilla, Boon Chong Goh, John E. Stone, and Klaus Schulten SC'15 Visualization and Data Analytics Showcase, 2015.
- **Atomic Detail Visualization of Photosynthetic Membranes with GPU-Accelerated Ray Tracing.** J. E. Stone, M. Sener, K. L. Vandivort, A. Barragan, A. Singharoy, I. Teo, J. V. Ribeiro, B. Isralewitz, B. Liu, B. Goh, J. C. Phillips, C. MacGregor-Chatwin, M. Johnson, L. F. Kourkoutis, C. N. Hunter, and K. Schulten. (submitted)
- **Visualization of Energy Conversion Processes in a Light Harvesting Organelle at Atomic Detail.** M. Sener, J. E. Stone, A. Barragan, A. Singharoy, I. Teo, K. L. Vandivort, B. Isralewitz, B. Liu, B. Goh, J. C. Phillips, L. F. Kourkoutis, C. N. Hunter, and K. Schulten. SC'14 Visualization and Data Analytics Showcase, 2014.
***Winner of the SC'14 Visualization and Data Analytics Showcase
- **Unlocking the Full Potential of the Cray XK7 Accelerator.** M. D. Klein and J. E. Stone. Cray Users Group, Lugano Switzerland, May 2014.
- **GPU-Accelerated Analysis and Visualization of Large Structures Solved by Molecular Dynamics Flexible Fitting.** J. E. Stone, R. McGreevy, B. Isralewitz, and K. Schulten. Faraday Discussions, 169:265-283, 2014.



Visualization Publications

<http://www.ks.uiuc.edu/Research/vmd/>

- **Stable Small Quantum Dots for Synaptic Receptor Tracking on Live Neurons.** E. Cai, P. Ge, S. Lee, O. Jeyifous, Y. Wang, Y. Liu, K. M. Wilson, S. Lim, M. A. Baird, J. E. Stone, K. Y. Lee, D. G. Fernig, M. W. Davidson, H. J. Chung, K. Schulten, A. M. Smith, W. N. Green, and P. R. Selvin. *Angewandte Chemie - International Edition in English*, 53(46):12484-12488, 2014.
- **Methodologies for the Analysis of Instantaneous Lipid Diffusion in MD Simulations of Large Membrane Systems.** Matthieu Chavent, Tyler Reddy, Joseph Goose, Anna Caroline E. Dahl, John E. Stone, Bruno Jobard, and Mark S.P. Sansom. *Faraday Discussions*, 169:455-475, 2014.
- **GPU-Accelerated Molecular Visualization on Petascale Supercomputing Platforms.** J. Stone, K. L. Vandivort, and K. Schulten. *UltraVis'13: Proceedings of the 8th International Workshop on Ultrascale Visualization*, pp. 6:1-6:8, 2013.
- **Early Experiences Scaling VMD Molecular Visualization and Analysis Jobs on Blue Waters.** J. Stone, B. Isralewitz, and K. Schulten. In *proceedings, Extreme Scaling Workshop*, 2013.



Visualization Publications

<http://www.ks.uiuc.edu/Research/vmd/>

- **Lattice Microbes: High-performance stochastic simulation method for the reaction-diffusion master equation.** E. Roberts, J. Stone, and Z. Luthey-Schulten. *J. Computational Chemistry* 34 (3), 245-255, 2013.
- **Fast Visualization of Gaussian Density Surfaces for Molecular Dynamics and Particle System Trajectories.** M. Krone, J. Stone, T. Ertl, and K. Schulten. *EuroVis Short Papers*, pp. 67-71, 2012.
- **Immersive Out-of-Core Visualization of Large-Size and Long-Timescale Molecular Dynamics Trajectories.** J. Stone, K. L. Vandivort, and K. Schulten. G. Bebis et al. (Eds.): *7th International Symposium on Visual Computing (ISVC 2011)*, LNCS 6939, pp. 1-12, 2011.
- **High Performance Computation and Interactive Display of Molecular Orbitals on GPUs and Multi-core CPUs.** J. Stone, J. Saam, D. Hardy, K. Vandivort, W. Hwu, K. Schulten, *2nd Workshop on General-Purpose Computation on Graphics Processing Units (GPGPU-2)*, *ACM International Conference Proceeding Series*, volume 383, pp. 9-18, 2009.
- **Visualization of Cyclic and Multi-branched Molecules with VMD.** Simon Cross, Michelle M. Kuttell, John E. Stone, and James E. Gain. *Journal of Molecular Graphics and Modelling*. 28:131-139, 2009.
- **A System for Interactive Molecular Dynamics Simulation.** John E. Stone, Justin Gullingsrud, Klaus Schulten, and Paul Grayson. In *2001 ACM Symposium on Interactive 3D Graphics*, John F. Hughes and Carlo H. Sequin, editors, pages 191-194, New York, 2001, ACM SIGGRAPH
- **An Efficient Library for Parallel Ray Tracing and Animation.** John E. Stone, Master's Thesis, University of Missouri-Rolla, Department of Computer Science, April 1998
- **Rendering of Numerical Flow Simulations Using MPI.** John Stone and Mark Underwood. Second MPI Developers Conference, pages 138-141, 1996.





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