Tinker-HP 1.2 : Release Notes

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1 Citing Tinker-HP

If you use Tinker-HP 1.2 please cite the following reference:

Tinker-HP: a Massively Parallel Molecular Dynamics Package for Multiscale Simulations of Large Complex Systems with Advanced Polarizable Force Fields. L. Lagardère, L.-H. Jolly, F. Lipparini, F. Aviat, B. Stamm, Z. F. Jing, M. Harger, H. Torabifard, G. A. Cisneros, M. J. Schnieders, N. Gresh, Y. Maday, P. Ren, J. W. Ponder, J.-P. Piquemal, Chem. Sci., 2018, 9, 956-972 (Open Access) DOI: 10.1039/C7SC04531J

If you use the AVX512 vectorized version of Tinker-HP 1.2, please also cite:

Raising the Performance of the Tinker-HP Molecular Modeling Package [Article v1.0]. L. H. Jolly, A. Duran, L. Lagardère, J. W. Ponder, P. Y. Ren, J.-P. Piquemal, LiveCoMS, 2019, 1 (2), 10409 (Open Access) DOI: 10.33011/livecoms.1.2.10409

2 New methods and improvement

Several new methods and improvements have been included in Tinker-HP version 1.2, they can be listed as follows:

• Truncated Conjugate Gradient (TCG) non iterative solver(s) for the polarization equations:

This method approximates the fully converged dipoles by a (systematically improvable) analytic expression which thus avoids the error in the computation of the related forces due to the use of a (never fully converged) iterative method. Furthermore, depending on its level of accuracy, TCG can be faster than the standard solution of the dipoles via an iterative method.

KEYWORDS: The use of TCG is set by using the line: **polar-alg 3** in the *.key file, additional keywords to control the parameters of TCG (preconditioner, peek step and guess) can be found in the **Readme** of Tinker-HP.

Because no analytical virial tensor is computed for TCG yet, NPT simulations with TCG are limited to the use of the Monte-Carlo barostat.

References

- Truncated Conjugate Gradient (TCG): an optimal strategy for the analytical evaluation of the many-body polarization energy and forces in molecular simulations. F. Aviat, A. Levitt, Y. Maday, B. Stamm, P. Y. Ren, J. W. Ponder, L. Lagardère, J.-P.Piquemal, J. Chem. Theory. Comput., 2017, 13, 180-190 (Open Access) DOI: 10.1021/acs.jctc.6b00981
- The Truncated Conjugate Gradient (TCG), a Non-iterative/Fixed-cost Strategy for Computing Polarization in Molecular Dynamics: Fast Evaluation of Analytical Forces. F. Aviat, L. Lagardère, J.P. Piquemal, J. Chem. Phys., 2017, 147, 161724 DOI: 10.1063/1.4985911
- New multi-timesteps integrators: Three-levels Respa1-like integrators have been introduced where the potential is evaluated at three different levels (and not just two as for the standard Respa integrator):

- the fast bonded terms
- the intermediate short-range non-bonded terms: short range van der Waals, short range (real space) electrostatics and short range (real space) polarization for polarizable force fields
- the long range non-bonded terms: long range van der Waals, long range + reciprocal space electrostatics, total polarization-short range (real space) polarization for polarizable force fields

KEYWORDS: These splittings can be used with a **BAOAB** inner loop for NVT simulations (keyword **baoabrespa1**) or with a velocity-verlet inner loop (keyword **respa1**), it has been showed that the BAOAB based respa1 integrators are always more stable than the Velocity-verlet based respa1 integrators.

The timesteps used for the three levels can be controlled in the key-file as well as the solver used for short-range polarization for polarizable force fields as described in the README of Tinker-HP. When used in conjunction with Hydrogen-Mass-Repartitioning (keyword **heavy-hydrogen**) and the use of a simple TCG1 short range polarization solver, AMOEBA computation can be made up to 7 times faster than with a standard 1fs Velocity Verlet integrator as described in the reference above.

Reference

Pushing the limits of Multiple-Timestep Strategies for Polarizable Point Dipole Molecular Dynamics.
 L. Lagardère, F. Aviat, J.-P. Piquemal, J. Phys. Chem. Lett., 2019, 10, 2593-2599
 DOI: 10.1021/acs.jpclett.9b00901

• Langevin Piston for NPT simulations:

The Langevin Piston extended Lagrangian method has been implemented with a baoab integration of the volume extended variable, with the keyword **barostat langevin** or **integrator baoabpiston**. In both cases, the only compatible integrator is a standard **BAOAB** limiting the usable timestep to around 1fs. The mass of the piston as well as the associated friction can be controlled by keyword reviewed in the Readme of Tinker-HP.

Reference

- Constant pressure molecular dynamics simulation: The Langevin piston method. Scott E. Feller, Yuhong Zhang, and Richard W. Pastor J. Chem. Phys., 1995, 103, 4613 DOI: 10.1063/1.470648

• Addition of Steered Molecular Dynamics:

Steered Molecular Dynamics (SMD) was added to Tinker-HP 1.2 version.

KEYWORDS: SMD can be set with the keywords **CVSMD** for constant velocity SMD and **CF-SMD** for constant force SMD. Details of how to use SMD within Tinker-HP can be found in the **SMD Tutorial** made by Frederic Célerse which can be found within in the tutorials/SMD/ directory of the release.

Reference

 Massively parallel implementation of Steered Molecular Dynamics in Tinker-HP: polarizable versus non-polarizable simulations. F. Célerse, L. Lagardère, E. Derat, J.-P.Piquemal, J. Chem. Theory. Comput. 2019, 15, 3694-3709 DOI: 10.1021/acs.jctc.9b00199

• Parallel version of the regular Tinker BAR program for Free Energy differences

The exact equivalent of the regular Tinker "BAR" program has been implemented within the massively parallel Tinker-HP framework. Given two trajectories stored in *.arc files and the corresponding two Hamiltonians characterized by to two different *key files, it allows to compute the free energy difference between the two states with the Bennett Acceptance Ratio method.

- Faster linked-cell method to compute neighbor lists
- Faster rattle algorithm in parallel

3 New compilation and installation method

Tinker-HP now uses a configure script built with autotools packages from GNU to ease the compilation and installation process. Apart from the usual options available with all configure scripts, there are specific options for Tinker-HP.

```
Usage: ./configure [OPTION]... [VAR=VALUE]...
```

Optional Features:

--enable-debug Enable debug mode (check array bounds, implicit

none, etc...). Should not be active in normal

operations [default is no]

--enable-skylake Enable AVX512 Optimization for Skylake Processors

[default is no]

--enable-knl Enable AVX512 Optimization for KNL (Xeon Phi)

Processors [default is no]

--enable-fft-generic Enable generic FFT mode [default is yes]
--enable-fft-mkl Enable MKL FFT mode [default is no]
--enable-fft-fftw3 Enable fftw3 FFT mode [default is no]
--enable-fft-fftw3_f03 Enable fftw3_f03 FFT mode [default is no]

Optional Packages:

--with-blaslib=<BLAS LIB> Specify BLAS library [mkl, lapack or

/absolute/path/to/BLAS_library]

--with-fftlib=<FFT LIB> Specify a library for FFT called by 2decomp [mkl or

fftw3 or /absolute/path/to/FFTW_library]

Users should now be able to have Tinker-HP running by doing:

```
./configure ; make ; make install
```

Please read the instructions on how to use the configure script in the Readme_v1.2.pdf.