Variational Principle for Mass Transport

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A variation principle for mass transport in solids is derived that recasts transport coefficients as minima of local thermodynamic average quantities. The result is independent of diffusion mechanisms and applies to amorphous and crystalline systems. This unifies different computational approaches for diffusion and provides a framework for the creation of new approximation methods with error estimation. It gives a different physical interpretation of the Green function. Finally, the variational principle quantifies the accuracy of competing approaches for a nontrivial diffusion problem.

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Mass transport in solids is the fundamental kinetic process controlling both the evolution of materials towards equilibrium and a variety of material properties [1]. Diffusion of atoms dictates everything from the stability of amorphous materials at finite temperature, design of nanoscaled semiconductor devices, processing of structural metals, including steels and superalloys, and performance of batteries and fuel cells to the degradation of materials due to corrosion or even irradiation. Since Einstein [2], diffusion has been understood as mesoscale motion arising from many individual atomic displacements, with significant effort over the last century to experimentally measure and model theoretically [3,4]. In the last 40 years, computation has played an increasingly important role, with different competing approximation methods developing, combined with increasingly accurate methods to compute transition state energies for atomic processes in transport [5–7]. However, while we have increasing accuracy in predicting atomic scale mechanisms, we lack a clear methodology to compare accuracy of theoretical models that derive mesoscale transport coefficients.

The modern macroscale description of mass transport comes from Onsager’s work on nonequilibrium thermodynamics [8], where atomic fluxes $J$ are linearly proportional to small driving forces. A general driving force is the gradient of chemical potential of species $\alpha$. Then, the Onsager transport coefficients are second-rank tensors $L^{(\alpha \beta)}$ that relate steady-state fluxes in species $\alpha$, where

$$J^\alpha = -\sum_\beta L^{(\alpha \beta)} \nabla \mu^\beta$$

are steady-state fluxes in response to perturbatively small driving forces in chemical species $\nabla \mu^\beta$. These transport coefficients can also be derived from a thermodynamic extremal principle [9,10] for maximum entropy production, making the Onsager matrix symmetric and positive semidefinite.

A brief, albeit incomplete, list of methods to compute transport coefficients from atomic mechanisms includes stochastic methods like kinetic Monte Carlo calculations [11–15], master-equation methods like the self-consistent mean-field method [16,17] and kinetic mean-field approximations [18–20], path probability methods for irreversible thermodynamics [21–23], Green function methods [24–27], and Ritz variational methods [28–30]. The different approaches all have different computational and theoretical complexity and rely on different approximations, which may or may not be controlled. However, the relationships between different approximations is not always clear, and it is difficult to determine which of two different calculations is more accurate, short of comparison to experimental results. In what follows, we derive a general expression for the mass transport coefficients in a solid system and then cast this nonlocal form into an equivalent minimization problem over thermodynamic averages of local quantities: a variational principle for mass transport, with a simple physical interpretation. We show that different computational approaches can be derived and compared with this principle, while also providing a framework for the development of new types of approximations for diffusion. We conclude with a quantitative comparison for a random alloy on a square lattice.

Consider a system with chemical species [31] $\alpha = A, B, \ldots$, with discrete microstates $\{x\}$ and transitions between states. For each state $\chi$ and species $\alpha$, $N^\alpha_\chi$ of that species are at positions $\{x^\alpha_i: i = 1, \ldots, N^\alpha_\chi\}$. Note that the $x^\alpha_i$ are themselves functions of the state $\chi$. If each state has an energy $E_\chi$, then in the grand canonical ensemble, the equilibrium probability of occupying a given microstate for chemical potentials $\mu^\alpha$ at temperature $T$ is

$$P^\chi_{\mu} := P^\chi_0(T, \mu^A, \ldots) = \exp \left[ \frac{1}{k_B T} \left( \Phi_0 + \sum_\alpha \mu^\alpha N^\alpha_\chi - E_\chi \right) \right] .$$

(2)
where $\Phi_0$ is a normalization constant—the grand potential—such that $\sum_{\chi} P_{\chi 0} = 1$. If the chemical potentials were spatially inhomogeneous, then the term corresponding to the sum over chemistry would be $\sum\alpha \sum_i \mu^\alpha(x_{\chi i})$. We assume that our system can achieve equilibrium through a Markovian process, with transition rates $W(\chi \to \chi') \geq 0$; then, by detailed balance, $P_{\chi 0} W(\chi \to \chi') = P_{\chi'} W(\chi' \to \chi)$.

If all nonzero rates conserve chemical species, then the rates $W(\chi \to \chi')$ are independent of the chemical potentials and can only depend on the initial and final states and temperature. The master equation for the evolution of a time-dependent probability $P_{\chi}(t)$ is

$$\frac{dP_{\chi}(t)}{dt} = \sum_{\chi'} P_{\chi'}(t) W_{\chi' \chi}.$$  (3)

where we introduce the shorthand matrix form

$$W_{\chi' \chi} = \begin{cases} W(\chi' \to \chi) : \chi \neq \chi' \\ -\sum_{\chi'} W(\chi \to \chi') : \chi = \chi'. \end{cases}$$  (4)

We identify steady-state solutions of Eq. (3)—which may not be equilibrium solutions—as distributions where the right-hand side is zero for every $\chi$; we are interested in steady-state solutions that maintain infinitesimal gradients in chemical potentials, for which we will compute fluxes.

What follows is a generalization of results derived previously for a lattice gas model [27]; details are available in the Supplemental Material [32]. Consider a steady-state probability distribution $P_{\chi ss} = P_{\chi 0}(T, \mu^\alpha, \ldots, \nabla \mu^\alpha, \ldots)$ in the presence of infinitesimally small chemical potential gradient vectors $\nabla \mu^\alpha$. This steady-state probability distribution can have time-independent fluxes $J^\alpha$ corresponding to mass transport. For any (nonzero rate) transition $\chi \to \chi'$, we define the mass transport vector for each species $\alpha$ as $\delta x_{\chi \chi'}^\alpha := \sum_i x_{\chi i}^\alpha - x_{\chi' i}^\alpha$. This is the net change in positions for all atoms of species $\alpha$, as $N_{\chi}^\alpha = N_{\chi'}^\alpha$ when $W(\chi \to \chi') \neq 0$. Then, the flux is

$$J^\alpha = V_0^{-1} \sum_{\chi' \chi} P_{\chi ss} W_{\chi' \chi} \delta x_{\chi \chi'}^\alpha.$$  (5)

for total system volume $V_0$. We make the ansatz that the steady-state probability distribution for infinitesimal gradients

$$P_{\chi ss} = P_{\chi 0} \left[ 1 + \frac{\delta \Phi_0}{k_B T} + \frac{1}{k_B T} \sum_{\alpha} \nabla \mu^\alpha \cdot \left( \eta_{\chi}^\alpha + \sum_{i=1}^{N_{\chi}} x_{\chi i}^\alpha \right) \right]$$  (6)

up to first order in $\nabla \mu^\alpha$, where $\delta \Phi_0$ is a change in the normalization relative to the equilibrium distribution, and introducing the “relaxation vectors” $\eta_{\chi}^\alpha$ that are to be determined for each state $\chi$. These vectors are a generalization of the rate-dependent relaxation in solute-vacancy exchange [33]. If we substitute Eq. (6) into Eq. (3), set $dP_{\chi ss}/dt = 0$, apply detailed balance, divide out by $P_{\chi 0}^\alpha$, and require that it hold for arbitrary $\nabla \mu^\alpha$, we find

$$\sum_{\chi'} W(\chi \to \chi') \delta x_{\chi \chi'}^\alpha = -\sum_{\chi'} W(\chi \to \chi') (\eta_{\chi}^\alpha - \eta_{\chi'}^\alpha).$$  (7)

We define the left-hand side as the velocity vector $v_{\chi}^\alpha := \sum_{\chi'} W_{\chi' \chi} \delta x_{\chi' \chi}^\alpha$, so that Eq. (7) becomes

$$v_{\chi}^\alpha = -\sum_{\chi'} W_{\chi' \chi} \eta_{\chi'}^\alpha.$$  (8)

for the steady-state ansatz solution to be time invariant. Then, the transport coefficients $L_{\chi \chi'}^{\alpha \beta}$ can be found by substituting the steady-state solution into Eq. (5), while explicitly symmetrizing the summation (rewriting as $\frac{1}{2} \sum_{\chi \chi'} + \sum_{\chi' \chi}$), which gives

$$L_{\chi \chi'}^{\alpha \beta} = \frac{1}{k_B T V_0} \left( \frac{1}{2} \sum_{\chi'} W_{\chi' \chi} \delta x_{\chi' \chi}^\alpha \otimes \delta x_{\chi' \chi}^\beta - \eta_{\chi}^\alpha \otimes \eta_{\chi}^\beta \right).$$  (9)

where the two terms are the “uncorrelated” and “correlated” contributions to diffusivity [33,34], and the average is the shorthand for $\sum_{\chi} P_{\chi ss}^{\alpha \beta}$.

While Eq. (9) has the form of a simple thermal average, the primary complication is the solution of Eq. (8), which requires the pseudoinversion of the singular rate matrix $W_{\chi' \chi}$ over the entire state space; this is the Green function $G_{\chi \chi'} := W_{\chi \chi'}^\dagger$. While the rate matrix is local—as there are only a finite number of final states $\chi'$ to transition from any state $\chi$—the Green function is known to be nonlocal and difficult to compute in general. However, the governing equation for the relaxation vectors $\eta_{\chi}^\alpha$ can be recast instead in a variational form by taking advantage of an invariance in Eq. (9).

First, the separation of Eq. (9) into correlated and uncorrelated terms is arbitrary [34,35]. We introduce changes to the positions of atoms in a state while leaving the rate matrix unchanged: let $y_{\chi}^\alpha$ be the sum of all displacements of atoms of species $\alpha$ in state $\chi$. We can, without loss of generality [36], consider only cases where $\sum_{\chi} y_{\chi}^\alpha = 0$. Then, the $y_{\chi}^\alpha$ change the displacement, velocity, and relaxation vectors

$$\tilde{\delta x}_{\chi \chi'}^\alpha = \delta x_{\chi \chi'}^\alpha + y_{\chi}^\alpha - y_{\chi'}^\alpha, \quad \tilde{v}_{\chi}^\alpha = v_{\chi}^\alpha + \sum_{\chi'} W_{\chi' \chi} y_{\chi'}^\alpha, \quad \tilde{\eta}_{\chi}^\alpha = -\sum_{\chi'} G_{\chi \chi'} \tilde{v}_{\chi}^\alpha = \eta_{\chi}^\alpha - \sum_{\chi'} G_{\chi \chi'} W_{\chi' \chi} y_{\chi'}^\alpha = \eta_{\chi}^\alpha - y_{\chi}^\alpha.$$
as $G$ is the pseudoinverse of $W$, and $y^a$ is orthogonal to the right null space of $W$. Then, the Onsager coefficients are

$$k_B T V_0 L^{(a)} = \left( \frac{1}{2} \sum_{x} W_{xx} (\delta x^a_{xx} + y^a_{xx} - y^0_{xx}) \otimes (\delta x^0_{xx} + y^0_{xx} - y^0_{xx}) - \left( b^a_x + \sum_{x} W_{xx} y^0_x \right) \otimes (\eta^a_x - y^0_x) \right)_x$$

$$W_{xx} = \sum_{yy} W_{xx} (y^a_{xx} - y^0_{xx}) \otimes (y^0_{xx} - y^0_{xx}) + \frac{1}{2} \sum_{x} P^0_x W_{xx} \delta x^a_{xx} \otimes (y^0_x - y^0_x) + \frac{1}{2} \sum_{x} P^0_x W_{xx} \delta x^0_{xx} \otimes (y^0_x - y^0_x)$$

$$+ \frac{1}{2} \sum_{x} P^0_x W_{xx} (y^a_{xx} - y^0_{xx}) \otimes (\delta x^0_{xx} - (b^a_x \otimes \eta^a_x)_x) + \sum_{x} P^0_x b^a_x \otimes y^0_x - \sum_{x} P^0_x W_{xx} y^0_x \otimes \eta^a_x + \sum_{x} P^0_x W_{xx} \delta x^a_{xx} \otimes y^0_x$$

$$= k_B T V_0 L^{(a)} - \sum_{xx} P^0_x W_{xx} (y^a_{xx} \otimes y^0_x - \eta^a_x \otimes y^0_x)_x - (y^0_x \otimes b^a_x)_x + \left( b^a_x \otimes y^0_x \right)_x + (y^a_x \otimes b^a_x)_x$$

$$+ \sum_{xx} P^0_x W_{xx} \delta x^a_{xx} \otimes \delta x^0_{xx} = k_B T V_0 L^{(a)}.$$

This requires detailed balance $P^0_x W_{xx} = P^0_x W_{xx}$ and the sum rule $\sum_{x} W_{xx} = 0$. Hence, the transport coefficients are invariant under arbitrary displacements, while the uncorrelated and correlated terms themselves change.

We can exploit this invariance by noting that, for $\alpha = \beta$, the uncorrelated contribution is positive semidefinite and the correlated contribution is negative semidefinite, as $W_{xx}$ and $\delta x_{xx}$ are negative semidefinite matrices. Thus, the maximum value of the correlated contribution is zero, which corresponds with the minimal value of the uncorrelated contribution, and so the equation for the transport coefficients can be rewritten as

$$L^{(a)} = \frac{1}{2 k_B T V_0} \inf_{\gamma^a} \left( \sum_{x} W_{xx} \tilde{\delta x}^a_{xx} \otimes \tilde{\delta x}^a_{xx} \right)_x,$$  \hspace{1cm} (10)

which is a variational principle for mass transport, involving only thermodynamic averages of local rate matrix. Here, the infimum of the tensor corresponds to the tensor with the smallest trace [37]. The values of $y^a_x$ that minimize Eq. (10) are found by making the generalized force from the gradient of $\epsilon^a := k_B T V_0 L^{(a)} = \frac{1}{2} \sum_{xx} W_{xx} (\tilde{\delta x}^a_{xx})^2$:

$$F^a_x := \frac{\partial \epsilon^a}{\partial y^a_x} = -\frac{1}{2} \frac{\partial}{\partial y^a_x} \sum_{xx} P^0_x W_{xx} (\delta x^a_{xx} + y^a_{xx} - y^0_{xx})^2$$

$$= -2 P^0_x \sum_{x} W_{xx} (\delta x^a_{xx} + y^a_{xx} - y^0_{xx}) = -2 P^0_x b^a_x$$  \hspace{1cm} (11)

equal to zero; this is satisfied when $y^a_x = \eta^a_x$. Moreover, the arguments $y^a_x$ that minimize $\epsilon^a$ can then be used to compute the off-diagonal contributions.

Hence, while the $\alpha \neq \beta$ terms are not variational, they are determined by the solution of the diagonal variational problem. Equation (10) is similar to the Varadhan-Spohn variational form [38], which Arita et al. note is a powerful, albeit abstract, result that is difficult to apply in practice, involving “cylinder” functions [39]; it is simpler than the alternate Ritz variational form, as there is no normalization of an eigenvector required [28–30].

This variational principle for mass transport has multiple consequences. First, it unifies multiple approaches for the computation of mass transport coefficients, including kinetic Monte Carlo calculations, Green function methods, and self-consistent mean-field theory. Moreover, it provides a direct way to compare the accuracy of different methods: outside of the convergence of stochastic sampling errors, once a mass transport method is recast in a variational form, the minimal value of the diagonal transport coefficients is necessarily closer to the true value. It also gives a simple physical explanation for the correlation contributions in mass transport: the $\eta^a_x$ values are displacements that map a correlated random walk into an equivalent uncorrelated random walk with identical transport coefficients. Finally, it provides a framework for the construction of new algorithms for the computation of mass transport that requires the minimizing of a thermal average; as it is based on minimization, different approximations for $y^a_x$ can be simultaneously introduced, while the process of minimization finds the optimal solution.

In the case of a linear expansion for the relaxation vectors, the variational principle for mass transport provides a simple general expression for diffusivity. Let $\{ \phi^a_{x,a} \}$ be a set of basis vectors so that we expand $y^a_x = \sum_a \phi^a_{x,a} \theta^a_x$ with coefficients $\theta^a_x$. Section S4 of the Supplemental Material [32] shows the most general solution of the...
linearized basis approximation method (LBAM); here, we include the solution for the case where the basis functions are chemistry and direction independent: \( \phi^{a}_{x,mi} = \hat{e}_{i} \phi_{x,n} \) for a Cartesian orthonormal basis \( \hat{e}_{1}, \hat{e}_{2}, \hat{e}_{3} \). Then, the coefficients that minimize Eq. (10) can be found by solving \( \sum_{nm} W_{nm} \theta_{mi} = \mathbf{b}_{n}^{a} \cdot \hat{e}_{i} \), where
\[
W_{nm} := \left\langle \sum_{x} W_{XX'} \phi^{a}_{x,n} \phi^{a}_{x',m} \right\rangle_{x}, \quad \mathbf{b}_{n}^{a} := \left\langle \phi^{a}_{x,n} \mathbf{b}^{a}_{x} \right\rangle_{x}. \tag{13}
\]

We can take the pseudoinverse of \( \tilde{G} := (\tilde{W})^{+} \), and then the transport coefficients are [cf. Eq. (S26)]
\[
L_{\text{LBAM}}^{(a\theta)} = \frac{1}{2k_{B}T_{0}} \left( \sum_{x} W_{XX'} \delta x^{a}_{x} \otimes \delta x^{\theta}_{x'} \right)_{x} + \frac{1}{k_{B}T_{0}} \sum_{nm} \left( \phi_{x,n} \mathbf{b}^{a}_{x} \right)_{x} \otimes \tilde{C}_{nm} \left( \phi_{x,m} \mathbf{b}^{\theta}_{x} \right)_{x}, \tag{14}
\]
where the diagonal transport coefficients \( L_{\text{LBAM}}^{(a\alpha)} \) are guaranteed to be an upper bound on the true coefficients, achieving equality when the basis \( \{ \phi_{x,n} \} \) spans \( \eta_{x}^{a} \).

We can now express existing computational approaches as attempts to solve the variational problem. For kinetic Monte Carlo calculations [11–15], each trajectory represents a single sample in the average, while the increasing length of a trajectory attempts to converge the relaxation vectors corresponding to that convergence starting state. In Sec. S3 of the Supplemental Material [32], the equivalence of kinetic Monte Carlo (KMC) calculations to the variation method is shown; moreover, the use of a finite length trajectory is variational. Assuming perfect sampling of initial states and with perfect sampling of trajectories of a finite length, the transport coefficients will be greater than the true transport coefficients. If one uses accelerated KMC methods [40–44], superbasins—a finite collection of states with fast internal transitions but slow escapes—are effectively collapsed onto a single position, which is an approximation to the relaxation vector \( \eta_{x}^{a} \). For vacancy-mediated diffusion, the dilute Green function method [26,27] and matrix methodology [24,25] work in a restricted state space \( \{ \chi \} \), where only one solute and vacancy are present, and then effectively construct a full basis in that state space. Finally, self-consistent [16,17] and kinetic mean-field [18–20] methods work with a cluster expansion of chemistry- and direction-independent basis functions \( \{ \phi_{x,n} \} \), which are products of site occupancies for different chemistries. It should be noted that these latter two methods derive their solution for the parameters \( \theta_{n} \) using a ladder of \( n \)-body correlation functions on which they invoke “closure approximations” for higher order correlation functions; in a variational framework, such closure approximations become unnecessary. Finally, when methods are framed in variational terms, we can quantitatively compare accuracy by identifying which method gives the smallest diagonal elements \( L^{(a\alpha)} \) and also estimate remaining error through the average residual bias \( \left\langle (\mathbf{b}^{a}_{x})^{2}/(-W_{XX'}) \right\rangle \) in Eq. (S29) or its ratio with \( \left\langle (\mathbf{b}^{a}_{x})^{2}/(-W_{XX'}) \right\rangle_{x} \).

In addition to providing a common frame for existing computational methods for mass transport, we now have a new framework to develop and test new approximations, including those that are more appropriate for amorphous systems that lack crystalline order but still possess well-defined microstates. A simple example is the basis function choice \( \phi^{a}_{x} = \mathbf{b}^{a}_{x} \); in Sec. S5 of the Supplemental Material [32], a closed-form approximation for transport coefficients is provided in Eq. (S32). This approximation involves inverting a matrix that has the same dimensionality as the number of independent chemical species; however, it only captures local correlations. We can also take the dilute Green function methodology for vacancy-mediated transport into finite solute concentrations by using the basis functions \( \phi_{x,\beta} \) that are equal to the occupancy (0 or 1) by chemistry \( \beta \) of a site at a vector \( x \) relative to a vacancy in state \( \chi \). This approximation exactly reproduces the dilute solute limit by being equivalent to an infinite range two-body-only version of the Green function.

For a quantitative comparison of these new approximations, we consider a random binary alloy on a square lattice with a single vacancy. In this model, there is no binding energy between any species, and the jump rate for the vacancy only depends on the chemistry of the species it is exchanging: either \( \nu_{A} \) (solvent exchange) or \( \nu_{B} \) (solute exchange). We take \( \nu_{A} = 1 \) and consider three cases: \( \nu_{B} = 1 \) (tracer), \( \nu_{B} = 4 \) (fast diffuser), and \( \nu_{B} = 0 \) (frozen solute). This system has nontrivial behavior, including a percolation limit [45,46] for \( \nu_{B} = 0 \), where the diffusivity of the solvent is zero for \( c_{B} < 1 \). To compute the transport coefficients, we use the following: (1) Kinetic Monte Carlo calculations on a \( 64 \times 64 \) periodic grid, 256 samples of trajectories are generated for 4096 vacancy jumps each; the transport coefficients are computed 32 separate times to get a mean and stochastic error estimate. (2) A two-body Green function (GF) approximation (cf. Sec. S6 [32]) has the analytic solution [cf. Eq. (S40) [32]],
\[
L_{\text{GF}}^{(AA)} = 1c_{A}a_{0}^{2} \left( c_{A}\nu_{A} - \frac{c_{A}c_{B}\nu_{A}^{2}}{\nu_{A} + \nu_{B} + 2(1-1/2)\left( c_{A}\nu_{A} + c_{B}\nu_{B} \right)} \right)
L_{\text{GF}}^{(BB)} = 1c_{B}a_{0}^{2} \left( c_{B}\nu_{B} - \frac{c_{A}c_{B}\nu_{B}^{2}}{\nu_{A} + \nu_{B} + 2(1-1/2)\left( c_{A}\nu_{A} + c_{B}\nu_{B} \right)} \right),
\tag{15}
\]
where \( f = (\pi - 1)^{-1} \approx 0.467 \) is the dilute tracer correlation coefficient for a square lattice. (3) A basis approximation has the same transport coefficients as Eq. (15) with the approximation \( f = 1-2/(\pi + 1) = 0.6 \). (4) A self-consistent mean-field approach is used with clusters of all orders within two jumps: \( \pm \hat{x}, \pm \hat{y}, \pm \hat{x} \pm \hat{y}, \pm 2\hat{x} \), and
solute concentrations from zero to one. The largest deviation is

SCMF methods are good approximations for both the Green function and self-consistent mean-field and only the self-consistent mean-field method should map all "vacancy is trapped and unable to diffuse over long limit. In this case, solutes are creating islands, where a but begin to break down as we approach the percolation
difference is seen for the percolation case \( \nu_B = 0 \), where the lowest diffusivity is the most accurate of the three approaches. The largest

\[ B_c \approx 0 \], which requires longer range correlation to capture the percolation limit near \( c_B \approx 0.5 \); the "percolation" simulations are averages of 256 Green function calculations on a 256 × 256 periodic cell.

Finally, for \( \nu_B = 0 \), we use a full Green function
solution for vacancy diffusivity with 256 configurations of a 256 × 256 periodic cell; and compute a residual bias correction (RBC) for the two-body Green function results.

Figure 1 shows the different accuracy for this binary system, where the lowest diffusivity is the most accurate [47,48]. The Green function approach captures the dilute \( A \) and \( B \) limits for the tracer and fast diffuser examples and is the most accurate of the three approaches. The largest
difference is seen for the percolation case \( \nu_B = 0 \), where both the Green function and self-consistent mean-field (SCMF) methods are good approximations for \( c_B \lesssim 0.2 \), but begin to break down as we approach the percolation limit. In this case, solutes are creating islands, where a vacancy is trapped and unable to diffuse over long distances; inside such an island, the relaxation vectors should map all “trapped” states onto the same position, producing no contribution to the diffusivity. We also see the direct simulations produce lower, more accurate, diffusivity. The size of these islands gets smaller as \( c_B \) increases, and only the self-consistent mean-field method—and only at large concentrations of solute—is able to reproduce the behavior seen by kinetic Monte Carlo calculations. This suggests the need to go beyond the two-body basis for the Green function approach, combining local multisite basis functions with long-range basis functions or perhaps new approximation methods all together. One such approach is the RBC, where following a linear basis approximation method the residual bias vectors serve as basis vectors for a correction to the diffusivity; in the case of \( \nu_B = 0 \), we derive an analytic expression [cf. Sec. S7, Eq. (S47) [32] ] that has a similar error to the SCMF result.

With a variational formulation of transport coefficients, we can develop new approximate methods for modeling diffusion in solids, including amorphous materials. If linear approximations are used, then basis functions provide a projection of the state space into a subspace, while the variational principle provides a lower bound on transport coefficients. The selection of basis functions can be guided by physical insight—such as diffusing quasiparticles—and systematic improvement is always possible. It is also possible to construct nonlinear approximations to the relaxation vectors \( \nu^2 \), which might require fewer parameters to describe; still, a variational principle permits relative comparisons of different methods and a lower bound on the result. While the fundamental insight for the variational formulation came from the invariance in Eq. (9), it can be derived as a thermodynamic extremum principle where the positions of atoms are “free” variables, connecting to Onsager’s original work.

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[31] The species can include vacancies as an independent chemical species.


[36] This is accomplished by subtracting a constant value from each $y_j^2$ to ensure that the sums vanish. This constant leaves the mass transport vectors unchanged.

[37] This follows as the correlated contribution is a symmetric negative semidefinite matrix, and the largest value possible is zero, which is achieved when the trace is similarly maximized.

[38] H. Spohn, Large Scale Dynamics of Interacting Particles (Springer-Verlag, Berlin, 1991).


[47] The python notebook for the random alloy results in Fig. 1 is available from D. R. Trinkle, “Onsager,” http://dallastrinkle.github.io/Onsager, which makes use of the algebraic multigrid solver PYAMG.