Role of the sampling weight in evaluating classical time autocorrelation functions

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We analyze how the choice of the sampling weight affects the efficiency of the Monte Carlo evaluation of classical time autocorrelation functions. Assuming uncorrelated sampling or sampling with constant correlation length, we propose a sampling weight for which the number of trajectories needed for convergence is independent of the correlated quantity, dimensionality, dynamics, and phase-space density. In contrast, it is shown that the computational cost of the “standard” intuitive algorithm which samples directly from the phase-space density may scale exponentially with the number of degrees of freedom. Yet, for the stationary Gaussian distribution of harmonic systems and for the autocorrelation function of a linear function of phase-space coordinates, the computational cost of this standard algorithm is also independent of dimensionality.

Introduction: Time-correlation functions. Many dynamical properties of stationary systems as well as the response of such systems to weak perturbations can be inferred from time autocorrelation functions \([1, 2]\). Examples include the optical absorption line shapes computed from the dipole time autocorrelation function, the diffusion coefficient computed from the velocity time autocorrelation function, and various relaxation properties \([3]\). More general time correlation functions are in fact the principal ingredients of semiclassical \([4, 5]\) and path-integral \([6–11]\) calculations of quantum dynamical properties. Trajectory-based methods for computing time correlation functions, however, often become too expensive in many-dimensional systems. Yet, dimensionality-independent algorithms have been found for special correlation functions, such as classical \([12]\) and semiclassical \([13]\) fidelity \([14]\). Motivated by the success in these special cases and by the importance of correlation functions in many areas of physics, in this Letter we explore how these functions can be computed more efficiently in general. In particular, we propose a sampling weight for which the number of trajectories needed for convergence of any classical normalized time autocorrelation function is independent of dimensionality.

Quantum mechanically, the unnormalized time autocorrelation function \(C_\mu^{QM}(t)\) of a vector operator \(\hat{A}\) may be written as

\[
C_\mu^{QM}(t) = \text{Tr}(\rho_0 \hat{A}^\dagger \cdot \hat{A} t),
\]

where \(\rho_0\) is the density operator of the state, \(\hat{A}^\dagger\) is the operator evaluated at time \(t = 0\), \(\hat{A} t = e^{iHt/\hbar} \hat{A} e^{-iHt/\hbar}\) is the operator \(\hat{A}\) evolved with Hamiltonian \(\hat{H}\) for time \(t\), and subscript \(\mu\) emphasizes that the correlation function is not normalized. The classical analog \(C_\mu^{CL}(t)\) of the quantum correlation function \([1]\) is

\[
C_\mu^{CL}(t) = h^{-D} \int dx \rho_0(x) \ A^0(x) \cdot A^t(x),
\]

where \(x := (q, p)\) is the 2D-dimensional phase-space coordinate, \(\rho_0(x)\) is the initial phase-space density, \(A^0(x)\) is the classical observable \(A\) evaluated at time \(t = 0\), and \(A^t(x) = e^{-Lt} A^0(x)\) is this function \(A\) evolved classically for time \(t\) with the Liouville operator \(L = \{H, \cdot\}\). Note that besides a 3-dimensional vector (such as the molecular dipole \(\mu\)), \(A\) can also be a scalar \((A)\) or a higher-dimensional phase-space vector. To make the connection between classical and quantum mechanical expressions explicit, the phase-space volume is measured in units of \(h^D\). Since our focus is on classical correlation functions, superscript CL will be omitted from this point forward.

The shape of the autocorrelation function is often more interesting than its overall magnitude \([15]\), and hence one often computes the time autocorrelation \(C(t)\) which is normalized with respect to its initial value:

\[
C(t) = \frac{C_\mu(t)}{C_\mu(0)}. \tag{3}
\]

Algorithms. Most common methods for evaluating Eqs. \((2)\) and \((3)\) in many-dimensional cases are based on classical trajectories. Two general approaches are currently used \([16]\): 1) the direct approach in which initial conditions for many trajectories are sampled from the stationary distribution \(\rho\) and the trajectories are subsequently evolved simultaneously in time; and 2) the single-trajectory approach in which only one trajectory is evolved in time and the desired autocorrelation function is computed as an average of many correlation functions computed using the same trajectory but initiated at different times. The direct approach is more general and does not require the ergodicity of the time evolution, whereas the single trajectory approach is generally simpler as it avoids explicit sampling of \(\rho\). Here we explore modifications of the direct approach using generalized sampling weights.

We start by expressing the correlation function \((2)\) in terms of trajectories,

\[
C_\mu(t) = h^{-D} \int dx \rho(x) A(x) \cdot A(x^{-t}), \tag{4}
\]

where \(x^t := \Phi^t(x_0)\) is the phase-space coordinate at time \(t\) of a trajectory of the Hamiltonian flow \(\Phi^t\) with initial condition \(x_0\). We further rewrite Eq. \((3)\) in a form suitable for Monte Carlo evaluation, i.e., as an average

\[
\langle E(x_0, t) \rangle_W := \frac{\int dx_0 E(x_0, t) W(x_0)}{\int dx_0 W(x_0)}, \tag{5}
\]
where the positive definite function $W$ is the sampling weight and $E$ is the estimator. In the Monte Carlo method, average (5) is evaluated numerically as an average

$$E_W(N, t) = \frac{1}{N} \sum_{j=1}^{N} E(x_0^j, t)$$

(6)

over $N$ trajectories whose initial conditions $x_0^j$ are sampled from the weight $W$.

The convergence rate of the sum (6) usually depends on $W$. Among many possible weights $W$, the form of Eq. (4) immediately suggests the following three: $W(x) = \rho(x)$, $W(x) = \rho(x)|A(x)|$, and $W(x) = \rho(x)A(x)^2$. These three weights lead to three different algorithms, which may be generally written as

$$C_{a,W}(t) = I_W \left( E_W(x_0, t) \right)_W,$$

(7)

where $I_W := \hbar^{-D} \int W(x) dx$ denotes the norm of $W$ and the estimators are

$$E_\rho(x_0, t) = A(x_0) \cdot A(x-t),$$

$$E_{\rho|A}(x_0, t) = \frac{A(x_0) \cdot A(x-t)}{|A(x_0)|^2},$$

$$E_{\rho A^2}(x_0, t) = \frac{A(x_0) \cdot A(x-t)}{|A(x_0)|^2}.$$  

(8)

(9)

(10)

Substitution of Eq. (7) into the definition (3) yields a Monte Carlo prescription for the normalized correlation function:

$$C_W(t) = \left( \frac{E_W(x_0, t)}{E_W(x_0, 0)} \right)_W.$$  

(11)

Note that since $E_{\rho A^2}(x_0, 0) = 1$, no normalization is needed for the $\rho A^2$ algorithm. The two averages in Eq. (11) may be evaluated either with two independent Monte Carlo simulations or during a single Monte Carlo simulation. Here we consider only the latter possibility, as it is computationally faster and normalizes both $C_\rho(0)$ and $C_{\rho|A}(0)$ exactly.

Statistical errors. The three algorithms differ by the sampling weight $W$ used and consequently also by the estimator $E_W$. The computational cost of all three algorithms is $O(c \Delta t N)$, where $N$ is the number of trajectories, $\Delta t$ the time step used, and $c$ the combined cost of a single evaluation of the force (needed for the dynamics) and of the estimator $E_W$. Usually, the cost of evaluating the estimator is or can be made negligible to that of evaluating the force. Therefore the costs of the algorithms differ mainly in the number $N$ of trajectories needed to achieve a desired precision (i.e., discretization error) $\sigma_{\text{discr}}$.

Alternatively, the algorithms can be compared by evaluating the discretization errors $\sigma_{\text{discr},W}$ resulting from a given number $N$ of trajectories. For an unbiased estimator, the discretization error $\sigma_{\text{discr}}$ is equal to the statistical error $\sigma_W$, where $\sigma_W(N, t)^2 = \overline{C_W(N, t)^2} - C_W(N, t)^2$ and the overline denotes an average over an infinite number of simulations with different sets of $N$ trajectories. Assuming for now that the $N$ trajectories are uncorrelated, one can show that the error of the unnormalized $C_W(t)$ satisfies

$$\sigma_{a,W}(N, t)^2 = \frac{1}{N} \sum_{j=1}^{N} \left( E_W(x_0, t) \right)_W - \left( E_W(x_0, t) \right)_W^2.$$  

(12)

For $W = \rho A^2$, the error of normalized $C(t)$ satisfies an analogous relation obtained by removing factors of $I_W$ from Eq. (12). Statistical errors of algorithms with weights $\rho$ and $\rho|A|$, which must be normalized according to Eq. (11), are found from the formula for the statistical error of a ratio of random variables:

$$\left( \frac{S/T}{S/T} \right)^2 = \left( \frac{S}{S} \right)^2 + \left( \frac{S}{T} \right)^2 - 2 \frac{ST - ST}{ST}.$$  

(13)

In our case, $S = C_{a,W}(N, t)$ and $T = C_{a,W}(N, 0)$. Realizing that $C_{a,W}(N, t) = C_W(t)$ we obtain the following general expression for the statistical errors of the three algorithms:

$$\sigma_W(N, t)^2 = \frac{1}{N \delta W} \left( a_W C(t)^2 - 2 b_W C(t) + c_W \right),$$  

(14)

where $a_W = \langle |A|^2 \rangle_{\rho/W}^4$, $b_W = \langle |A|^2 \rangle_{\rho/W}^2 (A \cdot A)^2$, $c_W = \langle |A|^2 \rangle_{\rho/W}^2 (A \cdot A)^2 (A \cdot A)^2$, and an abbreviated notation $A^2 := A(A^2)$ was used. The special cases are obtained by replacing $W$ with $\rho$, $\rho|A|$, or $\rho A^2$ in these expressions.

For $W = \rho A^2$, the coefficients can be rearranged as $a_{\rho A^2} = -d_{\rho A^2}$, $b_{\rho A^2} = 0$, $c_{\rho A^2} = \langle |A|^2 \rangle_{\rho/W}^2 (A \cdot A)^2$, and $d_{\rho A^2} = \langle |A|^2 \rangle_{\rho/W}^2$ using the Cauchy-Schwarz inequality $\langle A^0 \cdot A^0 \rangle^2 \leq |A|^2 |A|^2$ in the expression for $c_{\rho A^2}$ and the fact that for stationary distributions $\langle |A|^2 \rangle_W = \langle |A|^2 \rangle_W$. We find that $c_{\rho A^2} = \langle |A^2|^2 \rangle_{\rho/W} = d_{\rho A^2}$ and realize that for the weight $\rho A^2$ the upper bound for the statistical error depends only on $N$ and the value of the autocorrelation function $C(t)$:

$$\sigma_{\rho A^2}^2(N, t) \leq \frac{1}{N} [1 - C(t)^2].$$  

(15)

In particular, the error does not explicitly depend on the dimensionality $D$ of the system, chaoticity of its dynamics, the nature of the observable $A$, or time $t$. This remarkable fact is the main thesis of this paper.

Special cases. One cannot make a similar general statement about either of the algorithms using weight $\rho$ or $\rho|A|$. We therefore turn to two special cases permitting analytical evaluation of the statistical errors. Both examples involve a many-dimensional harmonic oscillator (HO) $H = (1/2)(p^2/m + q^2)$ and its stationary Gaussian distribution $\rho(x) = \left(2 \tanh(u/2)\right)^D \exp[- \tanh(u/2)(q^2/2 + \alpha^2q^2/2)]$, given by the Wigner transform of the Boltzmann density operator. Above, $u := \beta \hbar \omega$, $\omega^2 = k/m$, $\alpha^2 = \hbar/(\beta m \omega)$. [Note
that the ground state density and the classical Boltzmann distribution can be obtained as the limits of Eqs. (16) for \( \beta \to \infty \) and \( \beta \to 0 \), respectively.) The two examples differ in the choice of the observable \( \mathbf{A} \).

**Exponential growth of \( \sigma \) with \( D \).** First consider \( \mathbf{A} \) to be the product of coordinates: \( \mathbf{A} = q_1 q_2 \cdots q_D \). The statistical error for \( W = \rho A^2 \) is described by Eq. (15) in full generality and thus is independent of \( D \). On the other hand, straightforward but somewhat tedious calculations using Eq. (14) show that statistical errors for both weights \( \rho \) and \( \rho |A| \) grow exponentially with the number of dimensions \( D \):

\[
\sigma_{\rho} (N, t)^2 = \frac{1}{N} \left\{ \left[ 1 + \sqrt{C(t)^2} \right]^D - 3^D C(t)^2 \right\},
\]

(17)

\[
\sigma_{\rho |A|} (N, t)^2 = \frac{1}{N} \left( \frac{2}{\pi} \right)^D \left\{ \left[ 1 + \sqrt{C(t)^2} \right]^D - 2^D C(t)^2 \right\}.
\]

(18)

The fact that for \( W = \rho \) and \( \rho |A| \) there exist observables for which the error grows exponentially with \( D \) is our second main result. Similar behavior of \( \sigma \) is expected for any multiplicatively separable function \( \rho \) of phase-space coordinates, such as the Gaussian \( \rho = \exp(-q^2/\alpha^2) \).

**Independence of \( D \).** Yet, the situation is not always so bleak. Consider the correlated function \( \rho = \mu^i \cdot q \) to be a linear function of coordinates \( q (\mu^i \text{ is a } D \text{-dimensional vector}) \). In this important special case, all three sampling methods have statistical errors independent of dimensionality:

\[
\sigma_{\rho \rho A^2} (N, t)^2 = \frac{1}{N} [1 - C(t)^2],
\]

(19)

\[
\sigma_{\rho |A|} (N, t)^2 = \frac{2}{\pi N} [1 - C(t)^2].
\]

(20)

The proof of Eq. (20) for weight \( \rho |A| \) is somewhat involved and was done only for the case \( \mu_1 = \cdots = \mu_D \). On the other hand, Eq. (19) remains valid even for HOs with different frequencies in different dimensions. Note that the statistical error is slightly lower for \( W = \rho |A| \) than for \( W = \rho \) or \( \rho A^2 \).

**Sampling methods and correlation length.** Before presenting numerical examples, let us briefly discuss the sampling methods. In many dimensions, sampling from a general weight \( W \) is often performed with the Metropolis method [17,19]. Two variants are used here: The “original” Metropolis method proposes the new point \( x_{\text{new}} \) using a random walk step from the last accepted point \( x_{\text{old}} \): \( x_{\text{new}} \) is accepted with probability \( p_{\text{acc}} = \min[|W(x_{\text{new}})/W(x_{\text{old}})|, 1] \). If \( x_{\text{new}} \) is rejected, the last accepted point \( x_{\text{old}} \) is duplicated. In the “product” Metropolis method, \( W \) is factorized as \( W = Y Z \), where \( Y \) can be sampled “directly” to propose a new point \( x_{\text{new}} \) which is subsequently accepted with probability \( p_{\text{acc}} = \min[Z(x_{\text{new}})/Z(x_{\text{old}})], 1] \).

Unfortunately, except for a few distributions \( W \) (such as the uniform or normal distributions, which may be sampled “directly”), points generated by Metropolis methods are correlated, leading to a correlation length \( N_{\text{corr}} > 1 \) between samples. This increases the statistical error for a given number of samples \( N \). As a consequence, in all of our analytical expressions, \( N \) should be replaced by \( N/N_{\text{corr}} \), which can affect (slightly) the dependence of \( \sigma \) on \( D \). An important factor increasing \( N_{\text{corr}} \) is the rejection of proposed moves, which results in exactly identical samples. In a properly designed code, however, these repeated samples do not increase the computational cost; they are accounted for by increasing the statistical weight of the original (not yet duplicated) sample. Thus, strictly speaking, the efficiency of a sampling algorithm depends on the number \( N_{\text{uniq}} \) of unique trajectories needed for convergence rather than on the total number \( N \) of trajectories. While we took \( N_{\text{corr}} \) into account in the numerical calculations, a detailed analysis of \( N_{\text{corr}} \), which can both increase (slowly) or decrease (slowly) with \( D \), is beyond the scope of this paper.

**Numerical results.** We first confirmed our analytical results for HOs numerically using \( k = m = \hbar = \beta = 1 \). Numerical statistical errors were estimated by averaging these errors over 100 independent simulations, each with the same number of unique trajectories \( N_{\text{uniq}} = 5 \times 10^5 \). In order to compare with the analytical results, the effect of correlation was removed by converting the numerical statistical error \( \sigma \) to an error per trajectory \( \sigma_1 := (N/N_{\text{corr}})^{1/2} \sigma \). The correlation lengths \( N_{\text{corr}} \) were estimated using the method of block averages [20].

Figure 1 shows that for \( A = q_1 q_2 \cdots q_D \), the error \( \sigma_1 \) grows exponentially with \( D \) for both weights \( \rho \) and \( \rho |A| \) while it is independent of \( D \) for \( W = \rho A^2 \). Moreover, numerical results agree with the analytical predictions [15,17,18]. The original Metropolis method was used since the acceptance rate of the product Metropolis method was prohibitively low for high \( D \). The step size of the random walk was the same for all three weights but varied weakly with \( D \) for the sake of a reasonable acceptance rate. [Note that in our calculations \( \sigma_{\rho A^2} = (N/\rho A^2)^{1/2} \sigma_{1, \rho A^2} \) itself grew slightly with \( D \) due to a slow growth of the correlation length \( N_{\text{corr}} \) with \( D \). For \( W = \rho \), \( N_{\text{corr}} \) decreased slightly with \( D \) and for \( W = |A| \) it stayed approximately constant, but these effects did not cancel the overall exponential growth of the error. Even though \( N_{\text{corr}} \) can be varied to some extent by modifying the step size of the random walk, this was not explored in detail here.]

Figure 2 compares the analytical predictions with numerically computed errors for \( A = \mu^i \cdot q \), where \( \mu^i \) is a \( D \)-dimensional vector with all entries equal to 1. Such \( A \) can be interpreted as a linear approximation to the electric dipole of a nonpolar molecule. Figure 2 confirms that the statistical error \( \sigma_1 \) is independent of \( D \) for all three algorithms. Initial conditions were sampled using the product Metropolis algorithm with \( W = Y Z \) and \( Y = \rho \) in all cases. Function \( Z \) used in the acceptance criterion was equal to 1, \( |A| \), and \( A^2 \), for \( W = \rho, |A|, \) and \( \rho A^2 \), respectively. Therefore, for \( W = \rho \), \( N_{\text{corr}} = 1 \) and \( N = N_{\text{uniq}} \), while for \( W = |A| \) and \( \rho A^2 \), \( N_{\text{corr}} > 1 \) and \( N > N_{\text{uniq}} \).

Finally, we used the three algorithms to calculate the
performed the same way as in the previous example. The statistical error is independent of dimensionality for the algorithm with weight $W = \rho A^2$ and grows exponentially with $D$ for the other two weights. Time $t$ was chosen separately for each $D$ so that $C(t) \approx 0.5$.

vibrational spectrum of a 48-dimensional harmonic model of the ground electronic state of azulene computed at the CASSCF(4,6)/6-31G* level of theory. Observable $\mathbf{A}$ was a linear approximation of the dipole moment of azulene, $\mathbf{A} = \mu = \mu_0 + \mu' \cdot q$, where $\mu_0 := \mu(0)$ is the equilibrium dipole moment (a 3-dimensional vector) and $\mu'$ the $3 \times D$ matrix of derivatives of the dipole moment at $q = 0$. Sampling was performed the same way as in the previous example. The dipole autocorrelation function $C(t)$ was computed intentionally only up to time $t_{\text{tot}} = 1.45\text{ps}$, which is the minimum time needed to resolve all vibrational peaks, and with only $N_{\text{uniq}} = 10^4$ unique trajectories, for which $C(t)$ starts to converge. Prior to computing the spectrum via a Fourier transform, $C(t)$ was damped by a multiplication with the function $\cos(\pi t/2 t_{\text{tot}})^2$. After the transform, $\mathcal{F}[C(t)](\omega)$ was multiplied by the factor $2 \omega \tanh(\frac{\beta \omega}{2})$, which includes the standard “quantum correction” [2] for the lack of detailed balance in the classical $C(t)$. While this correction is not exact even for HOs if $\rho$ is the classical Boltzmann density, it becomes exact for harmonic systems if $\rho$ is the Wigner Boltzmann density [10]. Figure 3, showing the high-frequency region of the spectrum containing the C-H bond stretches, confirms that all three algorithms converge to the same result (agreeing, within the resolution, with the exact spectrum). Moreover, even in this slightly more general case than the one considered in Fig. 2, the statistical errors associated with all three sampling weights stayed approximately independent of $D$. (Systems with $D < 48$ were generated by progressively cutting off the lowest frequency normal modes of azulene.)

**Conclusions.** We have demonstrated the existence of a sampling weight for which the number of trajectories needed for convergence of the normalized time autocorrelation function of any phase-space function $A$ is independent of the dimensionality and the underlying dynamics of the system. This sampling weight is $W = \rho A^2$, which may not be surprising at time $t = 0$, when this $W$ represents the ideal importance sampling weight with all trajectories contributing unity to the sum (6). Here we have shown that this sampling weight retains its favorable properties also for $t > 0$ by proving that $\sigma_{\rho A^2}$ depends explicitly only on $C(t)$ itself, and not on other parameters of the system.

While best suited for normalized autocorrelation functions, weight $\rho A^2$ can also accelerate calculations of unnormalized autocorrelation functions $C_{\mu}(t)$ via the relation $C_{\mu}(t) = C_{\mu}(0)C(t)$. In the latter case, weight $\rho A^2$ is retained for the dynamical calculation of $C(t)$, which is usually the most time-consuming task by far. The initial norm $C_{\mu}(0)$ must be com-

![Figure 1: Expected statistical error per trajectory of the autocorrelation function $C(t)$ of the function $A = q_1 q_2 \cdots q_D$ in a many-dimensional harmonic oscillator. The statistical error is independent of dimensionality for the algorithm with weight $W = \rho A^2$ and grows exponentially with $D$ for the other two weights. Time $t$ was chosen separately for each $D$ so that $C(t) \approx 0.5$.](image1)

![Figure 2: Expected statistical error per trajectory of the autocorrelation function $C(t)$ of the linear operator $A = \mu' \cdot q$ in a many-dimensional harmonic oscillator. The statistical error is independent of dimensionality for all three sampling weights studied. Time $t$ was chosen separately for each $D$ so that $C(t) \approx 0.5$.](image2)

![Figure 3: The high frequency part of the vibrational spectrum of azulene computed via the Fourier transform of the dipole time autocorrelation function.](image3)
puted separately using a normalized sampling weight such as $\rho$. Yet, one can afford many more trajectories for computing $C_u(0)$ since this calculation does not require any dynamics.

To conclude, we hope that the dimensionality-independent sampling weight will find its use in other classical, semiclassical [4, 5], and even quantum mechanical trajectory-based applications, such as those using the centroid [6–11] or ring-polymer [7–11] molecular dynamics.

Acknowledgements. This research was supported by the Swiss NSF with grants No. 200021_124936 and NCCR MUST, and by EPFL. We thank C. Mollica and T. Prosen for discussions, and V. Sharma, D. Marcos Gonzalez, M. Wehrle, and M. Šulc for assistance with numerical calculations.

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