Advances in Interactive Molecular Dynamics



A VR Framework for Interacting with Molecular Simulations

N. FEREY[#], O. DELALANDE[#], G. GRASSEAU⁺, and M. BAADEN[#]

*Laboratoire de Biochimie Théorique - UPR 9080, Institut de Biologie Physico-Chimique (IBPC), 13 rue Pierre et Marie Curie F-75005 Paris [†]Institut du Développement et des Ressources en Informatique Scientifique (IDRIS), Bâtiment 506, F-91403 Orsay cedex

This work is supported by two ANR grants for software development (Project ANR-07-CIS7-003) and biological applications (Project ANR-06-PCVI-0025).

Introduction

The NAMD-VMD [1-3] ensemble provides access to Virtual Reality (VR) devices for interacting with a Molecular Dynamics (MD) simulation in real time. In this interactive approach, named Interactive Molecular Dynamics (IMD), the use of VR peripherals yields good results for the analysis and study of complex biological systems. On the given simulation timescale, reversible and non-reversible events can be examined during the simulation. Our goal is to improve this approach by designing a VR framework, for developing tools dedicated to IMD in a VR context. This framework should allow developers and users to add **new** VR interaction paradigms and additional feedback modalities in order to provide new ways for steering a simulation and analysing its results.

Software Architecture for IMD in a VR context

We use the *MDDriver* library for interfacing a molecular User Interface (UI) with an MD simulation engine (GROMACS [4]) through a network connection. Thanks to this protocol, the new atom positions computed for each simulation timestep by the simulation, are sent to the molecular UI. The user can apply forces on atoms via this UI, which are then sent to the simulation and are taken into account in the next simulation step. As alternative molecular UI, we chose the visualisation toolkit VTK, allowing to quickly design visualisation and interaction paradigms dedicated to IMD. The VRPN library is used to manage VR devices, such as tracking devices for 3D interactions or haptic devices to provide force feedback to the user.

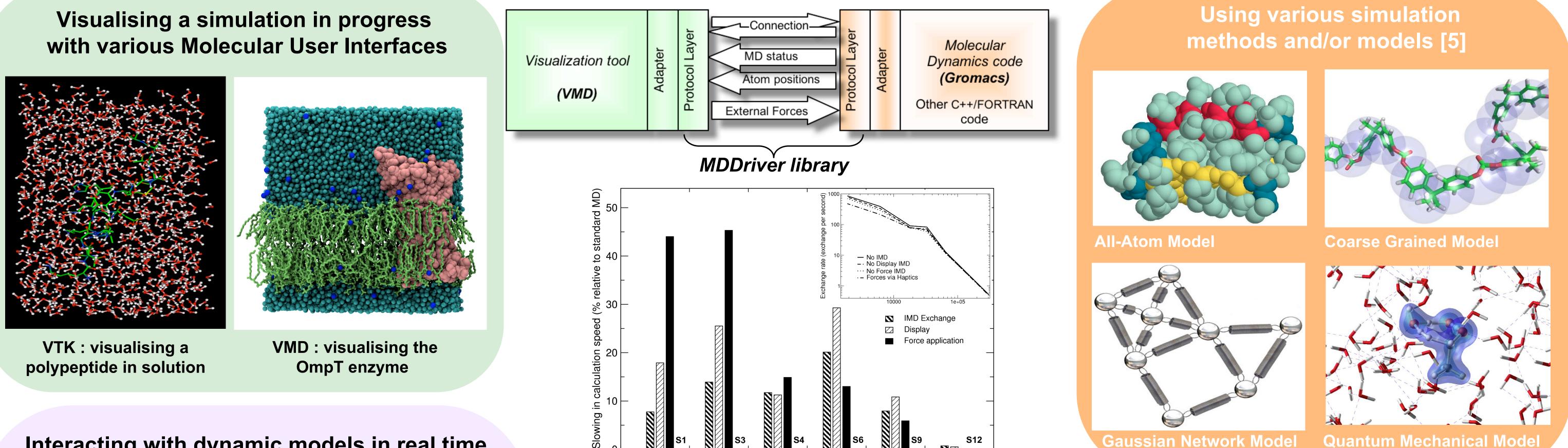
Why using VR techniques for Molecular Simulations

System exploration is easier using a 3D mouse or a haptic device to interact with molecular objects. Force feedback via haptics is especially well adapted to force-based methods such as molecular dynamics. It provides intuitive tactile feedback and enables the user to apply forces upon the virtual objects in order to guide or steer the simulation. A better perception of the three-dimensional organisation of large and complex biomolecular assemblies can be obtained by stereovision tools. The display surface can be extended on **multiple screens** in order to provide an immersive feeling to the user and create a collaborative work context.



European

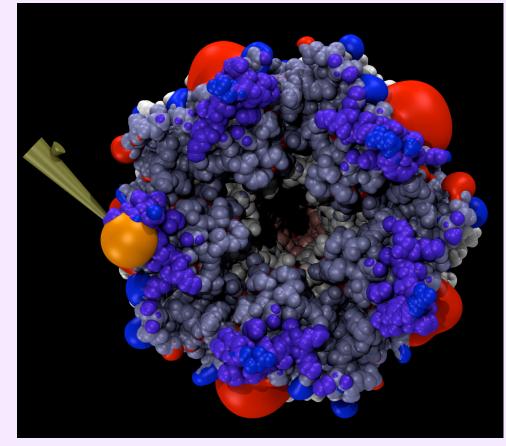
frastructure for



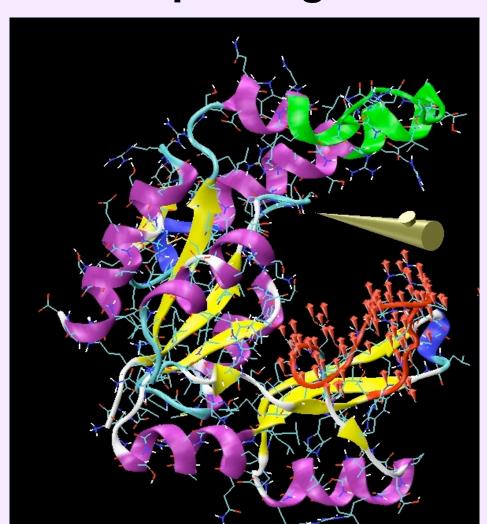
Interacting with dynamic models in real time using innovative devices and paradigms

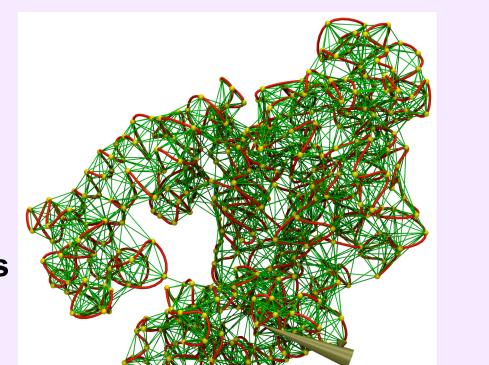
Interactive constraints (user applied forces) on a loop (red) of the GK enzyme

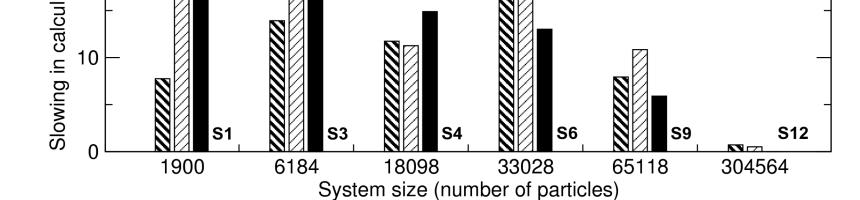
Exploring potential fields and isosurfaces of the **GLIC** ion channel



Improve mechanical properties of DNA transferase using elastic network (GNM)



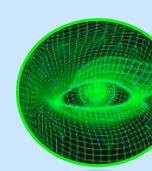




Quantum Mechanical Model

Relevant biological application : membrane fusion induced by the SNARE complex

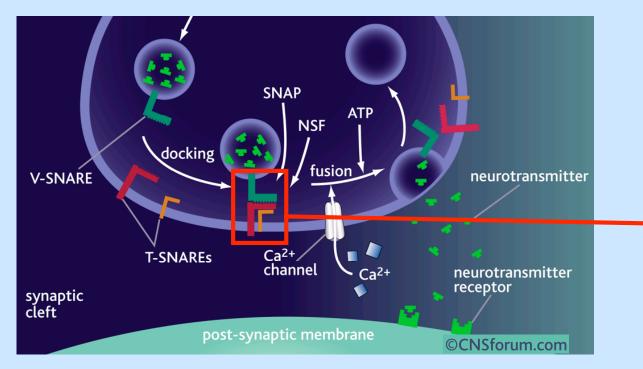
33028 particles



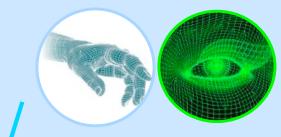
Instant visualization feedback (Low and High VR contexts)



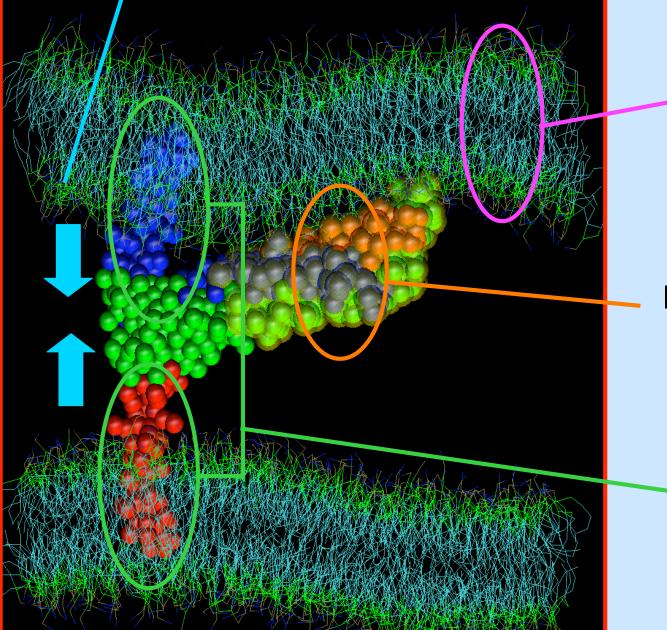
Haptic feedback (High VR context)



Synaptic membrane fusion



Mechanical process of SNARE action



Lipid bilayer behaviour

Inner structural coherence of the SNARE complex



Trans-membrane helices insertion (different in depth, orientation, strength)

Perspectives

Evolution of this VR framework is ongoing as a larger module, FVNano, to be encapsulated into the FlowVR middleware [6], a • FVNan library designed for exchanging and synchronizing VR events and data between computers.





[1] Humphrey, Dalke, and Schulten. VMD - Visual Molecular Dynamics. J. Mol. Graph. 14:33-38, 1996

[2] Phillips, Braun, Wang, Gumbart, Ta jkhorshid, Villa, Chipot, Skeel, Kale, and Schulten. Scalable molecular dynamics with NAMD. J. Comp. Chem. 26 :1781-1802, 2005 [3] Stone, Gullingsrud, and Schulten. A system for interactive molecular dynamics simulation. Proceedings of the 2001 symposium on Interactive 3D graphics, pages 191-194, 2001 [4] Hess, Kutzner, Vanderspoel, and Lindahl. GROMACS 4 : Algorithms for highly efficient, load-balanced, and scalable molecular simulation. J. of Chemical Theory and Comput., 2008 [5] Baaden and Lavery. There's plenty of room in the middle : Multiscale modelling of biological systems. Recent Advances in Structural Bioinformatics, 2007 [6] Allard, Gouranton, Lecointre, Limet, Melin, Raffin and Robert, FlowVR: a Middleware for Large Scale Virtual Reality Applications, Proceedings of Euro-par, 2004

Summary

We propose a virtual laboratory for IMD in a Virtual Reality context, composed of the following libraries and tools:

- **MDDriver** library provides new generic tools to connect calculation and visualization modules necessary for IMD simulations
- *vtk*-based interactive MUI allows rendering of molecular simulations
- vtkVRPN library manages Virtual Reality devices, such as trackers and haptic devices

See also http://mddriver.sourceforge.net and http://www.baaden.ibpc.fr/projects/fvnano/