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IMPLEMENTATION OF THE « BLUE-MOON » ENSEMBLE METHOD IN ABINIT

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- Many problems in physics involve RARE events, i.e
 <u>events occuring rarely w/r typical atomic vibrations</u>
- Yet, these events may control some **MACROSCOPIC** properties
- <u>Ex 1 :</u> diffusion of an ionic impurity/defect in a solid
 => diffusion coefficient / ionic conductivityin an <u>IONIC CONDUCTOR</u>
- Ex 2 : reorientation of a dipolar impurity / cluster => dielectric relaxation phenomena in the RF in a <u>RELAXOR</u>
- Events thermally activated => controlled by an activation energy E_a Occurrence rate :

$$r = r_0 e^{-E_a/k_B T}$$



1- CONTEXT : SIMULATION OF « RARE » EVENTS

$$r = r_0 e^{-E_a/k_B T}$$

How to obtain
$$E_a$$
 (at finite T) ?

1st possibility : simulating the dynamics of the system (MD), ... and counting the occurrences of the events



1- CONTEXT : SIMULATION OF « RARE » EVENTS

- 2nd possibility : « Blue-Moon » ensemble : Method to simulate the occurrence of RARE events within Molecular Dynamics (MD)



Blue-Moon Ensemble method in practise :

simulate each step of the event by « forcing » ξ to take
 a chosen value :

=> MD under (<u>holonomic</u>) constraint of FIXED ξ

Method of Lagrange multipliers

- Obtain for each value of ξ the derivative of the free energy « mean force » on the constraint : $-\frac{dF}{d\xi}$

- recover the (free) energy profile by (thermodynamic) integration :

$$\Delta F = F(\xi) - F(\xi_0) = \int_{\xi_0}^{\xi} \frac{dF}{d\xi}(\xi') d\xi'$$

Sprik, Cicotti, J. Chem. Phys. **109**, 7737, (1998)

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2 – THE « BLUE-MOON » ENSEMBLE METHOD

 $\frac{dF}{d\xi} = \frac{\langle Z^{-1/2}(-\lambda + k_B T G) \rangle_{\xi}}{\langle Z^{-1/2} \rangle_{\xi}}$

General case : $\xi({x_{i\alpha}})$

with :



Time average under fixed ξ (constrained MD) = « blue-moon » ensemble average

$$F(\xi_2) - F(\xi_1) = \int_{\xi_1}^{\xi_2} \frac{dF}{d\xi}(\xi') d\xi'$$

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CEA | MAY 2017 | PAGE 7

Implementation in ABINIT of the LINEAR constraint i.e. linear combination between atomic positions

$$\xi(\{x_{i\alpha}\}) = \sum_{i\alpha} a_{i\alpha} x_{i\alpha}$$

1) Lagrangian with constraints

Real coefficients

$$L = [T(\{\dot{x}_{i\alpha}\}) - V(\{x_{i\alpha}\})] - \lambda [\sum_{i\alpha} a_{i\alpha} x_{i\alpha} - \xi]$$

2) Apply *Euler-Lagrange equations* :

$$m_i \frac{d^2 x_{i\alpha}}{dt^2} = f_{i\alpha} - \lambda a_{i\alpha}$$

Lagrange multiplier computed at each step as

3) Derivative of the free energy : $\frac{dF}{d\xi} = -\langle \lambda \rangle_{\xi} = -\frac{\sum_{i\alpha} \frac{a_{i\alpha}}{m_i} \langle f_{i\alpha} \rangle_{\xi}}{\sum \frac{a_{i\alpha}^2}{\lambda_{i\alpha}}}$

$$\lambda(t) = \frac{\sum_{i\alpha} m_{i\alpha}}{\sum_{i\alpha} \frac{a_{i\alpha}^{2}}{m_{i}}}$$

 $\sum a \frac{f_{i\alpha}(t)}{dt}$

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- **Physically acceptable** linear constraints satisfy : $\sum_{i\alpha} a_{i\alpha} = 0$
- However, even if $\sum_{i\alpha} a_{i\alpha} \neq 0$, a supplemental constraint

has been added to ensure that center of mass does not move.

- Value of the constraint ξ : **FIXED** by **INITIAL** set of positions.

- Initialization of the velocities :

The initial velocities must be modified to ensure that the constraint is obeyed, i.e.

$$\sum_{i\alpha} a_{i\alpha} \frac{dx_{i\alpha}}{dt} = 0$$

- Implemented in the routine pimd_apply_constraint (m_pimd.F90).
- PIMD : the constraint is applied on the centroid.



The constraint is STRICTLY obeyed all along the MD trajectory (at EACH step, NOT on average)

Ex:

- Reaction coordinate = -36.5852098477
- the Center of mass is also strictly FIXED

Center of mass:

4.6876358799	6.5991695708	4.6808066009
Center of mass:		
4.6876358799	6.5991695708	4.6808066009
Center of mass:		
4.6876358799	6.5991695708	4.6808066009

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4 - EXAMPLE



In progress :

BaCeO₃ : orthorhombic perovskite *Pnma* space group

Excellent protonic conductor

Inter-octahedral transfer of H⁺

Cell = 20 atoms Reaction coordinate : $(O_1H)_y - (O_2H)_y$ = $y(H)-y(O_1) - [y(O_2) - y(H)]$

PIMD, T=200 K P=32 beads

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Input file for BaCeO3:

nline 6 npulayit 15

PARALLELIZATION # => 864 procs

paral_kgb 1 npkpt 9 npband 3 npfft 1 npimage 32

PIMD/MD

optcell 0 irandom 3 restartxf -1 **imgmov 9 # langevin PIMD** ntimimage 10000 **nimage 32 mdtemp 400 200 vis 5d-05** dynimage 32*1 nsym 1 pitransform 0 amu 140 16 138 1 # ce o ba h pimass 140 16 138 1

CONSTRAINTS

pimd_constraint 1 nconeq 1 natcon 3 iatcon 5 20 1 #ox1 ox2 hydrogen wtatcon 0.0 -1.0 0.0 0.0 -1.0 0.0 0.0 2.0 0.0

DFT

nband 111 occopt 3 tsmear 0.001 pawovlp -1 ixc 11 ecut 18.0 pawecutdg 25.0 kptopt 1 ngkpt 3 2 3 nstep 100 toldff 1.0d-05 prtden 0 prtwf 0 charge 1.0

ATOMS

GEOMETRY

acell 11.941471 16.813056 11.913748

xred 0.279 0.216 0.723 #yH to be varied 2.5108646807E-02 2.500000000E-01 -8.0469316958E-03 -1.3877787808E-17 -9.7144514655E-17 5.000000000E-01 4.8011508619E-01 2.500000000E-01 7.9658009587E-02 2.7974388262E-01 4.3160947462E-02 7.2102708776E-01 #ox1 9.7489135319E-01 7.500000000E-01 1.0080469317E+00 5.1988491381E-01 7.500000000E-01 9.2034199041E-01 7.2025611738E-01 9.5683905254E-01 2.7897291224E-01 5.2510864681E-01 2.500000000E-01 5.0804693170E-01 4.7489135319E-01 7.500000000E-01 4.9195306830E-01 5.00000000E-01 5.00000000E-01 -6.9388939039E-17 -1.3877787808E-17 5.000000000E-01 5.00000000E-01 5.000000000E-01 -9.7144514655E-17 -6.9388939039E-17 9.8011508619E-01 2.500000000E-01 4.2034199041E-01 1.9884913809E-02 7.500000000E-01 5.7965800959E-01 7.7974388262E-01 4.5683905254E-01 7.7897291224E-01 7.2025611738E-01 5.4316094746E-01 2.7897291224E-01 2.2025611738E-01 9.5683905254E-01 2.2102708776E-01 2.2025611738E-01 5.4316094746E-01 2.2102708776E-01 2.7974388262E-01 4.5683905254E-01 7.2102708776E-01 #ox2

7.7974388262E-01 4.3160947462E-02 7.7897291224E-01



5 – CONCLUSION : FUTURE IMPLEMENTATIONS

- 1) <u>Future developments</u> : other constraints, i.e.
- distance between 2 atoms
- difference of distance
- angle between directions defined by atoms
- 2) Extension to the NPT ensemble :
- => study of *Structural Phase Transitions*
- 3) Multiple constraints