Molecular Dynamics (MD) computer simulations have become a standard tool for investigating a variety of atomic and molecular systems ranging from solids to simple fluids to complex biomolecular assemblies. They are particularly attractive for dynamics and for equilibrium sampling in high density condensed matter systems where large scale collective modes may be significant. These modes may not be easily excited (and relaxed) by the alternative approach to phase space sampling, namely Monte Carlo (MC) simulations, because: (i) MC evolution is diffusive in nature, and (ii) MC tends to have low acceptance rates in high density regions. The MC method, however, possesses one significant advantage over MD - the ability to sample, at least in principle (i.e., for sufficiently long runs), almost any statistical ensemble in a fairly straightforward manner. This task is accomplished by performing MC moves that are ergodic and satisfy the detailed balance condition. In the canonical (N, V, T) ensemble (where N, V, and T denote the number of particles, volume, and temperature, respectively), the latter requirement is usually fulfilled by using the Metropolis criterion for the acceptance probability: $p_{\text{acc}} = \min[1, \exp(-\Delta U/k_B T)]$, where $k_B$ is the Boltzmann constant and $\Delta U$ is the change in the potential energy between the two states which are approached via opposite moves. Similarly, the isothermal-isobaric (N, P, T) (where P is the pressure) ensemble can be simulated by including coordinate displacements that change the volume of the system and scale the coordinates of the particles accordingly, and by redefining the potential energy to $U_{\text{eff}} = U + PV - Nk_B T \ln V$ [see, later, Eq. (13)].

Things become more complicated when it comes to MD simulations, which attempt to follow the dynamics of a molecular system by numerically solving Newton’s classical equations of motion for the constituent particles. This, supposedly, generates trajectories within the microcanonical ensemble (N, V, E) (where E is the internal energy of the system, which is the sum of potential and kinetic energies) although, due to round-off errors, one should not expect the computed trajectories to actually follow the real-time dynamics in many-particle systems. The most commonly used discrete-time integrator for MD simulations is the St"ormer-Verlet algorithm which (in its so-called “velocity-Verlet” form) reads:

$$r^{n+1} = r^n + v^n dt + \frac{dt^2}{2m} f^n$$
$$v^{n+1} = v^n + \frac{dt}{2m} (f^n + f^{n+1}),$$

where $r^n$, $v^n$, and $f^n$ denote, respectively, the coordinate, velocity, and the force acting on the particle with mass $m$ at time $t_n$, and $t_{n+1} = t_n + dt$. Notice that $r^n$, $v^n$, and $f^n$ represent Cartesian components, which means that for a system of N particles in a space with dimensionality d, the number of equations one needs to compute per time step $dt$ is $2dN$. The Verlet algorithm results in a trajectory which is accurate to second order in the time step $dt$. This deviation between the computed and the “correct” trajectories should not be a matter of concern if the simulations properly sample the correct statistical ensemble, or otherwise retain the measures of interest. Thus, the critical test for the performance of any numerical integrator must be its accuracy in measuring important thermodynamic quantities and the variations of the results with $dt$. Remarkably, the Verlet algorithm suffers from the problem that the total kinetic energy of a simulated system (which is supposed to be proportional to the temperature) becomes progressively depressed for...
increasing time step $dt$ compared to the potential energy (see also Appendix A for a harmonic oscillator demonstration of how the velocity in discrete time is not precisely the velocity of the corresponding spatial coordinate). Other thermodynamic observables also exhibit variations with $dt$, which makes a striking contrast with MC simulations in which the thermodynamic configurational averages are insensitive to the discrete steps.

The microcanonical ensemble sampled in MD simulations does not provide the best representation of experimental conditions, where the most common condition is that of constant temperature and pressure. Therefore, considerable effort has been devoted to the development of MD algorithms for simulations of the isothermal-isobaric ensemble. In the simplest method, proposed by Berendsen, the system is weakly coupled to external heat (“thermostat”) and pressure (“barostat”) baths, using the principle of least local perturbation \[7\]. This method has been criticized for failing to correctly sample the statistical ensemble, due to its tendency to suppress fluctuations in kinetic energy and volume. A second, more reliable method, pioneered by Andersen for fixed pressure \[8\], extended by Parrinello and Rahman \[9\] and by Nosé \[10\], and revised by Hoover \[11\] to fixed temperature MD simulations, is the extended Lagrangian formalism. The method is based on the idea of including additional degrees of freedom, corresponding to the volume and/or the kinetic energy of the system, together with their conjugate momenta variables. The new variables are coupled to the system in a manner which guarantees that the trajectory correctly samples the isothermal-isobaric ensemble. The latter constitutes a sub-space of the configuration space of the extended system. Within the extended phase space, the statistics is microcanonical and the equations of motion can be derived from the extended Hamiltonian, which is conserved in time. In principle, these Hamiltonian equations of motion can be integrated numerically using the Verlet algorithm. In practice, the implementation of the discrete-time Verlet algorithm raises several significant challenges and difficulties. Specifically, for the barostat part, the coupling between the particles’ degrees of freedom and the “piston” (introduced to control the volume fluctuations) is the source of the following problems:

1. When the piston moves, the particle coordinates must be rescaled, which leads to a metric problem with the algorithm. This problem has been addressed in Refs. 12-14.

2. The method is extremely sensitive to the value assigned to the mass of the piston. A low mass will result in rapid box size oscillations, which are not attenuated very efficiently by the motions of the molecules, while a large mass will give rise to a slow adjustment of the volume and may therefore be computationally inefficient.

3. The force on the piston [See Eq. (3) below] depends on the internal pressure of the system, the value of which depends on the instantaneous kinetic energy of the particles. This means that the velocity $v_{n+1}$ is the solution to an implicit equation, which therefore must be solved iteratively. This has several consequences, including that the computed trajectory is no longer time reversible - a feature that jeopardizes the (extended) energy conservation in long simulations. A set of explicit reversible integrators for the dynamics has been developed by Martyna and coworkers 15-17.

4. The dependence of the internal pressure on the kinetic energy leads to inaccurate determination of the pressure, since the kinetic contribution is derived from the particles’ velocities, which, as shown in Appendix A, deviate from the actual velocities.

The third approach to constant pressure and temperature MD simulations employs the Andersen extended Lagrangian formalism, i.e., it couples the system to a global piston which governs the volume fluctuations of the system. However, instead of using a Nosé-Hoover thermostat and solving the Hamiltonian equations of motion, the temperature is set by solving the Langevin equation 18:

\[
\dot{\dot{r}} = \frac{f(r,t)}{m} - \alpha v + \beta(t). \tag{3}
\]

The Langevin equation describes Newtonian dynamics where the conservative force field $f(r,t)$ is augmented by: (i) a friction force proportional to the velocity with friction coefficient $\alpha$, (ii) thermal white (“delta-function correlated”) noise, $\beta(t)$. The friction and noise terms represent the interactions with the implicit degrees of freedom of the heat bath. In order to satisfy Einstein’s fluctuation-dissipation theorem that relates the friction and noise to each other, it is usually assumed that the noise is Gaussian distributed and has the following statistical properties 19:

\[
\langle \beta(t) \rangle = 0 \tag{5}
\]

\[
\langle \beta(t)\beta(t') \rangle = 2\alpha k_BT\delta(t-t'). \tag{6}
\]

Historically, Langevin stochastic thermostats have been developed in parallel to the Nosé-Hoover deterministic thermostat, in the early 80’s 20. However, it was only in 1995, when Feller et al. proposed to simulate isothermal-isobaric conditions by considering Langevin dynamics for the piston’s equation of motion in Andersen’s extended system 21. This approach was improved a few years later by Kolb and Dünweg, who considered Langevin dynamics for both the particles and piston, and who developed a symplectic time-reversible integrator for this purpose 22. While many of the problems associated with the application of the Nosé-Hoover thermostat for $(N,P,T)$ simulations remained unsolved (especially those originating from the coupling between the movement of the piston and the particles), the idea of simulating the
extended system within the framework of Langevin dynamics appears to offer shorter correlation times and improved sampling.

In this paper we present a new method for Langevin dynamics simulations at constant pressure and temperature. The method, which is both effective and simple to implement, provides improvements compared to the method of Kolb and Dünweg (KD). Key distinctions between our algorithm and others, including KD, lie in the manner by which the displacements of the particles into “physical” and “scaled” components are decoupled. Within the traditional methods these two displacements are defined and separated prior to time-discretization, while our method is based on formulating the equations of motion for an already temporally discretized set of coordinates. Another change that we introduce in the method is to replace the kinetic energy term in the instantaneous pressure with its known thermodynamic average, which is precisely the ideal gas pressure $Nk_BT/V$. This change does not only resolve the aforementioned problems in implementations of Verlet-type integrators to the piston’s equation of motion, it also makes the extended Lagrangian dynamics more consistent with the statistical mechanics of the isothermal-isobaric ensemble that the simulations aims to sample. Finally, we take advantage of the recent advances in numerical integrators for Langevin dynamics and replace the old BBK (Brooks, Brünger, and Karplus) thermostat [20] with the recently introduced G-JF (Grønbech-Jensen and Farago) thermostat [6]. While the former has a simulated temperature that differs by $\Delta T = T_s - T$ from the correct one, the latter exhibits no detectable changes in the configurational sampling statistics as the time step is varied in the entire numerical stability range [6, 22].

The paper is organized as follows: In section II we derive the new method for isothermal-isobaric MD simulations. This section contains both a detailed discussion of the theoretical aspects of the method, as well as a derivation of the algorithm for isothermal-isobaric simulations. The new algorithm is tested against the method of Kolb and Dünweg in section III. For this purpose we present simulation results of both a one-dimensional toy model that can be solved analytically, and a three-dimensional Lennard-Jones system. We conclude the paper in section IV.

II. ISONTERMAL-ISOBARIC LANGEVIN DYNAMICS

A. Statistical mechanical considerations

In his seminal paper on the extended Lagrangian formalism, Andersen studied the statistical mechanics of $N$ particles within a box with a fluctuating volume $V$ subject to a constant external pressure $P$ [3]. He associated the volume fluctuations with the motion of a “piston”, and considered an extended phase space of $2(dN + 1)$ degrees of freedom, including (i) the $N\dot{d}$ coordinates of the particles, $\vec{r}_i$, and their $N\dot{d}$ conjugate momenta $\vec{p}_i$, and (ii) the volume $V$ representing the coordinate of a “piston” along with its conjugate momentum. The derivation of the extended Lagrangian formalism was done in the rather uncommon isenthalpic-isobaric ensemble $(N, P, H)$, where the enthalpy is $H = E + PV$. This is the equivalent of the microcanonical ensemble $(N, V, E)$ for fixed pressure.

The degrees of freedom $\vec{r}_i$ and $V$ in the extended system are not independent of each other because the particle coordinates are adjusted to volume fluctuations via simple scaling during MD simulations. In order to have independent statistical variables, one needs to define the scaled coordinates $\bar{s}_i$

\[ s_{i,\mu} = r_{i,\mu}/L_{\mu}, \]

where $\mu = x, y, z$, $\bar{s}_i = (s_{i,x} s_{i,y} s_{i,z})^T$, and $L_{\mu}$ is the linear size of the simulations box along the $\mu$-axis. For simplicity, we here assume that the simulation box is orthorhombic with fixed aspect ratios, such that $\prod \mu L_{\mu} = V$, and that all the particles have identical mass $m$, except for the piston, which is considered a coordinate with inertial constant $Q$. The “force” acting on the piston is derived from the extended Hamiltonian, which is obtained from the extended Lagrangian via Legendre transformation. It is given by (see Eq. (3.14C) in Ref. [3])

\[ f_P = \frac{1}{Vd} \sum_{i=1}^{N} \left( \bar{f}_i \cdot \bar{r}_i + \frac{\bar{p}_i^2}{m} \right) - P. \]

The transition from the isenthalpic-isobaric into the isothermal-isobaric ensemble requires the introduction of a “thermostat”, and as noted in section II the thermostat can be either “deterministic” (Nosé-Hoover) or “stochastic” (Langevin). In terms of the coordinates $\bar{s}_i$ and $V$, the isothermal-isobaric partition function reads:

\[ Z = \int_0^\infty dV V^N \int_0^1 \prod_{i=1}^{N} ds_{i,\mu} e^{-[U(\{L_{\mu}, s_{i,\mu}\}) + PV]/k_BT} \]

\[ = \int_0^\infty dV \int_0^1 \prod_{i=1}^{N} ds_{i,\mu} e^{-[U(\{L_{\mu}, s_{i,\mu}\}) + PV - Nk_BT \ln V]/k_BT}. \]

This partition function can be interpreted as if governing the canonical ensemble of a system consisting of $N$ particles confined to a three-dimensional unit cube ($0 \leq s_{i,\mu} \leq 1$), and a piston moving along an infinite line ($0 < V < \infty$), with the potential energy given by

\[ U_{\text{eff}}(\{s_{i,\mu}\}, \{L_{\mu}\}) = U(\{L_{\mu}, s_{i,\mu}\}) + PV - Nk_BT \ln V. \]

Notice that the partition function defined by Eq. (9) includes summation only over the spatial degrees of freedom (of the particles and piston), but not over their conjugate momenta. This deviation from Andersen’s extended Lagrangian formalism, where both the coordinates and momenta were included in the partition sum,
deserves an explanation. Andersen’s method describes Newtonian dynamics within a microcanonical ensemble. In this ensemble, the kinetic and potential energies are coupled by energy conservation. In contrast, Langevin dynamics occurs within an open system in contact with a heat bath. In this canonical ensemble, the kinetic and potential energies are decoupled, and the degrees of freedom of the coordinates can be integrated separately from their associated momenta. The momenta degrees of freedom follow a Maxwell-Boltzmann Gaussian distribution, while the coordinates (\(s_i\) and \(V\)) are governed by the Boltzmann distribution corresponding to \(U_{\text{eff}}\). The separation of the ensemble into two sub-spaces, corresponding to the coordinates and their associated momenta, is important because the goal of constant temperature and pressure simulations is to sample the phase space of the coordinates correctly. The momenta, i.e., the velocities, are only used in these simulations as a mean to access the simulated kinetic temperature. The average kinetic energy is a reasonable measure of the temperature, but not a good one in discrete-time because of the second order (in \(dt\)) deviation between the measured velocity relative to the trajectory of the corresponding coordinate (see Appendix A). Thus, numerical measures involving velocity are not reliable for non-vanishing time steps. In constant volume simulations, this problem is avoided if the Langevin dynamics is to sample the phase space of the ensemble into two sub-spaces, corresponding to the trajectory of the corresponding coordinate \(U\). In this ensemble, the kinetic and potential energies are decoupled, and the degrees of freedom relative to the trajectory of the corresponding coordinate \(U\) is accomplished through the scaled (normalized) coordinate \(\tilde{r}_i\), acting on this particle, is given by \(P\), which (in the velocity-Verlet form) is expressed by the following equations to calculate the coordinate (i.e., volume) \(V^{n+1}\) and velocity \(\dot{V}^{n+1}\) at time \(t_{n+1} = t_n + dt\) (See Eqs. (4)-(8) in Ref. 23):

\[
V^{n+1} = V^n + \tilde{b} dt \dot{V}^n + \frac{\tilde{b} dt^2}{2Q} f_p^n + \frac{\tilde{b} dt}{Q} \tilde{\beta}^{n+1},
\]

where \(\tilde{a} = 1 - \frac{\tilde{b} dt^2}{2Q}\)

\[
\tilde{b} = 1 + \frac{\tilde{a} dt}{Q},
\]

and \(\tilde{\beta}^n\) is a normally distributed random number with zero mean, and autocorrelation \(\langle \tilde{\beta}^n \tilde{\beta}^m \rangle = 2\tilde{a} k_B T dt \delta_{m,n}\).

2. Dynamics of the particles

The variation of the volume causes complications for the dynamics of the particles, which reside within the confinements of the defined, yet variable, volume. These complications are particularly apparent in systems with periodic boundary conditions since the simulated volume is associated with a lattice constant of a simulation box and not with a physical location of an actual piston. Thus, in order to preserve the translational invariance of the equations of motion in a bulk system with periodic boundary conditions, it is necessary to globally couple the dynamics of the volume to all the particles, regardless of particle location in the simulation cell, such that relative distances in the system are preserved. This is accomplished through the scaled (normalized) coordinate \(s_{i,\mu} = r_{i,\mu} / L_\mu\), which is understood to be constant for a simple expansion or contraction of \(L_\mu\). However, the physical velocity and acceleration of the coordinate \(r_{i,\mu}\) can then not be translational invariant without modifications. Andersen’s solution to the problem is to investigate the derivative

\[
\tilde{r}_{i,\mu} = \tilde{s}_{i,\mu} L_\mu + \tilde{s}_{i,\mu} \dot{L}_\mu.
\]

In Eq. (17) we can see the separation of two identifiable components to the motion: (i) the dynamics of the particle relative to the simulation cell (first term on the r.h.s.), and (ii) the dynamics due to the motion of the simulation cell (second term on the r.h.s.). Thus, defining

\[
v_{i,\mu} = \dot{s}_{i,\mu} L_\mu
\]
as the relevant physical velocity of the coordinate \(r_{i,\mu} = s_{i,\mu}L_\mu\) makes the particle dynamics invariant to the origin of the coordinate system – an essential necessity for meaningful dynamics. While this elegant observation has led to the advanced formulations of both deterministic \([10, 12, 14, 15]\) and stochastic \([21, 22]\) methods for NPT simulations, the inherent problem of time discretization persists.

We now arrive at the core of the derivation of the new method. We simplify the notation for brevity in the rest of this subsection, such that, e.g., the coordinate \(r\) refers to \(r_{i,\mu}, L_\mu\) to \(L_\mu\), etc, unless specifically indicated otherwise.

We reevaluate the particle equations of motion in discrete-time, starting with the definition of the scaled coordinate Eq. (17). The total particle displacement \(\Delta r_{n+1} = r_{n+1} - r_n\) in one time step is given by

\[
\Delta r_{n+1} = \int_{t_n}^{t_{n+1}} \frac{d}{dt}(sL)\ dt' = s_{n+1}L^{n+1} - s_nL^n
\]

(19)

\[
= (s_{n+1} - s_n)\frac{L^{n+1} + L_n}{2} + \frac{s_{n+1} + s_n}{2}(L^{n+1} - L^n). 
\]

(20)

We use the analogy between Eqs. (17) and (20) to define the relevant physical, spatially invariant, discrete-time particle displacement \(\Delta q_{n+1}\) from the first term in Eq. (20)

\[
\Delta q_{n+1} = (s_{n+1} - s_n)\frac{L^{n+1} + L_n}{2}.
\]

(21)

Notice that \(\Delta q \to sLdt\) for \(dt \to 0\), consistent with the usual continuous-time definition of the relevant velocity mentioned above \(v = sL\) \([15]\). Thus, we conclude that the discrete-time particle dynamics must involve the physical coordinate \(q\) and an associated velocity \(v\), which must relate to the discrete-time displacement through

\[
\Delta q_{n+1} = \int_{t_n}^{t_{n+1}} v\ dt'.
\]

(22)

The corresponding discrete-time velocity change \(\Delta v_{n+1} = v_{n+1} - v_n\) is obtained through the dt-integrated Langevin equation

\[
\int_{t_n}^{t_{n+1}} [m\ddot{v} + \alpha v]\ dt' = \int_{t_n}^{t_{n+1}} [f(r, t') + \beta(t')]\ dt',
\]

(23)

which, using (22) and with no approximation, can be written

\[
m\Delta v_{n+1} + \alpha\Delta q_{n+1} = \int_{t_n}^{t_{n+1}} f\ dt' + \beta_{n+1},
\]

(24)

where we have defined the Wiener process

\[
\beta^n = \int_{t_n}^{t_{n+1}} \beta(t')\ dt'.
\]

(25)

such that

\[
\langle \beta^n \rangle = 0, \quad \langle \beta^n \beta^m \rangle = 2\alpha k_BT dt \delta_{n,m}.
\]

(26)

(The noise autocorrelation reads with full notation: \(\langle \beta_{i,\mu}^n \beta_{j,\nu}^m \rangle = 2\alpha k_BT dt \delta_{i,j} \delta_{\mu,\nu}\).) Notice that the introduction of the discrete-time Langevin equation in Eq. (24), for linking the coordinate displacement \(\Delta q\) with its velocity change \(\Delta v\), ensures physically meaningful discrete-time evolution.

Starting from Eq. (22), and following our previous work \([3]\), we now choose the time-reversible relationship between the relative displacement and change in the associated velocity:

\[
\Delta q_{n+1} = b dt v_n + \frac{b dt}{2m} \int_{t_n}^{t_{n+1}} f\ dt' + \frac{b dt}{2m} \beta_{n+1}
\]

(28)

where

\[
b = \frac{1}{1 + \frac{\alpha dt}{2m}}.
\]

(29)

Equations (28) and (24) constitute a set of equations for determining \(\Delta q_{n+1}\) and \(\Delta v_{n+1}\). We then approximate the dt-integrals over the deterministic force \(f\) such that all terms in the equations become at least second order correct in \(dt\) (i.e., consistent with the traditional Verlet methods), which yields:

\[
\Delta q_{n+1} = b dt v_n + \frac{b dt^2}{2m} f_n' + \frac{b dt}{2m} \beta_{n+1}
\]

(30)

\[
\Delta v_{n+1} = -\frac{\alpha}{m} \Delta q_{n+1} + \frac{dt}{2m} (f_n + f_{n+1}) + \frac{1}{m} \beta_{n+1}
\]

(31)

These are explicit discrete-time equations for evaluating the evolution of the coordinates \(q^n\) and \(v^n\).

In order to express the equations in the most useful form for molecular simulations, we use the relationship \(\nu^n = s^nL^n\) (Eq. (7)) to combine Eqs. (21) and (30) for a direct expression of the dynamics of the physical coordinate \(\nu^n\):

\[
\nu_{n+1} = \frac{L_{n+1}^n}{L^n} \nu_n + \frac{2L_{n+1}^n}{L_{n+1}^n + L^n} b dt [v_n + \frac{dt}{2m} f_n + \frac{1}{2m} \beta_{n+1}].
\]

(32)

We also insert Eq. (30) into Eq. (31) in order to obtain an explicit equation for the dynamics of the velocity \(v^n\):

\[
v_{n+1} = a v_n + \frac{dt}{2m} (a f_n + f_{n+1}) + \frac{b}{m} \beta_{n+1},
\]

(33)
where

\[ a = \frac{1 - \frac{\alpha dt}{2m}}{1 + \frac{\alpha dt}{2m}} \]  

Equations 32 and 33 are the Verlet-type equations for particle updates in the stochastic G-JF thermostat/barostat, given a change \( \Delta L^{n+1} = L^{n+1} - L^n \) in simulation box dimension during the time step \( dt = t_{n+1} - t_n \). Notice that the velocity equation depends only indirectly on the change in the simulation dimension \( L \) through the force \( f_i^n = f(r_i^n, t_n, L^n) \).

To summarize, given \( r_i^n, v_i^n, f_i^n, f_P^n, V^n \), and \( V_n \), the discrete-time dynamics evolves according to the following protocol:

1. Compute \( V^{n+1} \) (and \( L_{\mu}^{n+1} \)) using Eq. (13).
2. Compute \( r^{n+1} \) (all \( r_i^{n+1} \)) using Eq. (32).
3. Evaluate the new forces \( f^{n+1} = f(r^{n+1}, t_{n+1}, L^{n+1}) \) (all \( f_i^{n+1} \)), and \( f_P(\{r_i^{n+1}\}, t_{n+1}, \{L_{\mu}^{n+1}\}) \).
4. Compute \( V^{n+1} \) using Eq. (34) and \( v^{n+1} \) (all \( v_i^{n+1} \)) using Eq. (33).

We reemphasize that the coordinates \( r^n, v^n, f^n \) here refer to each Cartesian coordinate of each particle, and that \( L \) refers to \( L_{\mu} \), such that \( V = \prod_{\mu} L_{\mu} \) is the volume of an isotropically varying, orthorhombic simulation box.

### III. TESTING THE ALGORITHM

In order to test the method we have applied it to two characteristic systems with a specific eye on the robustness against time-step variations. We compare our results with those generated by the KD method [22], which represents state-of-the-art of a sound approach to Langevin dynamics NPT simulations in atomic and molecular ensembles. The first system is a particular non-trivial, one-dimensional particle model for which we can analytically derive measurable thermodynamic quantities. This model therefore serves as a strict benchmark for the statistical accuracy of a numerical test simulation. The second system is the foundational model system in computational statistical mechanics, namely the three-dimensional ensemble of particles interacting with a Lennard-Jones force field. In this latter case, we do not have analytical expressions for the statistical measures, but we investigate the measures for different values of the discrete time step, and from that infer the quality of the applied numerical methods.

#### A. One-dimensional model system

We consider a one dimensional system of normalized length \( L \) (characteristic length \( r_0 \)) with periodic boundary conditions. \( N \) identical particles are located in order at \( \{x_1, x_2, \ldots, x_N : x_i < x_{i+1} \} \) such that the periodic boundary conditions ensure two neighbors for each particle (i.e., \( x_{i\pm N} = x_i \pm L \)). Each particle interacts with its two neighbors via a pair-potential that depends on the normalized pair distance \( r \). Expressing the energy in units of the thermal energy \( (E_0 = k_B T) \), the normalized pair-potential \( u(r) \) [related to the physical potential via \( U(r_0r) = E_0 u(r) \)] reads

\[ u(r) = \frac{\epsilon}{r} + \frac{1}{2} \ln r. \]  

The pair potential \( u(r) \) consists of two contributions: a repulsive part \( (\epsilon > 0) \), inversely proportional to \( r \), and an attractive logarithmic part. The latter may represent an entropic potential of mean force resulting from implicit degrees of freedom. Considering the isobaric-isothermal ensemble \( (N, P, T) \) (where \( P \) denotes the one-dimensional pressure, i.e., the force, and is expressed in units of \( E_0/r_0 \)), the partition function of the system is given by

\[ Z = \int dL \prod_{i=1}^{N} dx_i \exp \left[ - \sum_{i=1}^{N} u(x_{i+1} - x_i) - PL \right] \]  

Switching to the set of variables \( r_i = x_{i+1} - x_i \), the partition function reads

\[ Z = \prod_{i=1}^{N} \int_{0}^{\infty} dr_i \exp \left[ - \sum_{i=1}^{N} u(r_i) - P \sum_{i=1}^{N} r_i \right] \]

\[ = \left[ \int_{0}^{\infty} dr \exp(-\epsilon r + P r) \right]^N \]

\[ = 2 \int_{0}^{\infty} dy \exp(-\epsilon y^2 + Py^2)^N, \]  

where the last equality has been obtained by setting \( y = \sqrt{\frac{P}{\epsilon}} \). The value of the last integral is known, giving

\[ Z = \left[ \sqrt{\frac{P}{\epsilon}} e^{-2\sqrt{P\epsilon}} \right]^N. \]  

The normalized Gibbs free energy is given by \( G = -\ln Z \), and the mean nearest neighbor particle normalized distance, \( \langle l \rangle \), is then derived by

\[ \langle l \rangle = \frac{\langle L \rangle}{N} = -\frac{1}{N} \frac{\partial G}{\partial N} = \frac{1}{2P} + \sqrt{\frac{\epsilon}{P}}. \]  

The variance of the normalized length distribution is given by

\[ \sigma_l^2 = \frac{\left( \langle L^2 \rangle - \langle L \rangle^2 \right)}{N} = -\frac{\partial \langle l \rangle}{\partial P} = \frac{1}{2P^2} + \sqrt{\frac{\epsilon}{2P}}. \]  

For a system of \( N = 1000 \) particles, we simulate the evolution for a normalized transient time of \( 4.8 \times 10^4 \) units before producing statistical averages of \( \langle l \rangle \) and \( \sigma_l \) over the next \( 4.8 \times 10^5 \) normalized time units. Figures 3a and 3b show the resulting data for both the G-JF
FIG. 1: Results for $\epsilon = 10$. Simulated average length (a) and standard deviation $\sigma_l$ (b) for several values of applied 1D pressure (force) $P$. Markers represent the G-JF method of this paper (solid marker •) and the KD method (open marker ◦). At small time steps both methods produce the correct analytical values given by Eqs. (39) and (40). All simulations were done with $Q = 10^{-2}$ and $\tilde{\alpha} = 10^{-2}$. Lines serve as guides to the eye.

B. Three-dimensional Lennard-Jones model system

We now consider the simplest possible well-known system in the modeling of materials and liquids, namely a three-dimensional ensemble of identical spherical particles. Each particle has a normalized mass $m = 1$ (in units of $m_0$) and normalized friction coefficient $\alpha = 1$, and they all interact through the normalized potential $u(r)$ given by the physical pair-potential $U(\{r_{ij}\}) = E_0u(r)$, where $r = |\vec{r}|$ is the normalized pair-distance (in units if the characteristic length $r_0$) and $E_0$ is the characteristic energy. The normalized pair-potential reads

$$u(r) = \begin{cases} r^{-12} - 2r^{-6} & , \quad 0 < r \leq r_s \\ a_4(r - r_c)^4 + a_8(r - r_c)^8 & , \quad r_s < r < r_c \\ 0 & , \quad r_c \leq r \end{cases}$$

(41)
where

\begin{align}
    r_s &= \left( \frac{13}{7} \right)^{1/6} \approx 1.108683 \\
    r_c &= r_s - \frac{32u(r_s)}{11u'(r_s)} \approx 1.959794 \\
    a_4 &= \frac{8u(r_s) + (r_c - r_s)u'(r_s)}{4(r_c - r_s)^4} \\
    a_8 &= \frac{-4u(r_s) + (r_c - r_s)u'(r_s)}{4(r_c - r_s)^8}.
\end{align}

This function (see Fig. 3) is a short-range splined Lennard-Jones potential with continuity through the second derivative at \( r = r_s \) and continuity through third derivative at \( r = r_c \).

Conducting NPT simulations on a cubic system with \( N = 864 \) particles, we optimize the relaxation of the barostat degree of freedom \( V \) by choosing small values for the inertia \( Q \). By inspection, we find that values in the range \( Q = 10^{-4} \) and \( Q = 10^{-5} \) represent efficient relaxation. We also, by inspection, conclude that a small friction coefficient \( \dot{\alpha} = 10^{-4} \) helps relax the system (although this seems to be a weak effect) and, therefore, choose this value for our simulations. We have further chosen two characteristic normalized temperatures, for both solid \( (k_B T/E_0 = 0.3) \) and liquid \( (k_B T/E_0 = 0.7) \) phases. Finally, we have studied three different applied pressures \( (P = 0.01, 0.1, 1.0) \) (expressed in units of \( E_0/r_0^3 \)), and vary the discrete normalized time step \( dt \) in the entire range of stability to observe the behavior of the numerical methods. We only show the \( P = 0.1 \) data here since the results of all three applied pressures show the same characteristics. All statistical data are obtained by initiating the system in a perfect close-packed crystal near a zero-temperature ground state. We then simulate at least \( 2 \times 10^5 \) normalized time units before averages are acquired over the next \( 2 \times 10^5 \) time units. The normalizing time \( \tau_0 \) is given by \( E_0 \tau_0^2 = m_0 r_0^2 \). All the left axes of the figures display absolute results, while all the right vertical axes display the percentage deviation from the \( dt \to 0 \) value of the quantity shown in the plot.

Figure 4 shows the data for the volume \( V \) of the simulation box \( (Q = 10^{-4}) \): Fig. 4a; \( Q = 10^{-5} \): Fig. 4b) and its fluctuations \( (Q = 10^{-4}) \): Fig. 4c; \( Q = 10^{-5} \): Fig. 4d) as a function of the time step for a solid phase at \( k_B T/E_0 = 0.3 \) and external pressure \( P = 0.1 \). The new G-JF barostat results are displayed as solid markers (●), while the comparison KD method results are shown with open markers (○). The data clearly shows that the G-JF results are nearly independent of the time step \( dt \) for both the average volume and the corresponding fluctuations. In comparison, the KD method exhibits a consistent, albeit weak, increase in average volume. More dramatic is the increasing deviation of the volume fluctuations in the KD method. For \( Q = 10^{-4} \), this can be in excess of 10%, while we observe up to 70% discrepancy for \( Q = 10^{-5} \). Such discrepancies clearly change not only measured thermodynamic properties such as the elastic bulk modulus and heat expansion coefficient, but also the structure of the material under investigation. For example, close inspection (not shown) of the KD simulation shows that the excessive volume fluctuations induce crystal defects into the material for large \( dt > 0.016 \), before the numerical instability is found for \( dt \approx 0.019 \). Notice that the results of both methods converge to the same values in the KD method results are shown in Fig. 5. The results of the G-JF method are also here unimpressed with the simulated time step throughout the stability ranges, while the KD method shows a characteristic positive deviation. Since the KD method coincides with the BBK thermostat \( 20 \) when the volume is constant, this result is entirely expected in light of our previous work on the G-JF thermostat and comparisons \( 23 \) to other thermostats, including BBK.

These results are also consistent with the kinetic temperature \( T_k \) measurements shown in Figure 6 where \( T_k \) is defined as

\[ T_k = \frac{1}{3Nk_B} \sum_{i=1}^{N} \langle \dot{v}_i^2 \rangle. \]

In the G-JF method, the kinetic temperature decreases with increasing \( dt \). This is anticipated since, as mentioned in the introduction, it is known that the momentum \( m v_i^0 \) is not the conjugate variable to \( r_i^0 \) for \( dt > 0 \) (see, e.g., Refs. \( 3, 6, 23 \) as well as Appendix A below), and that the discrete-time second order approximations \( (v_i^0 \text{ and } V^0) \) to the velocity variables from the
The central difference approach in the Verlet formalism leaves kinetic and configurational measures mutually inconsistent. Thus, the very good configurational sampling properties of the G-JF method seen from the measurements of, e.g., volume and enthalpy (along with their fluctuations) inevitably mean that a measurement [such as $T_k$ in Eq. (47)] derived from the (incorrect) velocities will be incorrect to second order in $dt$. This is a simple consequence of $v_i^n$ being an approximation to the true velocity of $r_i^n$, which cannot be obtained. Figure 4 further displays the measured kinetic temperature $T_k$ for the KD method, and it is apparent that this quantity appears to confirm the required temperature, which is consistent with the incorrect configurational properties of this method seen for volume and enthalpy as $dt$ is increased. This artifact of discrete time emphasizes that one should refrain from using kinetic measures as reliable quantities in these types of simulations.

We now show results for a liquid phase at $k_B T/E_0 = 0.7$. Otherwise, all system and simulation parameters are exactly as for the $k_B T/E_0 = 0.3$ results shown above. The liquid phase is validated by structural analysis and the measured diffusion constant, which we derive from the Einstein definition

$$D = \frac{1}{N} \sum_{i=1}^{N} \lim_{n \to \infty} \langle V \rangle^2 \frac{(s_i^n - \bar{s}_i^n)^2}{6n dt}. \tag{48}$$

We use time averages over $n dt = 200,000$ for all chosen values of $dt$, and $s_i^n$ is understood to extend beyond the interval $0 \leq s < 1$ in this expression [24]. Figure 5 displays the non-zero measured diffusion coefficient of the liquid state as a function of the time step. It is clear that the migration at this temperature and pressure is weak, and that the diffusion measurement is noisy. Even so, the figure demonstrates that both G-JF and KD methods exhibit diffusion coefficients reasonably independent of the choice of the size of time step, although there may be a hint of a slight increase for the KD method for increasing $dt$.

Figure 5 shows the $k_B T/E_0 = 0.7$ data for the volume $V$ of the simulation box. The KD results for both average and fluctuation of the volumes exhibit significant increases with $dt$, consistent with the comparable $k_B T/E_0 = 0.3$ data. The G-JF results are much less impressed by the time step $dt$, but there is a small tendency for the volume and its fluctuations to decrease with increasing $dt$. However, the overall impression is clearly that the G-JF method is significantly less dependent on variations in $dt$ than the KD method is.

The $k_B T/E_0 = 0.7$ data for the potential energy are shown in Fig. 6 The KD results for this liquid phase exhibit the typical BBK behavior that was also seen in Fig. 4 for the solid phase at $k_B T/E_0 = 0.3$. In comparison, the potential energy shows only a slight decrease in both average and fluctuations for the G-JF method. It is again clear that the G-JF method produces simulated matter with configurational properties nearly independent of $dt$. The uncertainty on the acquired averages can be assessed from the associated standard deviations and the averaging time. We here also include a multiple of simulation data for the same parameters in order to indicate the magnitude of the statistical error that should

![Figure 4](image1.png)

![Figure 5](image2.png)
be associated with the presented standard deviations.

We confirm that the kinetic measurements of temperature and its fluctuations behave similarly in the liquid and solid phases by comparing Figs. 6 and 10. The latter shows the data for $k_B T / E_0 = 0.3$, which again demonstrates the signature of the momentum $mv^2$ not exactly being the conjugate variable to $r^n$. Thus, also for the liquid phase, we observe that the calculated discrete-time kinetic temperature is progressively short of the actual temperature of the configurational sampling statistics that can be inferred from the potential energy measurements in Figure 9.

We finally turn to investigating the pressure, which is the principal focus of the barostat. Clearly, one should expect that the average internal pressure $\langle P \rangle$ is controlled and equal to the imposed external pressure $P$, since this is the immediate purpose of the barostat. It is important to note that the internal pressure is defined differently in the G-JF and KD methods (see discussion above in section II). The former uses the canonical ensemble definition [see Eq. (11)]

$$P_{\text{can}} = \frac{1}{3V} \sum_{i=1}^{N} \bar{f}_i \cdot \bar{r}_i + \frac{N k_B T}{V}, \quad (49)$$

while the latter targets the microcanonical ensemble expression [see Eq. (8)]

$$P_{\text{micro}} = \frac{1}{3V} \sum_{i=1}^{N} \bar{f}_i \cdot \bar{r}_i + \frac{1}{3V} \sum_{i=1}^{N} m \bar{v}_i^2, \quad (50)$$

For each method we inspect the statistics of the relevant internal pressure. The results for the solid phase simulations at $k_B T / E_0 = 0.3$ are displayed in Figure 11. From the data, it is obvious that the imposed pressure is correctly adopted by the G-JF method presented here. The KD method, however, displays a curious and perhaps significant deviation from the expected. The origin of these deviations is the time reversible discretization used in the KD method (see sequential steps (1)-(7) in Sec. V of Ref. [22]), that applies a trapezoidal approximation $(r^n f^n + r^{n+1} f^{n+1})/2$ to the configurational pressure contribution [first term on rhs of Eq. (50)], while
consistent with the enforced value $P_n$ [(c) and (d)] for $Q = 10^{-4}$ [(a) and (c)] and $Q = 10^{-5}$ [(b) and (d)]. Markers represent the G-JF method of this paper (solid •) and the KD method (open ◦). Horizontal dotted lines are leveled at $\langle T_k \rangle = 0.3$ [(a) and (b)], and at $\sigma_{T_k}$ for $Q = 10^{-4}$ and $dt = 0.001$ [(c) and (d)]. All figures show axes with absolute quantities on the left and percentage deviation on the right axes.

The new G-JF method is simple and stable, and simulating instantaneous pressure (at times $t_n$) with its fluctuations as a function of the time step for the liquid phase at $k_B T/E_0 = 0.7$. The overall behavior of the methods is the same for liquid and solid phases with direct averages of the instantaneous pressure $\langle E_t \rangle$ being significantly depressed for the KD method as $dt$ is increased. We also see that the fluctuations of the KD pressure is fairly independent of $dt$ for $Q = 10^{-4}$, while the fluctuations of the pressure $P_H$ increase dramatically for $Q = 10^{-5}$. The G-JF method is generally robust, although we do observe some increase in pressure fluctuations for $Q = 10^{-5}$.

We note that improved statistical accuracy of kinetics using half-step velocities $v^{n+\frac{1}{2}}$ in the so-called leapfrog versions of the Verlet method have been investigated for deterministic Nosé-Hoover control of temperature and pressure. However, while the half-step velocities may be able to produce better consistency for the averaged kinetic temperature in deterministic dynamics, these approaches may neither translate to stochastic dynamics nor resolve the fundamental question of calculating instantaneous pressure (at times $t_n$) for statistical averages and fluctuations, as illustrated above.

**IV. DISCUSSION**

We have presented and demonstrated a new thermostat-barostat pair for simulating atomic and molecular dynamics with periodic boundary conditions. The new G-JF method is simple and stable, and simu-
lations of thermodynamic properties produce data with very little dependency on the applied numerical time step. We have investigated the method in the context of two characteristic models - a one-dimensional toy model with known statistical solutions, and the classical three-dimensional Lennard-Jones material, simulated in both crystalline and liquid phases. In all cases the G-JF method behaves extremely well for measured averages as well as for their fluctuations. In comparison, the state-of-the-art KD method, which is also representative of other commonly used methods, may exhibit significant deviations in both averages and fluctuations for increasing time steps. As we have emphasized throughout this paper, and specifically in Appendix A, it is crucial to appreciate that discrete time invalidates the conjugate relationship between the coordinate \( r \) and its simulated velocity \( v \). Consequently, one cannot expect accurate simulation measures for both configurational and kinetic quantities using any given method. This interesting and essential feature becomes apparent when comparing the behavior of kinetic and potential energies as a function of time step variations. We submit that the G-JF thermostat and barostat are advantageous in that they consistently provide proper configurational properties (such as Boltzmann distributions, Einstein diffusion, potential energy, pressure, system volume, as well as their fluctuations), while leaving kinetic measures (such as measured kinetic energy and the derived kinetic temperature) with predictable deviations. In contrast, most other methods (e.g., KD, Nosé-Hoover, etc.) enforce the expected kinetic measures, thereby sacrificing the accuracy of proper configurational sampling. The latter is unfortunate, since most molecular simulations are conducted in order to obtain configurational information.

We close by noting that the G-JF method is easily extended to non-isotropic volume adjustments, and that the algorithm is not only simple, but also in a form that makes it easy to implement into existing molecular dynamics codes that have thermodynamic temperature and pressure control. Specifically, the method can also conveniently be expressed in the so-called leap-frog and position-Verlet forms, as outlined for the thermostat in Ref. 23.

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Appendix A: Discrete-time relationship between position and velocity

The continuous-time expectation, that the momentum \( p^n = m v^n \) is the conjugate variable to the spatial coordinate \( r^n \), is not fulfilled in discrete-time Verlet methods. This unfortunate consequence of time discretization has significant implications for the use and interpretation of
simulations, and it can be illuminated by considering a the simple analysis of a simulated harmonic oscillator

\[ m \ddot{r} = -\kappa r, \] (A1)

where \( \kappa > 0 \) is a Hooke’s spring constant. The continuous-time solution to this equation is, of course, 
\( r(t) \propto \exp(\pm i \Omega_0 t) \) and \( v(t) = \dot{r} = \pm i \Omega_0 r(t) \), where \( \Omega_0^2 = \kappa/m \). (We use \( i \) for complex notation in this Appendix.)

The discrete-time Verlet equations for Eq. (A1) are found from Eqs. (1) and (2):

\[ r^{n+1} = 2r^n - r^{n-1} - \frac{\kappa dt^2}{m} r^n \] (A2)
\[ v^n = \frac{r^{n+1} - r^{n-1}}{2dt}. \] (A3)

The solution is

\[ r^n \propto \exp(\pm i \Omega_V n dt) \] (A4)

with the oscillation frequency \( \Omega_V \) of the discrete-time Verlet oscillator given by

\[ \sin(\Omega_V dt) = \Omega_0 dt \sqrt{1 - \left( \frac{\Omega_0 dt}{2} \right)^2} \] (A5)
\[ \cos(\Omega_V dt) = 1 - \frac{\Omega_0^2 dt^2}{2} \] (A6)

for \( \Omega_0 dt \leq 2 \) (see, e.g., Refs. [6, 27]). Inserting Eq. (A4) into Eq. (A3), we get the Verlet velocity

\[ v^n = \pm \sin(\Omega_V dt) \frac{\Omega_V}{\Omega_0 dt} i \Omega_V r^n, \] (A7)

which, when compared to Eq. (A4), shows that the Verlet velocity is always depressed in magnitude compared to the true velocity of \( r^n \) when \( \Omega_V dt \neq 0 \). For the extreme case of the stability limit, \( \Omega_0 dt = 2 \) (\( \Omega_V dt = \pi \)), we find \( v^n = 0 \), even if \( r^n = (-1)^n r^0 \neq 0 \) changes every time step. It is obvious from the above analysis that the Verlet velocity is not an accurate representation of the discrete time trajectory, and that any kinetic measure, such as kinetic energy, temperature or diffusion derived from velocity autocorrelations, cannot reliably represent the behavior of the simulated configurational evolution see also discussion in the appendix of Ref. [25].

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