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# A Three-Dimensional Visualization and Projection Workbench for the Pace HPC Initiative

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## **Progress Report: ThinkFinity Grant**

A Three-Dimensional Visualization and Projection Workbench for the Pace HPC Initiative

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## Purpose

The purpose of this project is to build a visualization workbench that supports the Pace High Performance Computing (HPC) Initiative in several ways:

#### A. Visualization

- 1. Allow design and building of molecular models and systems using force-feedback and 3D stereographics that will be submitted to the HPC cluster for further computation and refinement.
- 2. Allow real-time interactive force-feedback sensing and 3D stereographic visual analysis of molecular systems computed on the HPC cluster that includes the output of quantum mechanical, molecular mechanical and molecular dynamic simulations.
- 3. Provide an experimental test bed for investigating real-time molecular docking and interactive molecular dynamics of prototype systems.
- 4. Supports classroom demonstrations of above.

#### **B.** Projection

- 1. Develop a portable system for 3-D stereographic projection for seminar projection
- 2. Develop a local system for interactive 3-D stereographics in intimate setting.

## **Progress to Date**

Parts of the prototype system were ordered in January and February 2012. By mid-March all essential components for a basic test system had arrived and initial assembly began. The prototype system is comprised of a: INTEL XEON W3565 processor, with 8 GB RAM, 500 GB Hard Drive, 2GB Nvidia Quadro 4000 stereographic video card, Acer HN274H 27" Stereo 3D LCD Display, Windows 7 64 bit OS, and a Sensable Phantom Omni 6DOF haptic interface. System software was upgraded and drivers for all special hardware were installed.

Molecular visualization software was installed on the computer system, including the programs: VMD [1], PyMOL [2], Swiss PDB Viewer (DeepView) [3], UCSF Chimera [4], and Accelrys DS Visualizer [5]. All these programs were tested with the basic 3D stereographic display hardware. In addition, the HaptiMOL program by Stocks, Laycock, and Hayward was installed and tested [6]. HaptiMOL uses the Omni Phantom haptic device to allow a user to feel a molecule's solvent accessible surface, and apply forces to a molecular model and view its resulting deformation.

Work began on building software so the system could be used for interactive molecular dynamics (IMD) and steered molecular dynamics (SMD) [7]. This system requires three components: the NAMD molecular dynamics package [8], VMD, and VRPN, the Virtual Reality Peripheral Network [9], which provides a haptic interface to the Omni Phantom. Currently, the VRPN server is being built to interface with both the VMD molecular visualization software and the NAMD molecular dynamics package.

#### **Future Work**

Summer 2012 will be used to complete the construction of the force feedback system and test real-time docking. During Fall 2012 the system will be expanded to meet the requirements for 3D projection and demonstration.

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