



Relaxation of terrace-width distributions: Physical information from Fokker–Planck time

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ABSTRACT

Recently some of us have constructed a Fokker–Planck formalism to describe the equilibration of the terrace-width distribution of a vicinal surface from an arbitrary initial configuration. However, the meaning of the associated relaxation time, related to the strength of the random noise in the underlying Langevin equation, was rather unclear. Here we present a set of careful kinetic Monte Carlo simulations that demonstrate convincingly that the time constant shows activated behavior with a barrier that has a physically plausible dependence on the energies of the governing microscopic model. Remarkably, the rate-limiting step for relaxation in the far-from-equilibrium regime is the generation of kink–antikink pairs, involving the breaking of three lateral bonds on a cubic {001} surface, in contrast to the processes breaking two bonds that dominate equilibrium fluctuations. After an initial regime, the Fokker–Planck time at least semiquantitatively tracks the actual physical time.

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1. Introduction

With equilibrium properties of vicinal surfaces—especially the form of the terrace width distribution (TWD)—now relatively well understood [1], much attention is focusing on non-equilibrium aspects, which have long been of interest. In a previous paper some of us [2] derived the following Fokker–Planck (FP) equation (Eq. (1)) to describe the distribution of spacings between steps on a vicinal surface during relaxation to equilibrium. The goal was to describe the relaxational evolution of this spacing distribution rather than the evolution of the positions of individual steps as in a previous investigation [3–6]. As in all those papers, we simplify to a one-dimensional (1D) model, in which a step is represented by its position in the \hat{x} direction (the downstairs direction in “Maryland” notation), averaged over the \hat{y} direction (along the mean direction of the step, the “time-like” direction in fermionic formulations) [7]. This picture implicitly assumes that one is investigating time scales longer than that of fluctuations along the step.

We started with the Dyson Coulomb gas/Brownian motion model [8,9]; made the mean-field-like assumption, when comput-

ing interactions, that all but adjacent steps are separated by the appropriate integer multiple of the mean spacing; and set the width of the confining [parabolic] potential in the model to produce a self-consistent solution. Details are provided in the Appendix, which expands the earlier derivation and corrects some inconsequential errors in intermediate stages [2]. We found the following:

$$\frac{\partial P(s, \tilde{t})}{\partial \tilde{t}} = \frac{\partial}{\partial s} \left[\left(2b_\varrho s - \frac{\varrho}{s} \right) P(s, \tilde{t}) \right] + \frac{\partial^2}{\partial s^2} [P(s, \tilde{t})], \quad (1)$$

where s is the distance w between adjacent steps divided by its average value $\langle w \rangle$, determined by the slope of the vicinal surface.

The steady-state solution of Eq. (1) has the form of the generalized Wigner surmise (GWS), thus

$$P_\varrho(s) = a_\varrho s^\varrho e^{-b_\varrho s^2} \quad a_\varrho = \frac{2 \left[\Gamma\left(\frac{\varrho+2}{2}\right) \right]^{2\varrho+1}}{\left[\Gamma\left(\frac{\varrho+1}{2}\right) \right]^{2\varrho+2}} \quad b_\varrho = \left[\frac{\Gamma\left(\frac{\varrho+2}{2}\right)}{\Gamma\left(\frac{\varrho+1}{2}\right)} \right]^2 \quad (2)$$

where the constants b_ϱ and a_ϱ assure unit mean and normalization, respectively. (The Wigner surmise, Eq. (2) pertains to the special cases $\varrho = 1, 2, 4$; the generalization is to use this expression for arbitrary $\varrho \geq 1$.) The dimensionless variable ϱ gauges the strength A of the A/w^2 energetic repulsion between steps: $(\varrho - 1)^2 = 1 + 4A\tilde{\beta}/(k_B T)^2$, where $\tilde{\beta}$ is the step stiffness. The dimensionless FP time \tilde{t} can be written as t/τ ; here the relaxation time τ is $\langle w \rangle^2/\Gamma$, where $\sqrt{\Gamma}$ is the strength of the white noise in the Langevin equation (for the step position) underlying the FP equation [2].

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To confront data, both experimental and simulational, one typically investigates the variance $\sigma^2(t)$ or standard deviation $\sigma(t)$ of this distribution. If the initial configuration of the vicinal surface is “perfect” (i.e., has uniformly-spaced straight steps), then $\sigma(t)$ obeys [2]

$$\ln \left[1 - \left(\frac{\sigma(t)}{\sigma_{\text{sat}}} \right)^2 \right] \propto -t/\tau; \quad \sigma_{\text{sat}}^2 = \frac{(\varrho + 1)}{2b_\varrho} - 1 \quad (3)$$

where $\sigma_{\text{sat}}^2 \equiv \sigma(\infty)$ is the variance for an infinite system at long time. When dealing with numerical data, we take the variance to be normalized by the mean spacing, so divided by the squared mean terrace width, to mimic the formal analysis. The precise value of the proportionality constant is not of importance to our analysis, since we view τ as the source of information for an activated process, with any prefactor therefore insignificant. As discussed in the Appendix (esp. Eq. (A15)), one might expect the prefactor to be unity when the first moment has the assumed GWS value of one, but with the approximations we make to obtain a compact solution, the prefactor seems better described as two.

Time in this formulation is not the natural fermionic time associated with the direction along the steps (\hat{y} in “Maryland notation”), i.e., that resulting from the standard mapping between a 2D classical model and a (1 + 1)D quantum model. Instead it measures the evolution of the 2D or (1 + 1)D system toward equilibrium and the thermal fluctuations underlying dynamics. Since the time constant τ enters rather obliquely through the noise force of the Langevin equation, a key investigational objective in the previous Letter [2] and in this paper is whether τ corresponds to a physically significant rate. Monte Carlo simulations allow the examination of a well-controlled numerical experiment. In the former we used our well-tested Metropolis algorithm to study a terrace-step-kink (TSK) model of the surface. We found a satisfactory fit to the form of Eq. (3), from which we obtained $\tau \approx 714$ MCS (Monte Carlo steps per site) for $\varrho = 2$ (or $A = 0$, only entropic repulsions) while $\tau \approx 222$ MCS for $\varrho \approx 4.47$. This result is in qualitative agreement with the understanding that Γ should increase (and, so, τ should decrease) with increasing ϱ , as discussed in Ref. [2].

In this paper, we confront more systematically and thoroughly the above-noted crucial issue, showing that the time constant associated with the FP transcription can be related to the atomistic processes underlying the relaxation to equilibrium and that the FP time in some sense tracks (though of course does not replicate) the literal physical time of the relaxing system. We use a standard, simple lattice model that embodies the basic atomistic properties of these surfaces. We report far more extensive simulations, using kinetic Monte Carlo (KMC) [10,11] rather than the Metropolis algorithm, for a solid-on-solid (SOS) rather than a TSK model, so that we have real mass transport. Since atomic energies in this generic model are proportional to the number of lateral nearest neighbors, detailed-balance is satisfied. To simplify the analytic expressions and, especially, the simulations, we concentrate in this paper on the special case $\varrho = 2$, corresponding to steps with only entropic repulsions, “free fermions”. We find that the time constant, extracted from the numerical data by fitting to the time correlation function in the form predicted by the FP analysis, has an activated form that can be related to an atomistic rate-limiting process in the simulations. Our goal is not to find the best accounting for the dynamics of a real stepped surface, nor even of our model surface. It is to show that the FP approach offers a relatively simple and physically viable approach to accounting for the relaxation of artificial initial configurations toward equilibrium.

The second section describes the model and KMC algorithm that we use. The third presents our numerical results. The fourth discusses them, with one subsection describing the crucial role played

by the creation of kink–antikink pairs and another investigating the evolution of the shape of the distribution. The fifth makes comparisons with the venerable mean-field treatment of step distributions, and the final section sums up our findings. In an Appendix we expand the derivation of the key Fokker–Planck equation given in Ref. [2]; we present some new results for the evolving moments of the $P_2(s, \hat{t})$ and correct some inconsequential errors in Ref. [2].

2. Model

Our SOS model assigns an integer height $h_{\mathbf{r}}$ to each point \mathbf{r} on a square grid of dimensions $L_x \times L_y$. We use periodic boundary conditions in the \hat{y} direction. On our vicinal (001) simple cubic crystal, we create N close-packed [100] steps, with mean separation $L = L_x/N$, via screw periodic boundary conditions in the \hat{x} direction. In our simulations we take $N = 5$ in the initial simulations [7] and $N = 20$ in later investigations. The energy of a configuration is given by the standard absolute SOS prescription:

$$\mathcal{H} = \frac{1}{2} E_a \sum_{\mathbf{r}\delta} h_{\mathbf{r}} h_{\mathbf{r}+\delta} \quad (4)$$

where δ runs over the four nearest-neighbors of a site, and the factor 1/2 cancels the double-counting of bonds.

In our SOS model, which has been described elsewhere [12], we use barriers determined by the standard longstanding simple rule [13–15] of bond-counting: the barrier energy E_b is a diffusion barrier E_d plus a bond energy E_a times the number of lateral nearest neighbors in the initial state. This number is 1 for an edge atom leaving a straight segment of step edge for the terrace, 3 for a detaching atom that originally was part of this edge (leaving a notch or kink–antikink pair in the step), or 2 for a kink atom detaching, either to the step edge or the terrace. Processes that break four bonds, in particular the removal of an atom from a flat terrace plane, are forbidden, as is any form of sublimation. No Ehrlich–Schwoebel barrier hinders atoms from crossing steps. We chose values $0.9 \leq E_d \leq 1.1$ eV and $0.3 \leq E_a \leq 0.4$ eV, using temperatures $520 \text{ K} \leq T \leq 580 \text{ K}$. At these temperatures we expect no significant finite-size effects in the \hat{y} direction for the values of the mean terrace width L (in lattice spacings) that we use: $4 \leq L \leq 15$.

The width L_y of the lattice should be greater than the “collision length” y_{coll} , the distance along \hat{y} for a step to wander a distance $L/2$ in \hat{x} . For a TSK model, estimates using a random-walk model give $y_{\text{coll}} = (L^2/2) \sinh^2(E_k/2k_B T)$ [16], where E_k is the formation energy of a kink. At the temperatures and energies used in our simulation, y_{coll} is of order 10^2 for $L = 6$ and 10^3 for $L = 15$. E.g., for $T = 580 \text{ K}$, $E_k = E_d/2 = 0.175$ eV, and $L = 6$, $y_{\text{coll}} \approx 140$. In almost all simulations reported here, we use $L_y = 10^4$. While L_y may often be larger than necessary, it allows for some self-averaging, decreasing the number of runs we need to carry out to get good statistics.

In our rejection-free KMC, we separate all top-layer sites into four classes, those with $i = 0, 1, 2, 3$ nearest neighbors (NNs). (Those with $i = 4$ are not allowed to move and are not considered when updating.) Typical realizations of these four classes are isolated adatoms, atoms protruding from a straight step edge, atoms at kink sites, and atoms at the edge of a step, respectively. We compute probabilities for each of the movable classes: $P_i = f_i / \sum_{i=0}^3 f_i$, where $f_i = N_i \times \exp[-(E_d + iE_a)/k_B T]$, and N_i is the number of sites with i NNs. (Of course, the four exponentiations are done once and for all at the beginning for each set of energies.) For each update we need four random numbers— r_1, r_2, r_3, r_4 —uniformly distributed between 0 and 1. We use r_1 to pick which of the 4 movable classes will have the move. For the “winning” class, r_2 determines which of the N_i possible atoms will move. Then r_3 determines in which of the 4 NN directions the atom moves. In this rejection-free scheme, we then decrease the height (the z value) of the initial

position by one and increase the height of the chosen direction move from this initial site by one. This scheme can be (and has been, elsewhere) modified to allow for an Ehrlich–Schwoebel barrier. Finally r_4 is used to advance the clock in standard KMC fashion, similar to the n -fold way or BKL [17] approach, using the prescription $\Delta t = -\ln(r_4)/R$, where R is the total rate for a transition from the initial state [10]. Explicitly, $R = \nu_0 \sum_{i=0}^3 f_i = \nu_0 \exp[-E_d/k_B T] \sum_{i=0}^3 N_i \times \exp[-i \cdot E_a/k_B T]$, where we take the hopping frequency $\nu_0 = 10^{13} \text{ s}^{-1}$.

We saved essentially every hundredth update; that interval corresponds to our unit of time, which is about 1 s. for the selected temperature and energies.³ This update interval is long enough so that the sum of the KMC update times varies insignificantly ($\pm 0.01\%$) but short enough to capture the behavior during the steep initial rise.

In our model, the mass carriers are atoms rather than vacancies (or both). Since atoms with $i = 4$ are frozen in our model, atom–vacancy pairs cannot form spontaneously on a terrace. (More generally, this mechanism is highly improbable.) Mass carriers are thus created at step edges. If MC moves depend on the difference between final and initial energies, as in Metropolis schemes, then there is equivalence between atom and vacancy creation and transport. (If one goes beyond a strict SOS model and allows local relaxation, vacancies tend to be favored somewhat [18,19].) An atom quitting a step edge for the lower terrace costs $3E_a$ if it leaves a straight step and $2E_a$ if it leaves from a kink. At the upper side of a step, a vacancy can be spawned if a step-edge atom moves out one spacing onto the lower terrace (with the same energy cost as just given) and its inner neighbor happens to move in the same direction before the initial atom returns to its initial position. In kinetic Monte Carlo, however, rates are determined just by the difference between the barrier energy and the initial-state energy. This does not change the energy to produce an atom, but adds a cost of $3E_a$ for the move of the second, inner-neighbor atom. Moreover, while the energy for an atom to hop along the terrace is E_d , for a vacancy it is $E_d + 3E_a$. Indeed, we never observed the unlikely concerted process for vacancy creation in our simulations nor, for that matter, did we see any vacancies. The number of isolated atoms was also very small, with N_0 being in single digits, and they moved very rapidly, rarely appearing in successive saved images.

The freezing of $i = 4$ processes marks a violation of detailed balance (since such a vacancy, if it existed, could be filled by a roving adatom); however, given the negligible occurrence of such vacancies in our simulations, the violation should be insignificant. In some physical systems, motion of surface vacancies does evidently dominate mass transport [20]. Again, our goal in these calculations is not to account generally for experiments but to create a fully-controlled data set to see how well the dynamics can be described using our Fokker–Planck formalism.

3. Computed results

We extract a characteristic time (or inverse rate) τ from numerical data by fitting the dimensionless width using Eq. (3), as illustrated in Fig. 1. The fit is notably better than that found in the Metropolis/TSK study in Ref. [2]. [However, the saturation value is notably higher than in Ref. [2], with the normalized standard deviation σ (the value in the simulation divided by L) approaching ~ 0.48 , or a dimensionless variance of 0.24, rather than 0.18 as found in Ref. [2] and anticipated from Eq. (3). This difference arises because the present algorithm allows steps to make contact along

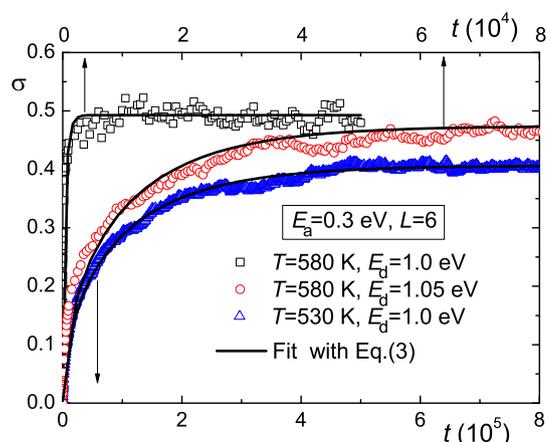


Fig. 1. Three examples of fits using Eq. (3), used to extract τ . Note that the data are very well fit in all three cases. The plotted $\sigma(t)$ is the standard deviation of the KMC data divided by the mean step spacing L (listed in lattice constants), and E_d and E_a are the energy barriers for diffusion and for breaking a bond, respectively. Time t is essentially in seconds (see text). In all cases here and in later figures, $\rho = 2$. Here $N = 5$. (For interpretation of the references to color in this figure, the reader is referred to the web version of this article.)

edge links rather than just at corners as in the usual fermion simulations. The variance of 0.18 is appropriate to “free fermions” with $\rho = 2$. As we discuss in detail elsewhere [21], the present algorithm leads to a smaller (and L -dependent) effective ρ as the steps come in contact more frequently, i.e., for smaller L and higher T (cf. Fig. 1). For the present choice of parameters ($L = 6$, $k_B T/E_d \approx 1/20$), the TWD has close to $\rho = 1$, for which the dimensionless variance is 0.27. This feature is inconsequential for the arguments in this paper.]

We expect that the decay time exhibits Arrhenius behavior: $\tau \propto \exp(E_b/k_B T)$. We investigate E_b closely in the two traces of Fig. 2. We show typical runs at $T = 580 \text{ K}$, corresponding to $k_B T \approx 1/20 \text{ eV}$. First, we ramped E_d , holding E_a fixed at 0.35 eV (open squares, red). In the semi-log plot of reduced energies (energies/ $k_B T$), we find a slope of 0.99 ± 0.02 , indicating that in the effective barrier, the multiplier of $E_d/k_B T$, is essentially unity, as expected. In a second set of runs, we ramped E_a , holding E_d fixed

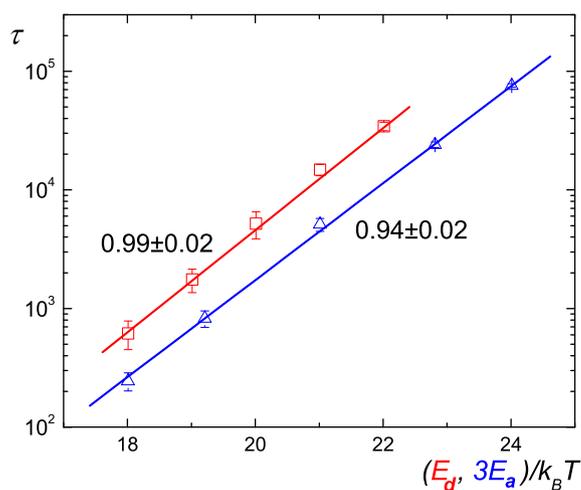


Fig. 2. Semilog plots of the relaxation time τ (in sec.) vs. the diffusion barrier E_d (squares, upper line, red) or thrice the bond energy E_a (triangles, blue), with the other held fixed, both in eV, with $k_B T = 0.05 \text{ eV}$ and $N = 5$. The numbers indicate the slopes; both are essentially unity. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

³ The mean of the Poisson-distributed update intervals [11] was about 0.01 s. We saved when the counter reached 100 times this mean. The mean overshoot was also about 0.01 s.

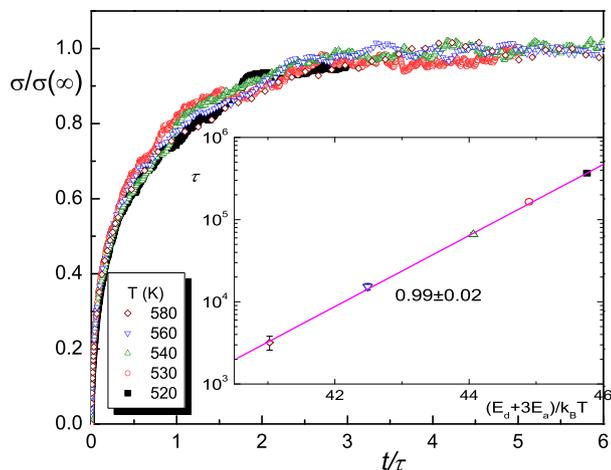


Fig. 3. Demonstration of the robustness of the form of the evolving standard deviation, $\sigma(t) = \sigma(\infty)[1 - \exp(-t/\tau)]^{1/2}$, for five temperatures. In the inset, the five values of τ by which the curves are rescaled are plotted vs. $(E_d + 3E_a)/k_B T$, showing their Arrhenius form. Here $E_d = 1.0$ eV and $E_a = 0.35$ eV. In this and subsequent figures, $N = 20$.

at 1.0 eV (open triangles, blue). Plotting now vs. $3E_a/k_B T$, we find a slope 0.94 ± 0.02 , indicating that the effective energy barrier E_b is $E_d + 3E_a$. To corroborate this idea, we ramped T from 520 to 580 K, fixing $E_a = 0.35$ eV and $E_d = 1.0$ eV. As illustrated in the inset of Fig. 3, we determine the fitted activation energy to be 2.03 ± 0.03 , in excellent agreement with $E_d + 3E_a = 2.05$ [eV]. Evidently the rate-determining process is the removal of a 3-bonded atom from a straight step, creating a pair of kinks (i.e., a kink and an antikink [22]) rather than the presumably more frequent process, with energy $E_d + 2E_a$, in which an atom leaves a kink position of a step [22,23]. (Of course, kink–antikink pairs also arise with a lower barrier when an atom from the terrace or from a kink site attaches to a step edge or splits off from a kink site. However, as members of class $i = 1$, such edge-atom structures are likely to be very short-lived.) The main part of Fig. 3 shows the standard deviation (\propto TWD width) vs. time scaled by the relaxation time of each of the five temperatures. Evidently the fit to $\sigma(t) = \sigma(\infty)[1 - \exp(-t/\tau)]^{1/2}$ is robust.

Initially the steps retreat as atoms are emitted. There is also an asymmetry in fluctuations from a straight step, since retreating moves involve higher barriers than advancing fluctuations. Once the continuum picture becomes applicable, the fluctuations appear to be symmetric, with typical configurations shown in Fig. 4.

To check consistency, we compare the intercepts of the linear fits in the two semilog plots, i.e., the prefactors of the exponential term in which the particular energy is ramped. In addition to the activation components there is the leading factor $\tau_0 \equiv \langle w \rangle^2 / 4v_0$ [2],⁴ where we make the standard assignment for the hopping frequency, $v_0 = 10^{13}$ Hz. Since $\langle w \rangle \equiv L = 6$ in our simulations, τ_0 is 9×10^{-13} s. In the ramp of E_d , the prefactor is $\tau_0 \exp(3E_d/k_B T)$, predicted to be 1.19×10^{-3} s. The value we find from the simulations is $(1.2 \pm 0.1) \times 10^{-3}$ s, in excellent agreement. Similarly in the ramp of E_a , the prefactor $\tau_0 \exp(E_d/k_B T)$ is predicted to be 4.366×10^{-4} s and measured from the fit as $(4.36 \pm 0.15) \times 10^{-4}$ s.

We also varied the system size L_x , holding the number of steps fixed, and thereby ramping $\langle w \rangle$. From the random-walk analogy,

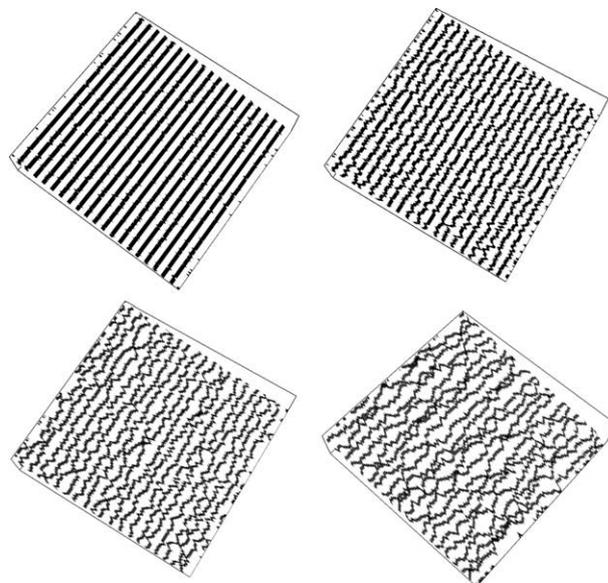


Fig. 4. Typical step configurations during the evolution of initially straight steps in Fig. 3. The panels are 120×5000 site portions extracted from the full 120×10000 net; there is considerable compression in the \hat{y} direction. The panels range from early time to near saturation. Specifically, the ratios of the image time to τ are: $\sim 1/80$, $\sim 1/8$, $\sim 1/2$, and somewhat over 3. From the step images alone, one would be hard pressed to distinguish the uphill direction, which is to the left.

the prediction is that $\tau \propto \langle w \rangle^2$. We find tolerable agreement, with a slope 18% below the expected value. We suspect that the reason behind the poorer agreement than for $L = 6$ above originates in the L -dependence of the variance associated with the peculiar algorithm used in our simulations, which allows steps to touch [21].

Independently, another argument corroborates that the kink creation rate has an activation energy of $E_d + 3E_a$: At equilibrium the creation and the annihilation rates are equal, so we compute the latter. Annihilation of kinks requires that an adatom diffuses to a step-edge notch—a kink–antikink pair, whose density is $n_{k-ak} \approx \exp(-2E_k/k_B T) \approx \exp(-2(E_d/2)/k_B T)$. Since the equilibrium adatom density is $c_{eq} = \exp(-2E_d/k_B T)$, the annihilation rate of kinks at a step edge is proportional to $Dc_{eq}n_{k-ak} \sim \exp[-(E_d + 2E_a + E_a)/k_B T]$ (cf. Ref. [24]). This in turn implies that the activation energy for the kink creation rate is $E_d + 3E_a$.

4. Discussion of results

4.1. Crucial role of kink creation

The key energy in the relaxation time is that for detaching 3-bonded atoms rather than kink atoms, a remarkable observation. Neither equilibrium nor growth processes involve 3-bonded atoms. At equilibrium, step fluctuations are controlled by the so-called step mobility, which is proportional to the emission rate of adatom from kinks [25], which involves 2-bonded atoms. Hence, the relaxation towards equilibrium should also be controlled by the step mobility. However, our results evidently contradict this notion in the far-from-equilibrium regime.

One possible explanation of our remarkable finding is that kinks have to be formed first, requiring the extraction of atoms from straight steps. (As noted earlier, the addition of an atom to a step edge also creates kink–antikink pairs, but the “tooth” is of class N_1 , so very short-lived.) In that case, our initial configuration, in which steps are perfectly parallel and straight may be introducing a bias in the results. To investigate this possibility, we considered two other initial states with equal numbers of kinks and antikinks,

⁴ The factor of 4 accounts for the 4 directions in which an atom can move from a general position on the surface. In some treatments it is absorbed into v_0 .

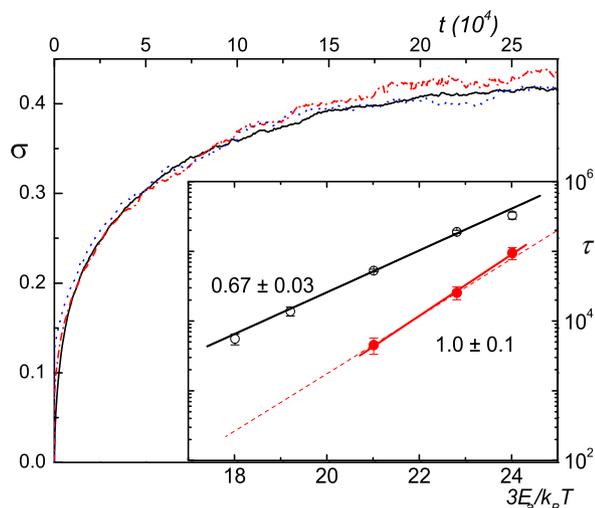


Fig. 5. Checks of dependencies on initial conditions and update moves. Evolution of the standard deviation σ of the TWD for three initial configurations: straight steps (solid, black), “decimated” edge (dash-dotted, red), and crenelated (dotted, blue) edge. The $120 \times 10,000$ lattice has 20 steps, with $L=6$; we choose $T=580$ K, $E_d=1$ eV, and $E_a=0.4$ eV. For equilibrated non-interacting (free-fermion-like, $q=2$) steps, σ approaches $0.42=\sigma_w$, as expected. The smooth curve is a fit to an exponential approach to saturation. The smooth curve is $\sigma(t)=\sigma(\infty)[1-\exp(-t/74852)]^{1/2}$. Inset: Surface azimuthally misoriented by 0.0005 radians, forcing 5 kinks along the 10^4 -site steps. The three filled (red) circles represent runs with an initial “perfect” configuration of 5 straight 2000 -site segments; the reduced slope, indicated by the solid line, is 3.0 ± 0.3 , in excellent agreement with the steeper line in Fig. 2, indicated here with a dashed line, having reduced slope of 2.9 ± 0.1 . Thus, processes in which 3 bonds are broken govern the scaling of the relaxation time τ (given in sec. as in Fig. 1). The open circles are from runs with 3-bonded atoms immobile; the corresponding reduced slope is 2.0 ± 0.1 , so that now 2-bonded atoms control the (much larger) τ . (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

i.e., in which one produces kinks by adding atoms to (or removing atoms from) a straight edge.⁵ In the “decimated” case every tenth atom along a straight step is removed; in the other case every other atom is removed to create a “fully kinked” step, so that the edge resembles dentil molding or castle crenelations. In Fig. 5 we plot the resulting evolution of the standard deviation σ of the TWD for the three cases. During the early-time rapid spreading of the initial sharp TWD, the slope increases with the number of initial kinks. In this regime, our model is not expected to apply (nor should any other 1D, continuous model); indeed, Eq. (3) does not describe the steep initial part of the traces very well. After about 4×10^4 MCS the curves are essentially indistinguishable within the noise level. The smooth curve is a one-parameter best fit of the data for the initially straight step by $\sigma(t)/\sigma(\infty) = 1 - \exp(-t/\tau)$. For this case we find $\tau \approx 7.5 \times 10^4$ (in units that are essentially sec.).

We formulated the crenelated configuration because it creates at the outset a high density of atoms of class N_1 : 8×10^{-2} atoms/site, three orders of magnitude greater than the equilibrium density of 5×10^{-5} . These atoms quickly lead to a burst of adatoms that should quickly thermalize the step configurations. If it were only the supply of adatoms that limits equilibration, then for this scenario the subsequent creation of kinks would not be crucial. Evidently this is not so; even for the crenelated case, the relaxation

kinetics are determined by the rate of creating kinks–antikinks pairs, with the usual energy barrier.

To corroborate that kink–antikink creation is indeed the rate-limiting process, we computed the relaxation rate of a surface with steps azimuthally misoriented so as to create kinks via screw boundary conditions in the \hat{y} direction. Specifically, in the initial state the in-plane misorientation slope was set at 0.0005 , so that geometry forces the existence of 5 kinks for $L_y=10,000$. Keeping the diffusion barrier fixed at 1 eV, we varied E_a . The results are shown in the inset of Fig. 5 as filled circles. We computed just three points, but clearly, essentially no difference is found with respect to the relaxation rate of straight $[100]$ steps (the red line in Fig. 2). The latter is drawn as a dashed line in the inset. The fitted slope to the data (times $k_B T$) is 3.0 ± 0.3 , fully consistent with 3-bonded ledge atoms being responsible for the rate-limiting process. We also checked that the relaxation rate is enormously slowed if 3-bonded atoms are kept immobile: Fitting the distribution width with Eq. (3), we ramped E_a while holding fixed $E_d=1$ eV. The extracted relaxation times are shown in the inset of Fig. 5 as open circles. The reduced slope is 2.0 ± 0.1 , consistent with 2-bonded kink atoms providing the rate-limiting process for the step motion in this case. The characteristic time is at least an order of magnitude larger than the previous case, which can be interpreted as due to the inability to create new kink sites, so that the number of sources for 2-bond escape of atoms to the straight segments of the step is limited to the initial 5 kinks. Without the azimuthal misorientation, this surface would be inert. Furthermore, the eventual width of the distribution, σ_{sat} is only about half the size of the 3-bond case. Thus, at least over the course of our long runs, the surface is never able to equilibrate.

We considered the number of N_2 sites, typically kinks along steps. This quantity rose much more rapidly than the variance. Referring to the main plot of Fig. 5, N_2 achieves its saturation value by $t \approx 3 \times 10^4$. Thus, the process controlling τ is not the initial formation of an adequate number of kinks but rather the maintenance of this number. For our chosen energies, about 1 in 30 sites along a step was a kink, far higher than in our azimuthally slightly-misoriented case.

For [unphysically] large values of E_a , the two straight lines in the inset of Fig. 5 cross, so that processes breaking two bonds become rate-limiting. However, since τ depends on the dimensionless ratio $E_a/k_B T$, this crossing should also happen at low temperatures. For our far-from-equilibrium conditions, we expect that the two-bond-breaking rate is roughly proportional to the number of forced kinks, and so the azimuthal misorientation, while the three-bond-breaking rate is relatively insensitive to this angle. Hence, as one gets closer to straight initial steps, the crossover occurs at ever lower temperatures. To model this behavior, we could take the overall rate to be N_2 and N_3 times their respective rates from the inset of Fig. 5. Unfortunately, the slow evolution of the lattice at the low temperatures precludes our investigating this crossover issue with KMC.

In a different limit, we applied a similar analysis of the variance of the TWD to a vicinal (001) surface misoriented in along an azimuth rotated 45° so as to have zig-zag $[110]$ steps. For such steps, every outer atom has $i=2$ lateral neighbors. Our analysis then shows that these 2-bond kink atoms produce the rate-limiting step, with a slope of 2 in the equivalent of the plot of τ vs. E_a in the inset of Fig. 5. This system has some idiosyncratic behavior due to the ease of creating fluctuations of steps from their mean configuration. Discussions of these subtleties would cloud the focus of this paper. Hence, we defer details to a future communication [21].

In short, we reach the striking conclusion that the equilibration of a terrace width on a vicinal (001) simple cubic crystal with close-packed $[100]$ steps (or steps not far from close-packed)

⁵ For initial states small polar misorientations and so with small densities of kinks with one orientation, equilibration of the step shape still involves creation of new kinks, and this in turn implies detachment of 3-bonded atoms, since detachment of an atom from a kink does not change the number of kinks. For larger misorientations and kink densities, more subtle effects—with profound implications—come into play. Since these findings distract from the focus of this Communication, we defer discussion to a separate paper [21].

and the fluctuations of the same terrace width at equilibrium are qualitatively different phenomena in the time regime of our KMC calculations. The latter can take place with a constant number of kinks, while the former requires creation of new kink–antikink pairs. This single well-defined process sets the time scale for far-from-equilibrium relaxation. Fluctuations from the equilibrium distribution, having the form of Eq. (2), will not lead to arbitrary initial configurations such as a perfect cleaved crystal with straight, uniformly-spaced steps. It seems that the near-equilibrium regime is reached only at time scales several times longer than considered in our simulations, when the variance is relatively close to saturation.

4.2. Higher moments of the TWD

So far, we have only considered the first two moments of the TWD. Several different distributions might account for these two moments. As a further check that $P_2(s, \tilde{t})$ describes the KMC data well, we study higher moments of the TWD in comparison with the analytical expressions in Eqs. (A16) and (A17). This is a far more demanding test than considering just the variance; nonetheless, in the temporal regime under study, semiquantitative accounting for the KMC data is achieved.

In Fig. 6 we plot the second, third, and fourth moments (with respect to the origin) for initially-straight steps. (Since we wish to compare with $P_2(s, \tilde{t})$, we divide the “raw” KMC j th moment by the j th power of the mean spacing to determine $\mu_j(\tilde{t})$.) For each moment there is a steady rise (from unity) that approaches the equilibrium value $\mu_j(\infty)$ exponentially. (Steps which are initially decimated or crenelated behave similarly, though in somewhat “noisier” fashion.) For μ_2 , μ_3 , and μ_4 , these saturation values agree well with the analytic results $3\pi/8 \approx 1.18$, $\pi/2 \approx 1.57$, $15\pi^2/64 \approx 2.3$, respectively, as which can be read off Eqs. (A12), (A16), and (A17). For ease of comparison, we plot $\mu_j(\tilde{t})/\mu_j(\infty)$ in Fig. 6, so that each normalized moment approaches unity. The approach to saturation is evidently slower for successively higher moments.

To make contact between the moments extracted from the KMC data and the analytic expressions in dimensionless units arising

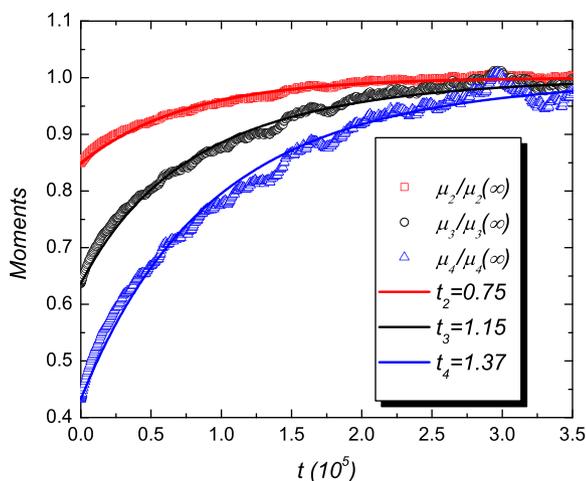


Fig. 6. From top to bottom, the second (squares, red), third (circles, black), and fourth (triangles, blue) moments (with respect to the origin) of the evolving KMC-generated TWDs for initially straight configurations and the same parameters as in Fig. 5; for ease of comparison, the j th moment $\mu_j(\tilde{t})$ is divided by its equilibrium value $\mu_j(\infty)$. To make contact with the exponential approach of these moments to their saturation values, one must rescale the KMC time. For μ_2 the rescaling factor, as in Fig. 5, is 7.5×10^4 . For μ_3 and μ_4 , the approach to saturation is progressively slower; the rescaling times are 1.15×10^5 and 1.37×10^5 , $3/2$ and $9/5$ as large, respectively. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

from our Fokker–Planck analysis, we seek whether by rescaling KMC times by some t_j leads to a good description of the data by the deduced moments. For μ_2 such a rescaling factor t_2 , with $t_2 = 7.5 \times 10^4$, was already used in analyzing the data in Fig. 5. In other words, we adjust t_2 so that $\mu_2(t/t_2)/\mu_2(\infty)$ from Eq. (A12) fits the data as well as possible, as illustrated in Fig. 6. For μ_3 and μ_4 , the approach to saturation is progressively slower; the rescaling times t_3 and t_4 , similarly obtained, are 1.15×10^5 and 1.37×10^5 , $3/2$ and $9/5$ as large, respectively.

The analytic expressions for the four moments, given in Eqs. (A12), (A13), (A16), and (A17), all approach saturation asymptotically from below like $\exp(-\tilde{t})$. Indeed, $P(s, \tilde{t})$ itself approaches $P_2(s)$ like $\exp(-\tilde{t})$, so all its moments must do likewise. However, in the temporal regime corresponding to the KMC simulations, higher-order terms cause the evident exponential-like approach to depend on \tilde{t}/τ rather than simply \tilde{t} , where τ is some effective time constant of order unity.

To determine τ_j we consider in Fig. 7 the evolution of $-\tilde{t}/\ln[1 - \mu_j(\tilde{t})/\mu_j(\infty)]$ for each moment ($j=2, 3, 4$). To the degree that this trace is horizontal, the Ansatz is appropriate. The thin curves in Fig. 7 show that this assumption becomes progressively better as time advances. While all three curves eventually converge to unity, in the time regime under consideration the three moments have significantly different time constants, with the higher moments having progressively larger magnitudes, consistent with the KMC findings displayed in Fig. 6.

The inset of Fig. 7 displays the first moment of $\mu_1(\tilde{t})$. As described in the Appendix, it is a couple percent smaller than the proper value of unity in the region around $\tilde{t} \approx 1$. (This is clearly a deficiency only of the analytic results. The KMC data have $\mu_1(\tilde{t}) \equiv 1$ by construction.) In order to estimate how much the mean-field approximation affects higher-order moments, we compute the “corrected” moments by recasting $P(s, \tilde{t})$ as a function of $s/\mu_1(\tilde{t})$ rather than s . Then the “corrected” moments are $\mu_j^{\text{corr}}(\tilde{t}) = \mu_j(\tilde{t})/\mu_1^j(\tilde{t})$, where, obviously, $\mu_1^{\text{corr}}(\tilde{t}) \equiv 1$. The higher moments $\mu_2^{\text{corr}}(\tilde{t})$, $\mu_3^{\text{corr}}(\tilde{t})$, and $\mu_4^{\text{corr}}(\tilde{t})$ are displayed as the thick curves in Fig. 7. These curves flatten considerably sooner than the thin curves, and to a value $\sim 1/2$, reminiscent of Eq. (A15). For these

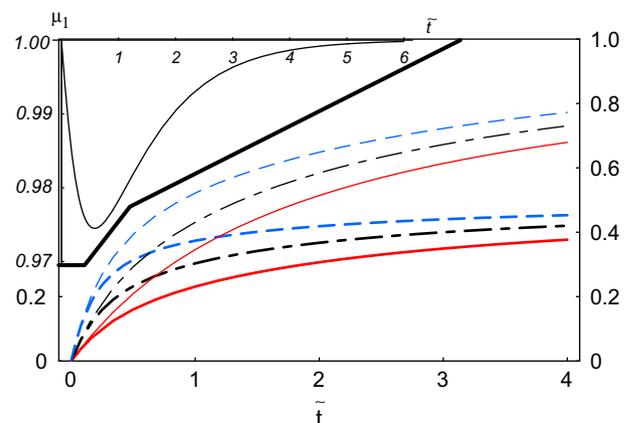


Fig. 7. Analytic results for the moments of the evolving TWD predicted by the (1D continuum) Fokker–Planck theory: Plot of the effective exponential decay time of the difference between the evolving second (solid, red), third (long-short-dashed, black), and fourth (dashed, blue) moments of the solution to Eq. (1), $P_{q=2}(s, \tilde{t})$, given explicitly in the appendix in Eqs. (A12), (A16), and (A17), and their steady-state, asymptotic values associated with Eq. (2). As in the KMC data in Fig. 6, the decay is significantly slower for the higher moments. The thicker set of curves corrects for the modest deficiency of the the first moment $\mu_1(\tilde{t})$; $\mu_1(\tilde{t})$ is depicted in the inset (upper left, italicized axes labels) and given analytically in Eq. (A13). See text for details. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

curves, $(\tau_3 - \tau_2)/\tau_2$ is somewhat over 0.1 while $(\tau_4 - \tau_2)/\tau_2$ is about twice as large, capturing the trend of the numerical data. If we use the analytic expressions for the corrected moments as the rescaling factors, then the time rescaling factors are 1.75×10^5 , 1.9×10^5 , and 2.0×10^5 , respectively. Then $(\tau_3 - \tau_2)/\tau_2 \approx 0.09$ and $(\tau_4 - \tau_2)/\tau_2 \approx 0.14$, closer to the KMC values.

As detailed in the Appendix, all moments eventually approach their saturation value by a term proportional to $\exp(-\bar{t})$, the leading term of a power series in that exponential. Those higher-order terms, which underlie the effective decay times we find in the computed time regime, become insignificant eventually, but only on a time scale several times that in the KMC simulations, when the variance is relatively close to its saturation value. One might speculate whether the characteristic time after which the higher moments behave as $\exp(-\bar{t})$ corresponds to the time for which one is close enough to equilibrium that two-bond-breaking processes control the relaxation rate. Meaningfully addressing that possibility, e.g. by scrutinizing N_2 at times several times as great as in our reported simulations, is beyond the scope of our simulations.

Accounting for the skewness and the kurtosis poses a more difficult test for our kinetic Monte Carlo simulation of our model. Due to the interplay of several moments in computing these statistics, our numerical data is not adequate to test these predicted behaviors meaningfully. Such analyses are arguably the most stringent tests, in which we seek differences from random-walk, Gaussian behavior; they corroborate the conclusion above that the surface never fully equilibrates. Far more extensive computations might clarify this matter, but are beyond the scope of our present analysis.

5. Comparison with Gruber–Mullins

To help place our approach in context, we compare it with previous approximations in the literature based on the fermion description of the fluctuating steps. The celebrated Gruber–Mullins (GM) approximation [26] considers a fluctuating step between two fixed neighbors treated as rigid boundaries. In fermion language, the step is the 1D trajectory of a quantum particle confined to the segment $(0, 2\langle w \rangle)$ by an infinite potential. Two cases are easily treated: For non-interacting steps, the fluctuating step is then the trajectory of a free fermion, and is equivalent to a classical particle performing a random walk in a potential of the form [27]

$$V(x) = -2 \ln[\sin(\pi x/2\langle w \rangle)]. \quad (5)$$

Then $s \equiv w/\langle w \rangle$ obeys the Langevin equation

$$\dot{s} = \frac{\pi\gamma}{\langle w \rangle^2} \frac{1}{\tan(\pi s/2)} + \eta. \quad (6)$$

This approximation preserves the logarithmic behavior of the repulsive potential at short range; even the amplitude is correct: $(2b_0s - q/s)|_{q=2} = (8s/\pi - 2/s)$ in Eq. (1) is replaced by $-\pi \cot(\pi s/2)$, nearly the same for $s < 0.7$. However, the GM potential has bogus symmetry about $\langle w \rangle$, truncating the long-range tail of $P(s)$.

For strongly interacting steps, the fluctuating step feels an (approximately) quadratic confining potential, and the TWD distribution is predicted to be Gaussian [26]. In our formalism the replacement is now $(s-1)/\sigma_G^2$, where σ_G^2 is the variance of the Gaussian TWD (so half the variance of the associated ground-state wavefunction, which we used to construct the FP potential [27].) This replacement should be compared with $(2b_0s - q/s) \approx (q + \frac{1}{2})s - q/s$ [28]. In the GM approximation, $\sigma_G^{-2} = [12q(q-2)]^{1/2}$, with $(12)^{1/2} \approx 3.5$ replaced by $(2\pi^4/15)^{1/2} \approx 3.6$ if the interactions with all steps rather than just the two bounding steps are considered [28]. An improved approximation (“modified

Grenoble”) gives $\sigma_G^{-2} \approx 2.1q$ [28]. Our expression for the FP potential now differs at small s from that derived for these approximations because the Gaussians actually extend (unphysically, albeit with insignificant amplitude) to negative values of s . Our approach is globally superior to the celebrated GM approximation (as well as to the usual alternatives [1]), both quantitatively and qualitatively, for all physical values of the step–step interaction strength [29]. Moreover, our FP Eq. (1) is fully soluble, so that the TWD can be obtained analytically as a function of time.

6. Summary

In summary, we have shown that the relaxation time of the variance of the solution of our Fokker–Planck equation for step relaxation on a vicinal surface can be fit to the comparable variance in a kinetic Monte Carlo simulation of the standard simple model of atomistic processes at surfaces. This time has Arrhenius behavior that is related to microscopic processes, substantiating that this FP approach can offer useful physical insight into the evolution of complex surface structures toward equilibrium. Thus, once the continuum formalism becomes appropriate, the FP time in some sense tracks actual time in our model of an evolving physical system of steps with no energetic repulsion. The formalism also readily allows such repulsions, inviting future simulations to test how well the Fokker–Planck formalism describes such systems. Since the steps communicate from the outset, the continuum formalism might apply sooner. For the situation we have considered, we have presented several pieces of evidence that the rate-determining process in step relaxation is the creation of kink–antikink pairs, distinctly different from the processes associated with step meandering near equilibrium. We have also examined higher moments of the distribution, both analytically and with simulations. While we make no pretense that our approach is either exact or a formal theory, we have shown that it can be a fruitful way to treat relaxation of steps on surfaces. Many avenues of extension are possible.

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Appendix. Derivation of Eq. (1) and some consequent results

In this appendix we expand the derivation of the Fokker–Planck equation given in Ref. [2], as well as correct some algebra oversights in intermediate steps presented there. We also present some simpler expressions for quantities of interest that arise for the case of non-interacting [energetically] steps ($q=2$) investigated in the reported computations.

As in Ref. [2], we begin with the correspondence found by Dyson between RMT and his Coulomb gas model [8]: N classical particles on a line, interacting with a logarithmic potential, and confined by an overall harmonic potential. Dyson’s model helps our understanding of the fluctuation properties of the spectrum of complex conserved systems. This model can be generalized to the dynamic Brownian motion model, in which the N particles

are subject, besides the mutual Coulomb repulsions, to dissipative forces [30]. The particle positions x_i then obey Langevin equations

$$\dot{x}_i = -\gamma x_i + \sum_{i \neq j} \frac{\hat{Q}}{x_i - x_j} + \sqrt{\Gamma} \eta, \quad (\text{A1})$$

where η is a delta-correlated white noise and $\hat{Q} (\propto Q)$ is the “charge” of each particle. The probability of finding the particles at the positions $\{x_n\}$ at time t is the solution of the multidimensional FPE

$$\frac{\partial P(\{x_n\}, t)}{\partial t} = \sum_i \frac{\partial}{\partial x_i} \left[\frac{\partial}{\partial x_i} P(\{x_n\}, t) + \gamma x_i P(\{x_n\}, t) \right] - \sum_{i \neq j} \frac{\partial}{\partial x_i} \left[\frac{\hat{Q}}{x_i - x_j} P(\{x_n\}, t) \right]. \quad (\text{A2})$$

In the 1D case, γ^{-1} would essentially be the variance of the stationary distribution. Narayan and Shastry [9] showed that the CS model is equivalent to Dyson’s Brownian motion model, in the sense that the solution of the FPE (A2) may be written as $P(\{x_n\}, t) = \psi(\{x_n\}, t) \psi_0(\{x_n\}, t)$, where $\psi(\{x_n\}, t)$ is the solution of a Schrödinger equation with imaginary time, derived from the CS Hamiltonian. The deterministic force of Eq. (A1)

$$F(x_m) = -\gamma x_m - \sum_{k>m} \frac{\hat{Q}}{x_k - x_m} + \sum_{q<m} \frac{\hat{Q}}{x_m - x_q} \quad (\text{A3})$$

so that

$$F(x_{m+1}) - F(x_m) = -\gamma(x_{m+1} - x_m) - \hat{Q} \left[\frac{-2}{x_{m+1} - x_m} + \sum_{k>m+1} \frac{x_{m+1} - x_m}{(x_k - x_{m+1})(x_k - x_m)} + \sum_{q<m} \frac{x_{m+1} - x_m}{(x_{m+1} - x_q)(x_m - x_q)} \right]. \quad (\text{A4})$$

Our goal is to find the distribution of widths w . Mindful of the Gruber–Mullins approach [26], we construct a single-“particle,” mean-field approximation in which the dynamical variable is the nearest-neighbor distance $w_m \equiv x_{m+1} - x_m$. To decouple the force on w_m from the other particles, we assume—in the spirit of GM—that the denominators $(x_k - x_{m+1})(x_k - x_m)$ in Eq. (A4) are replaced by their mean values, the average being taken in the stationary state:

$$\langle (x_k - x_{m+1})(x_k - x_m) \rangle_{st} = \langle w^2 \rangle_{st} (k - m - 1)(k - m), \quad (\text{A5})$$

Each of the two sums in Eq. (A4) then simplifies greatly, taking the form

$$\frac{(x_{m+1} - x_m)}{\langle w^2 \rangle_{st}} \times \left(\sum_{p=1}^N \frac{1}{(p+1)p} = \frac{N}{N+1} \xrightarrow{N \rightarrow \infty} 1 \right). \quad (\text{A6})$$

Hence, the interaction of a particle pair with all other particles acts on average as a harmonic potential, increasing the “spring constant” of the external confining potential. We arrive at a single-particle Langevin equation for the terrace width w :

$$\frac{dw}{dt} = -2 \left[\left(\frac{\gamma}{2} + \frac{\hat{Q}}{\langle w^2 \rangle_{st}} \right) w - \frac{\hat{Q}}{w} \right] + \sqrt{2\Gamma} \eta. \quad (\text{A7})$$

Our goal is to convert Eq. (A7) into a FPE for which Eq. (2) is a steady-state solution. We change to dimensionless variables $s \equiv w/\langle w \rangle_{st}$ and $\tilde{t} \equiv \Gamma t / \langle w \rangle_{st}^2$. Treating γ as a self-consistency parameter and recognizing $\hat{Q} = Q\Gamma/2$, we set $\gamma = \Gamma/\langle w \rangle_{st}^2$. Then the coefficient in parentheses in Eq. (A7) becomes

$$\frac{(1+Q)\Gamma}{2\langle w^2 \rangle_{st}} = \frac{b_Q \Gamma}{\langle w \rangle_{st}^2}, \quad (\text{A8})$$

using the second moment of $P_Q(s) [(s^2) = (Q+1)/(2b_Q)]$. Furthermore, if $\langle \eta(t)\eta(\tilde{t}) \rangle = \delta(t - \tilde{t})$, then $\tilde{\eta}(\tilde{t}) \equiv \sqrt{2/\Gamma} \langle w \rangle_{st} \eta(t)$ satisfies

$\langle \tilde{\eta}(\tilde{t})\tilde{\eta}(\tilde{t}') \rangle = \delta(\tilde{t} - \tilde{t}')$. With these results, we recast Eq. (A7) into the Langevin equation

$$\frac{ds}{d\tilde{t}} = - \left[2b_Q s - \frac{Q}{s} \right] + \tilde{\eta} \quad (\text{A9})$$

and thence the sought-after FPE given in Eq. (1).

To solve Eq. (1) we must specify the initial distribution in s_0 . For an initial (at $\tilde{t} = 0$) sharp distribution $\delta(s - 1)$, corresponding to a perfectly cleaved crystal, the solution is essentially written down by Montroll and West [31,32]:⁶

$$P(s, \tilde{t}) = 2\tilde{b}_Q s^{\frac{Q+1}{2}} e^{\frac{(Q-1)\tilde{t}}{4}} I_{\frac{Q-1}{2}} \left(2\tilde{b}_Q s e^{-\frac{\tilde{t}}{2}} \right) e^{-\tilde{b}_Q (s^2 + e^{-\tilde{t}})}, \quad (\text{A10})$$

where $\tilde{b}_Q \equiv b_Q/(1 - e^{-\tilde{t}})$. In the limit of long times, we showed in Ref. [2] that, as \tilde{t} increases, this $P(s, \tilde{t})$ approaches $P_Q(s)$ of Eq. (2).

For the particular case $Q=2$, \tilde{b}_Q becomes $4/[\pi(1 - e^{-\tilde{t}})]$, while $I_{\frac{1}{2}}(z) = \sqrt{2/(\pi z)} \sinh(z)$. Then Eq. (A10) simplifies to

$$P(s, \tilde{t}) = \frac{2^{\frac{3}{2}} e^{\frac{3\tilde{t}}{4}} s}{\pi \sin h^{\frac{1}{2}}(\frac{\tilde{t}}{2})} \sinh \left(\frac{(4/\pi)s}{\sin h(\frac{\tilde{t}}{2})} \right) \exp \left[-\frac{4(s^2 + e^{-\tilde{t}})}{\pi(1 - e^{-\tilde{t}})} \right]. \quad (\text{A11})$$

In experiments, $P(s)$ is generally characterized just by its variance $\sigma^2 \equiv \mu_2 - \mu_1^2$, which can be calculated from its second and first moments, μ_2 and μ_1 , respectively:

$$\mu_2(\tilde{t}) = \frac{3\pi}{8} u_{\tilde{t}} + e^{-\tilde{t}} = \left(\frac{3\pi}{8} - 1 \right) (1 - e^{-\tilde{t}}) + 1 \quad (\text{A12})$$

$$\mu_1(\tilde{t}) = \frac{1}{2} \left[u_{\tilde{t}}^{\frac{1}{2}} \exp \left(\frac{-4/\pi}{e^{\tilde{t}} - 1} \right) + \left\{ 1 + \left(\frac{8}{\pi} - 1 \right) e^{-\tilde{t}} \right\} \Upsilon(\tilde{t}) \right] \quad (\text{A13})$$

where, for brevity, we take $u_{\tilde{t}} \equiv 1 - \exp(-\tilde{t})$, which obviously approaches unity exponentially from below. Furthermore, we write $\Upsilon(\tilde{t}) \equiv (\pi/4) \exp(\tilde{t}/2) \text{erf}(2/[\pi(\exp(\tilde{t}) - 1)]^{1/2})$, where erf is the error function [33]; $\Upsilon(\tilde{t})$ also approaches unity exponentially, but from above, after rising initially from $\pi/4$ to about 1.01.

Scrutiny of Eq. (A13) reveals that each of the two summands in the square brackets approaches 1 for large \tilde{t} . As \tilde{t} approaches 0, the first summand vanishes while the second rises to 2. Thus, $\mu_1(\tilde{t})$ has the expected value for vanishing and large \tilde{t} , as illustrated in the inset of Fig. 7. There is, however, an initial rapid drop, reaching a minimum of about 0.9745 around $\tilde{t} = 0.582$, and then rising smoothly, reaching 0.99 by $\tilde{t} = 1.95$, 0.995 by $\tilde{t} = 2.685$, and 0.999 by $\tilde{t} = 4.33$. The small deviation from unity is presumably due to the approximations in using Eq. (A5) to reach Eq. (A7), which apparently break the symmetry of the fluctuations of the steps (m and $m+1$) bounding w_m [34].

To the extent that this deviation is negligible (and in any case for qualitative purposes), we get

$$\sigma^2(\tilde{t})|_{\mu_1 \equiv 1} = \sigma_W^2 (1 - e^{-\tilde{t}}). \quad (\text{A14})$$

If we numerically evaluate $\sigma^2(\tilde{t})$ using Eqs. (A12) and (A13), we find a similar expression but with a more rapid rise to the equilibrium result; remarkably, it is well approximated by

$$\sigma^2(\tilde{t}) = \sigma_W^2 (1 - e^{-2\tilde{t}}). \quad (\text{A15})$$

reminiscent of the solution of the Fokker–Planck equation for a Brownian particle in a quadratic potential.⁷ In any case, the Arrhenius behavior of the characteristic time of the exponential will not be affected by such modest changes in the prefactor.

⁶ For stationary initial distributions, Stratonovich proceeds by separation of variables, finding the spatial eigenfunctions in terms of Laguerre polynomials. Taking into account Stratonovich’s unconventional normalization of his Laguerre polynomials and, eventually, removing the initial stationary distribution in his Eq. (4.77), we can recapture Montroll and West’s [31] Eq. (3.32).

⁷ Specifically, Eq. (3.21) of Ref. [31] shows that the variance of the Gaussian distribution is proportional to $1 - \exp(-2t)$.

If the Fokker–Planck description of step relaxation is robust, then the higher moments of $P(s, \tilde{t})$ should also characterize those moments extracted from the KMC data, as displayed in the text in Fig. 6. Thus, we present explicit analytic formulas for the third and fourth moments:

$$\mu_3(\tilde{t}) = \left(\frac{1}{2} e^{-\tilde{t}} u_i^{1/2} + \frac{5\pi}{16} u_i^{3/2} \right) \exp\left(-\frac{4/\pi}{e^{\tilde{t}} - 1}\right) + \left(\frac{4}{\pi} e^{-2\tilde{t}} + 3u_i e^{-\tilde{t}} + \frac{3\pi}{16} u_i^2 \right) \Upsilon(\tilde{t}) \quad (\text{A16})$$

$$\sim \frac{\pi}{2} - \left(\frac{3\pi - 8}{4} \right) e^{-\tilde{t}} - \left(1 - \frac{3\pi}{16} - \frac{4}{5\pi} \right) e^{-2\tilde{t}} + \mathcal{O}(e^{-3\tilde{t}}) \quad (\text{A16a})$$

$$\approx 1 + \left(\frac{\pi}{2} - 1 \right) (1 - e^{-\tilde{t}/\tau_3}) \quad (\text{A16b})$$

$$\mu_4(\tilde{t}) = \frac{15\pi^2}{64} u_i^2 + \frac{5\pi}{4} e^{-\tilde{t}} u_i + e^{-2\tilde{t}} \quad (\text{A17})$$

$$= \frac{15\pi^2}{64} - \frac{5\pi}{32} (3\pi - 8) e^{-\tilde{t}} + \left(1 - \frac{5\pi}{32} + \frac{15\pi^2}{64} \right) e^{-2\tilde{t}} \quad (\text{A17a})$$

$$\approx 1 + \left(\frac{15\pi^2}{64} - 1 \right) (1 - e^{-\tilde{t}/\tau_4}) \quad (\text{A17b})$$

In the long-time limit, $\mu_2(\tilde{t})$, $\mu_3(\tilde{t})$, and $\mu_4(\tilde{t})$ all asymptotically approach their limit $\mu_j(\infty)$ from below as $e^{-\tilde{t}}$, as shown explicitly in Eqs. (A16a) and (A17a) and by quick inspection of Eq. (A12). Indeed, all moments should approach their asymptotic limit in this way, since we can expand Eq. (A11) in its entirety as a power series in $e^{-\tilde{t}}$. Such a more detailed exploration of the crossover to the asymptotic limit is beyond the scope of this paper.

We also can write approximate expressions for $\mu_3(\tilde{t})$ and $\mu_4(\tilde{t})$ in Eqs. (A16b) and (A17b) in the form of the exact result for $\mu_2(\tilde{t})$ in Eq. (A12). By setting $\tau_3 = \tau_4 = 1$ one obtains a mediocre approximation which underestimates $\mu_3(\tilde{t})$ and $\mu_4(\tilde{t})$ by as much as 6% and 15%, respectively. A far better accounting is obtained by taking $\tau_3 \approx 0.79$ and $\tau_4 \approx 0.76$; the best values of these time constants depends weakly on the temporal range over which one seeks to optimize the agreement. The approximate expressions then underestimate the actual $\mu_j(\tilde{t})$ (by at most 2% and 3%) up to $\tilde{t} \approx 3/2$ and then overestimate it (by at most $\frac{1}{2}\%$ and 1%), respectively. Thus, $\mu_3(\tilde{t})$ and $\mu_4(\tilde{t})$ can be well described by curves starting rising smoothly from unity and decaying exponentially toward their long-time limit, but with values of t_j that are smaller than unity. Evidently the time regime relevant to the KMC simulations is before the (single-term) asymptotic limit. Finally, note that the approximate expressions based on Eqs. (A16b) and (A17b) are not used in the analysis of the moments in Section 4.2; hence, the values of the τ_j play no role there.

From these results the skewness can be expressed analytically but has an unwieldy form. However, it is semiquantitatively described by $0.4857 \tanh(\tilde{t})$ (i.e., to within $\pm 4\%$ for $\tilde{t} \geq 0.46$ and within a percent for $\tilde{t} \geq 1.9$). In other words, the skewness rises smoothly and monotonically from 0 initially to the equilibrium va-

lue. Since for large \tilde{t} , $\tanh(\tilde{t}) \sim 1 - 2 \exp(-2\tilde{t})$ we find the same approach to saturation as for the variance in Eq. (A15). The kurtosis begins at 3 but dips (to about 2.87 near $\tilde{t} = 0.5$) before rising to its equilibrium value of 3.1082. The approach to this asymptotic value is well approximated by $3.11(1 - 0.58e^{-2\tilde{t}})$.

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