

# HPC Molecular Dynamics Course

17-19 November 2014

## PRACE link

<http://events.prace-ri.eu/conferenceDisplay.py?confId=313>

<https://hpc->

[forge.cineca.it/files/CoursesDev/public/2014/Molecular\\_dynamics\\_in\\_an\\_HPC\\_environment/Bologna/](https://hpc-forge.cineca.it/files/CoursesDev/public/2014/Molecular_dynamics_in_an_HPC_environment/Bologna/)

## Programme

	Day-1	Day -2	Day -3
9.30 - 10.15	Registration. L1 HPC architectures I ( <b>A.Emerson</b> )	L4. Running MD on HPC architectures I. Clusters and Hybrid clusters. ( <b>A. Grottesi</b> )	L5 Running MD on HPC architectures II. Bluegene/Q. ( <b>G. Chillemi</b> )
10.15 - 11.00	L1. Introduction to HPC architectures II ( <b>A. Emerson</b> )	Invited lesson 1. ( <b>R. Gnudi, Uni. Modena</b> ) Porting of an MD code on Xeon PHI.	Tutorial T3. Analysis of MD trajectories: Essential Dynamics ( <b>A. Grottesi</b> )
11.00 - 11.30	Break	Break	Break
11.30- 12.30	L2. Introduction to Classical Molecular Dynamics ( <b>G. Chillemi</b> )	Invited Lesson 2. ( <b>I. Eberini, Uni. Milano</b> ) Simulation of urea- induced protein unfolding: a lesson from bovine $\beta$ - lactoglobulin	Invited Lesson 3. PLUMED ( <b>G. Bussi, SISSA</b> )
12.30 - 14.00	LUNCH	LUNCH	LUNCH
14.00 - 15.00	L3. Parallel Molecular Dynamics: Atom, Force and Domain decomposition ( <b>A. Emerson</b> )	Tutorial T2. Running and analysing Gromacs on a GPU cluster. ( <b>A. Grottesi</b> )	L6. A guide for accessing computer resources from funding bodies such as PRACE. ( <b>A. Emerson</b> )
15.00 - 17.30	Tutorial T1. Analysis of a parallel program.	T2. continued ( <b>A. Grottesi</b> )	Practical session or FREE