

GRASSMANN EXTRAPOLATION AS A TOOL TO ACCELERATE BORN-OPPENHEIMER MOLECULAR DYNAMICS

F. Lipparini, F. Pes, P. Mazzeo, É. Polack, G. Dusson, and B. Stamm
Dipartimento di Chimica e Chimica Industriale, Università di Pisa
Via G. Moruzzi 13, 56124 Pisa, Italy
`filippo.lipparini@unipi.it`

Born-Oppenheimer molecular dynamics (BOMD), is a powerful, yet expensive computational tool, where the system's nuclei are propagated classically on a potential energy surface generated on the fly with Density Functional Theory (DFT). The iterative solution to the DFT non-linear equations is the most computationally demanding step in BOMD: to speed up such calculations, various extrapolation strategies have been developed to use information available at previous simulation steps as a guess for the iterative procedure.

In this contribution, we present an approach to extrapolate density matrices, rank n projectors that codify the quantum mechanical information, by using computable maps between the Grassmann manifold where they are defined to its tangent space and back [1]. After comparing the new approach to the state of the art, we discuss its current limitations, with particular focus on energy conservation problems. We introduce a time-reversible extrapolation and an approximately reversible one, that exhibits optimal performance and good numerical stability.

References

- [1] É Polack et al., *Grassmann Extrapolation of Density Matrices for Born–Oppenheimer Molecular Dynamics*, *J. Chem. Theory Comput.* 17 (2021), pp. 6965–6973.