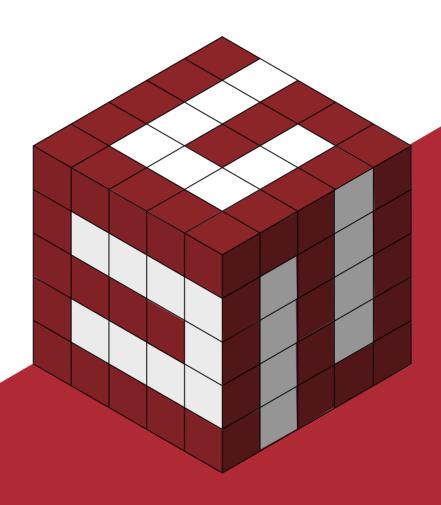




GENESIS

Overview and Demo at SC17



01

14th Nov. AM11:00-11:10

General introduction of GENESIS

Chigusa Kobayashi (RIKEN, AICS)

02

14th Nov. AM11:10-11:20

High performance scheme in GENESIS

Jaewoon Jung (RIKEN, AICS)





Molecular dynamics (MD) software for bimolecular systems.

Open source software distributed under the GPLv2 license.

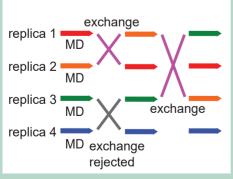


Please check 'GENESIS' website URL: http://www.aics.riken.jp/labs/cbrt/

Feature

- High performance schemes using hybrid parallelization (MPI+OpenMP) and/or GPGPU
- 2. Enhanced sampling algorithm for free-energy calculation
- 3. Multi-scale MD (All atom, Coarse-Grained, QM/MM)
- 4. Data assimilation

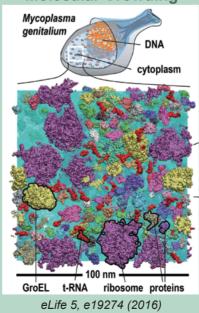
Enhanced sampling



Target Application

- 1. Molecular crowding
- 2. Structure/Dynamics of biomolecules
- 3. Data driven simulations

Molecular Crowding



Performance of GENESIS on K computer

