Microscopic origin of excess wings in relaxation spectra of supercooled liquids

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Glass formation is encountered in diverse materials. Experiments have revealed that the dynamic relaxation spectra of supercooled liquids generically become asymmetric near the glass transition temperature \( T_g \) where an extended power law emerges at high frequencies. The microscopic origin of this ‘wing’ remains unknown, and has so far been inaccessible to simulations. Here we develop a novel computational approach and study the equilibrium dynamics of model supercooled liquids near \( T_g \). We demonstrate the emergence of a power-law wing in the numerical spectra, which originates from relaxation at rare, localized regions over broadly distributed timescales. We rationalize the asymmetric shape of the relaxation spectra by constructing an empirical model associating heterogeneous activated dynamics with dynamic facilitation, which are the two minimal physical ingredients revealed by our simulations. Our work offers a glimpse into the molecular motion responsible for glass formation at relevant experimental conditions.

The formation of amorphous solids results from the rapid growth of structural relaxation time \( \tau \) of a supercooled liquid\(^1\). Molecular motion occurs on a timescale of about \( 10^{-15} \) s at the onset temperature of glassy behaviour, but it takes about 100 s at the experimental glass transition temperature \( T_g \) (ref. \(^2\)). Over the last decades, dielectric, mechanical and light-scattering experiments kept developing to probe molecular motion over a broader frequency range with increased accuracy\(^3\). This progress reveals that the temperature evolution of \( \tau \) is just the tip of the iceberg, as relaxation spectra \( \chi''(\omega) \) measured near \( T_g \) exhibit relaxation processes taking place over an extremely large frequency window\(^4\). The overall shift in the relaxation spectra is accompanied by an equivalent broadening of about 12 decades, which is the other side of the same coin. A microscopic explanation of these slow dynamics is at the heart of glass transition research\(^1\).

High-temperature spectra reflect near-exponential relaxation in the picosecond range, but low-temperature spectra broaden into a two-step process with a stretched exponential relaxation at low frequency \( \omega \approx 1/\tau \) and a microscopic peak remaining at the picosecond timescale. In 1990, more precise experiments\(^5\)–\(^8\) showed that for a number of molecular liquids, the structural relaxation peak extends much further at higher frequencies \( \omega \tau \gg 1 \) and transforms into a power law, namely, \( \chi''(\omega) \approx \omega^{-\alpha} \), with a small exponent \( \sigma(T) \in [0.2, 0.4] \) decreasing with temperature\(^14\). Using logarithmic scales, this resembles a ‘wing’ in ‘excess’ of the \( \alpha \) peak. At \( T_g \), the wing extends over the millihertz–megahertz range with an amplitude about 100 times smaller than the \( \alpha \) peak. A universal scaling comprising the excess wing was proposed\(^14\), which can be altered by further microscopic processes\(^5\). Although this universality is debated\(^5,\)\(^10\), the presence of an excess contribution often taking the form of a wing is not\(^4,\)\(^7\).

Elucidating the nature of molecular motion responsible for the small signal in these excess wings appears daunting. Yet, experiments managed to characterize its heterogeneous nature\(^5\) and aging properties\(^9\). So far, computer simulations were unable to access the required range of equilibration temperatures and timescales to even address the question. Physical interpretations and empirical models have been proposed to explain the shape of the relaxation spectra. Some of them couple slow translational motion with an ‘additional’ degree of freedom (for example, rotational)\(^12,\)\(^13\). Others invoke spatially heterogeneous dynamics to construct a broad distribution of timescales of static\(^12,\)\(^15\)–\(^17\) or kinetic\(^18\) origin. The winged asymmetric shape then requires specific physics, such as geometric frustration\(^3\), length-scale-dependent dynamics\(^5\) or dynamic facilitation\(^9\). With specific choices, these approaches yield relaxation spectra comprising excess wings, but direct microscopic investigations testing the underlying hypotheses are still lacking.

Here we show that computer simulations can now directly observe excess wings and assess their microscopic origin. We take advantage of the recent swap Monte Carlo algorithm\(^29\) to efficiently produce equilibrated configurations of a supercooled liquid with \( \tau \approx 100 \) s. We observe their physical relaxation dynamics over 10 decades in time, up to 20 ms. We are, thus, able to probe the temperature and time regimes where excess wings are observed in experiments. We report the emergence of a power law (a wing) in the numerical spectra with the same characteristics as in experiments. We demonstrate that it is caused by a sparse population of localized regions, whose relaxation times follow a power-law distribution. These relaxed regions then coarsen by dynamic facilitation.

We construct an empirical model to illustrate how heterogeneous dynamics and dynamic facilitation generically lead to asymmetric, winged relaxation spectra.

We study size-polydisperse mixtures of \( N \) soft repulsive spheres in two and three dimensions (Methods). These models are representative computational glass formers\(^30,\)\(^31\). We use the swap Monte Carlo algorithm (designed in ref. \(^32\)) to generate \( n \in [200, 450] \) independent equilibrium configurations at temperature \( T \) down to the extrapolated experimental glass transition temperature \( T_g \). Each equilibrium configuration is then taken as the initial condition of a multiple central processing unit (CPU) molecular dynamics (MD) simulation (without swap). The \( n \) independent simulations run for up to a simulation time \( t_{\text{max}} \) of \( 1.5 \times 10^7 \) in.

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three dimensions (one week on two CPUs for $N=1,200$). We push a few two-dimensional (2D) simulations to unprecedentedly long times, up to $t_{\text{max,2D}} = 6 \times 10^4$, representing a computational time of several months. By using the relaxation time at the onset of glassy dynamics to relate the numerical and experimental timescales, our longest simulations translate into a physical time of about 20 ms for systems having an equilibrium relaxation time $\tau_o \approx 10^2$ s. This strategy is the key to observe excess wings, which would otherwise be buried underneath structural relaxation in conventional approaches\(^\text{31}\). The 2D and three-dimensional (3D) models behave similarly; therefore, we present quantitative results for the 3D model ($N=1,200$) in Figs. 1 and 3 and illustrate the relaxation process in Fig. 2 with 2D snapshots ($N=10,000$), which are easier to visualize. Quantitative results for the 2D model are provided in the Supplementary Information.

We investigate the spatiotemporal evolution of the relaxation dynamics using averaged and particle-resolved dynamic observables. In three dimensions, we measure the self-intermediate scattering function $F_\omega(t)$, averaged over the $n_i$ independent runs. We define the relaxation time $\tau_o$ by $F(\tau_o) = e^{-1}$. In two dimensions, collective long-ranged fluctuations affect the measurement of $F(t)$. We instead focus on observables that are independent of these fluctuations\(^\text{34}\) and define $\tau_o$ via the bond-orientational correlation function\(^\text{\textit{35}}\). In both two and three dimensions, we investigate the relaxation process at the particle scale via the bond-breaking correlation $C_{bb}(t)$ that quantifies the fraction of nearest neighbours lost by particle $i$ after time $t$. Starting from $C_{bb}(t = 0) = 1$, it decreases as rearrangements take place close to particle $i$, and reaches zero when its local environment is completely renewed. Precise definitions of the correlation functions are provided in Methods.

To connect with experimental results obtained in the frequency domain, we compute the dynamic susceptibility $\chi''(\omega)$ from a distribution of relaxation times $G(\log \tau)$ (refs.\(^\text{17,28})$):

$$\chi''(\omega) = \int_{-\infty}^{\infty} G(\log \tau) \frac{\omega \tau}{1 + (\omega \tau)^2} d \log \tau,$$

(1)

where distribution $G$ is related to the derivative of a time correlation function, $G(\log \tau) \approx -dF_\omega(t)/d\log t$ in three dimensions. We use the bond-breaking correlation function instead of $F_\omega$ in two dimensions. We discuss the numerical evaluation of $\chi''$ in Methods, and discussion on the statistical noise and comparison with direct Fourier transforms are provided in Supplementary Section 1 and Supplementary Figs. 1 and 2.

We start by presenting the equilibrium measurements of $F_\omega(t)$ in three dimensions (Fig. 1a), concentrating on the unexplored low-$T$ regime below the mode-coupling crossover $T_{\text{mct}} \approx 0.095$. The latter is determined by a power-law fit of $\tau_o(T)$ in the range $\tau_o / \tau_o < 10^3$, where $\tau_o \approx 3$ is the value of $\tau_o$ at the onset temperature $T_o \approx 0.20$ (ref.\(^\text{29})$. At all the temperatures, the correlations display a fast initial decay near $t \approx \tau_o$ due to fast dynamical processes. At larger times, we observe
a much slower decay to zero. As \( T \) decreases, the relaxation time grows and eventually exits the numerically accessible time window. At the lowest investigated temperatures near \( T_\alpha \), correlations appear almost constant over more than seven decades in time, suggesting a near-complete dynamic arrest. We recall that owing to the swap algorithm, all the measurements reflect genuine equilibrium dynamics, even when \( \tau_\alpha \) is larger than the simulated time by many orders of magnitude.

Our strategy allows us to directly observe the \( \alpha \) relaxation when \( \tau_\alpha < \tau_{\text{max}} \), equivalently, \( \tau_\alpha / \tau_o \leq 5 \times 10^5 \) down to \( T = 0.0755 \) (Figs. 1a,b). In this regime, the relaxation is well described by a stretched exponential \( F(t) \propto e^{(-t/\tau_\alpha)\beta} \) with an almost constant stretching exponent \( \beta \approx 0.56 \), and amplitude \( F_0 \) that changes modestly with temperature. We use this time–temperature superposition (TTS) property to estimate \( \tau_\alpha \) for \( 0.0700 \leq T \leq 0.0755 \), where the decorrelation of \( F(t) \) is sufficient\(^6\), and obtain \( \tau_\alpha \) over roughly two more decades (Fig. 1b). We finally use an Arrhenius law to extrapolate \( \tau_\alpha \) over four more decades to get a safe lower bound for the experimental glass transition temperature \( T_g \approx 0.056 \), defined by \( r_c(T_g)/r_o = 10^{12} \) (ref. \(^{23} \); Methods).

The corresponding relaxation spectra are shown in Fig. 1c for the 3D model. All of them display a peak at high frequency \( \omega_\alpha \approx 1/\tau_\alpha \), corresponding to the short-time decay of \( F(t) \). A low-frequency peak near \( \omega \approx 1/\tau_o \) is also visible. As \( T \) decreases, this peak shifts to lower frequencies and eventually exits the accessible frequency window. When the \( \alpha \) peak is not directly measured, we extrapolate its shape by inserting the above stretched exponential form for \( F(t) \) into equation (1). We use \( \beta = 0.56 \); \( \tau_o \) is given by the Arrhenius extrapolation and constant \( F_0 \). The tiny temperature dependence of \( F_0 \) is immaterial on a logarithmic scale (Fig. 1c). The resulting \( \alpha \) peaks are shown in Fig. 1c with dashed lines that smoothly merge into the measured data at the highest temperatures, validating our procedure.

As \( T \) decreases, the measured susceptibility and \( \alpha \) peak deviate increasingly from one another, the data being systematically in excess of the \( \alpha \) peak. Since the Arrhenius extrapolation underestimates \( \tau_\alpha \), this excess is at worst slightly underestimated and cannot be accounted for by a vertical shift that would require unphysical values of \( F_0 \) and \( \beta \). At the lowest \( T \), where the \( \alpha \) peak no longer interferes with the measurements, the spectra are well described by a power law \( \chi' \propto \omega^{-\nu} \) at low frequencies, with exponent \( \nu \approx 0.38 \) slightly decreasing with \( T \), and an amplitude about 100 times smaller than the \( \alpha \) peak. The relaxation spectra of the 2D model (Supplementary Fig. 3) exhibit similar features with exponent \( \nu_{2D} \approx 0.45 \), which is fairly close to the one found in three dimensions. In our simulations, the measured spectra do not exhibit a secondary peak separated from the \( \alpha \) relaxation, and cannot be interpreted using an additive \( \beta \) process\(^{31} \). Therefore, close to \( T_g \), the numerical spectra follow a power law over a similar frequency range, with a similar exponent and similar amplitude as the excess wings obtained experimentally, suggesting that simulated glass formers display excess wings resembling observations in molecular liquids.

We take advantage of the atomistic resolution offered by simulations to explore the microscopic origin of excess wings and provide a physical interpretation of the spectral shapes. We illustrate the relaxation dynamics with 2D snapshots, which are easier to render and interpret. We confirm that the same mechanisms are observed in three dimensions. In Fig. 2, we show the 2D snapshots illustrating how structural relaxation proceeds at temperature \( T_{2D} \approx 0.09 \) (we estimate \( T_{2D} \approx 0.07 \)) for which \( \tau_\alpha / \tau_o \approx 10^7 \), which corresponds to around 10 ms in physical time. This temperature is the lowest for which the \( \alpha \) relaxation can be observed in the numerical window, and is considerably lower than the mode-coupling crossover near \( T_{\text{MC;2D}} \approx 0.12 \). The images are shown at logarithmically spaced times \( t \) in the range \( t/\tau_o \in [10^{-4}, 1] \). The particles are coloured according to \( C_0(t) \): red particles have relaxed and blue ones have not. Supplementary Fig. 3 shows the relaxation spectrum measured at this temperature.

For \( t/\tau_o \ll 1 \), relaxation starts at a sparse population of localized regions that independently distributed throughout the sample over broadly distributed times. This conclusion holds over a large range of temperatures down to \( T_g \) for both \( d = 2 \) and \( d = 3 \). As time increases, newly relaxed regions continue to appear, but a second mechanism becomes apparent (Fig. 2) as regions that have relaxed in one frame typically appear larger in the next. This growth of relaxed regions (Fig. 2) is the signature of dynamic facilitation\(^5\). More precisely, we observe that from one frame to the next, relaxation events keep accumulating at similar locations, which results in mobile particles undergoing multiple relaxations and mobility propagating to nearby particles. Also, the slowest regions are typically ‘invaded’ at \( t \approx \tau_\alpha \) from their faster boundaries. Dynamic facilitation has been identified before at high temperatures above the mode-coupling crossover\(^{37,38} \). Our investigations show that it becomes a central physical mechanism for structural relaxation near \( T_g \).

We concentrate on the early times when the power-law spectra are observed. The visualization suggests that clusters of relaxed particles appear at sparse locations. We now establish that these early relaxation events are responsible for the excess wing. To this end, we define mobile (\( C_0(t) < 0.55 \)) and immobile (\( C_0(t) \geq 0.55 \)) particles; the threshold value near 0.50 is determined requiring self-consistency with alternative mobility definitions based on displacements. We identify connected clusters of mobile particles by performing a nearest-neighbour analysis (Methods), and investigate the statistical properties of relaxed clusters. In particular, we find that the excess wing regime at \( t/\tau_o \ll 1 \) is dominated by the appearance of new clusters, whereas the growth of existing clusters dominates at later times. Figure 3 shows the distribution \( \Pi(\tau) \) of waiting times \( \tau \) for the appearance of new clusters in three dimensions. For \( T \leq 0.07 \), we cannot measure the entire distribution, which is, thus, determined up to an uninteresting prefactor. The corresponding 2D results are shown in Supplementary Fig. 4.

At the highest investigated temperature, near \( T_{\text{MC;2D}} \), the distribution \( \Pi(\log_{10}(\tau)) \) in Fig. 3 is already very broad, with clusters appearing as early as \( 10^{-4} \tau_o \). The distribution peaks near 0.1\( \tau_o \), when dynamic facilitation starts to dominate, and has a cutoff at around 10\( \tau_o \). As \( T \) decreases below the mode-coupling crossover, a power-law tail emerges at \( t/\tau_o \ll 1 \). For \( T \leq 0.07 \), the power law extends over at least six decades, with a nearly constant exponent \( \Pi(B) \approx \tau^{-0.38} \) for the 3D model. The relaxation of localized clusters at early times is extremely broadly distributed, presumably stemming from an equally broad distribution of activation energies.

Remarkably, if we plug the measured distribution of waiting times in equation (1), the power law \( \Pi(\log_{10}(\tau)) \approx \tau^{-0.38} \) directly
relaxation spectra, we analyse the simplest version of such a model traps (Fig. 4b). To provide a qualitative, generic description of the a given local relaxation event may now affect the state of the other stems from the relaxation of a sparse population of clusters characterized by trap models. To introduce dynamic facilitation as the second observations. We first imagine that the liquid can be decomposed into we construct an empirical model based on our numerical observations, with the exponent . This microscopic view of the power-law wing alone does not explain why it appears in excess of the peak well fitted by the spectrum of a stretched exponential (dashed). The high-frequency part of the underlying spectrum, unaffected by facilitation, is well described by a power law, namely, . The excess wing, thus, corresponds to the beginning of the relaxation process.

This microscopic view of the power-law wing alone does not explain why it appears in excess of the peak observed at larger times where dynamic facilitation is observed. To explain this point, we construct an empirical model based on our numerical observations. We first imagine that the liquid can be decomposed into independent domains characterized by a local relaxation time (Fig. 4a). This heterogeneous viewpoint is mathematically captured by trap models. To introduce dynamic facilitation as the second key ingredient, we construct a facilitated trap model, assuming that a given local relaxation event may now affect the state of the other traps (Fig. 4b). To provide a qualitative, generic description of the relaxation spectra, we analyse the simplest version of such a model and assume, in a mean-field spirit, that dynamic facilitation equally affects all the traps. A more local version was designed elsewhere for different purposes.

We consider traps with energy levels drawn from distribution , and assume an activated dynamics. The energy of a trap is renewed after a Poisson-distributed timescale of mean . Since deep traps take much longer to relax than shallow ones, the system is dynamically heterogeneous. Following ref. , we use , where , to smoothly interpolate between the much-studied Gaussian and exponential distributions. The dynamics at temperature leads to the equilibrium energy distribution , where and are the mean and variance of the energy distribution in the interval , using a Metropolis filter to leave the equilibrium distribution unchanged. This coupling between traps mimics dynamic facilitation. The relaxation spectra is computed either analytically or by simulating the facilitated model. The model is specified by two parameters, namely, and , for which equilibrium dynamics can be studied at any temperature . We have systematically investigated this parameter space, and find spectra with quantitative differences but generic features. In Fig. 4c, we select and , one obtains the blue spectrum (Fig. 4c), with the same high-frequency behaviour, but which extends much further at low frequencies. Indeed, without facilitation, each trap relaxes independently, and the equilibrium distribution determines the dynamic spectrum, which is broad and relatively symmetric. In the presence of facilitation, shallow traps still relax independently and are essentially unaffected. Crucially, deep traps now receive small kicks whenever a shallow trap relaxes, and their energies slowly diffuse towards the most probable value. This accelerates their relaxation, which eventually affects the tail of the relaxation time distribution. As a result, dynamic facilitation ‘compresses’ the low-frequency part of the underlying spectrum (blue), as hinted in ref. , and highlighted by the arrow in Fig. 4c. We, thus, interpret the winged, asymmetric spectrum as a broad underlying distribution of relaxation timescales (well described by a power law at early times) compressed by dynamic facilitation at long times. Ironically, in our picture, the peak itself is in ‘excess’ of a much broader underlying time distribution with a high-frequency power-law shape. In this view, the excess wing forms an integral part of the structural relaxation.

Our study frontally attacks a central question regarding the relaxation dynamics of supercooled liquids near the experimental glass transition and paves the way for many more studies of a totally unexplored territory now made accessible to modern computer studies. Further enlarging the family of available computer glass formers would also help filling the gap with more complex molecular systems studied experimentally.

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Mobility at the single-particle level. When analysing the mobility at the single-particle level, we first need a criterion to distinguish between mobile and immobile particles. In three dimensions, we have considered several mobility definitions that give quantitatively similar results. The first mobility definition is based on displacements. To remove fast dynamical processes, we use the conjugate gradient method and find the inherent structure (IS) of a configuration at time \( t \), namely, \( \{ \mathbf{r}_i^\text{g} (t) \} \). Particle \( i \) is defined as mobile at time \( t \) if \( | \mathbf{r}_i^\text{g} (t) | > d \) (ref. 48). This cutoff is between the first minimum and second maximum of the self-part of the Van Hove function \( G_s (\mathbf{r}, t) = \langle | \delta (\mathbf{r} - \mathbf{r}_i^\text{g} (t)) | \rangle \) in the time regime where \( F_s (t) \) is almost constant. This first mobility definition is, however, not convenient in two dimensions because of the collective long-ranged fluctuations that affect the translational dynamics. A second mobility definition is based on changes in the particle’s local environment. At time \( t = 0 \), we find number \( n_i \) and the identity of particle \( i \)’s neighbours, defined as particles \( j \) with \( r_{ij}^\text{g} < 1.485 \) in three dimensions (1.300 in two dimensions), corresponding to the first minimum in the rescaled pair correlation function \( g_r (r) n_i \). We define the bond-breaking correlation as the fraction of the remaining neighbours at time \( t \):

\[
C_b (t) = \left( \frac{1}{N} \frac{\sum_{i=1}^{N} C_b (t_i)}{n_i} \right)_{n_i}
\]

which is averaged over \( n_i \) independent runs.

A particle is defined as mobile at \( t \) if \( C_b (t) > 0.55 \), that is, if it has lost half of its initial neighbours. The cutoff value ensures that the set of particles identified as mobile in this way substantially overlap with that identified via the displacement criterion. We then introduce clusters of mobile particles. Two particles \( i \) and \( j \) that are mobile at time \( t \) belong to the same cluster if \( r_{ij} < 1.5 \) in three dimensions and \( r_{ij} < 1.4 \) in two dimensions, close to the first minimum of \( g_r (r) \).

Relevant temperature scales. We determine three temperature scales relevant to the glassy slowdown: the onset temperature of glassy dynamics \( T_o \), the mode-coupling crossover temperature \( T_m \), below which conventional MD simulations cannot reach equilibrium and the extrapolated experimental glass transition temperature \( T_m \). We, thus, need to extrapolate our data over five decades to locate \( T_m \) (ref. 25). We fit the high-temperature \( T_m \), data to an Arrhenius law, and identify the onset temperature \( T_o \) as the temperature below which \( \tau \) is super-Arrhenius. We note \( \tau_s = \tau (T_o) \). The mode-coupling crossover temperature \( T_m \) is obtained by fitting the data with a power law \( T_m (T - T_o) \sim \log (10) \) (ref. 8), where \( g = 2.7 \) and 2.5 for \( d = 2 \) and 3, respectively. Given that \( \log (\log (t/\tau^s)) = 4 \) at \( T_m \), this temperature delimits the regime \( T > T_m \) (where MD alone can reach equilibrium) from the regime \( T < T_m \) (where the swap algorithm is needed to perform equilibrium simulations). The experimental glass transition temperature \( T_o \) is defined by \( \log (\log (t/\tau^s)) = 12 \). In three dimensions, the longest simulation time \( t_m = 1.5 \times 10^7 \) is independent of \( T_o \), therefore, we can directly access \( \log (\log (t/\tau^s)) \leq 5 \). We, thus, need to extrapolate our data over five decades to locate \( T_o \). We increase the accuracy of extrapolation by using TTS, which is well obeyed in our model (ref. 39). In the temperature regime where correlation functions reach \( e^{-1} \), the second step of relaxation is well fitted by a stretched exponential \( \exp (-\beta | t - t_o | ^\gamma) \). The stretching exponent \( \beta \) is 0.56 in three dimensions (in two dimensions, \( \beta = 0.60 \) for \( C_2 \) and \( \beta = 0.67 \) for \( C_2 \)) is almost independent of temperature, and amplitude \( E \) slowly increases with decreasing temperature. Fixing \( \beta \), we estimate \( \tau \) at temperatures where decorrelation is sufficient to perform accurate TTS, extending our measurements over approximately two decades. We extrapolate \( \tau \) over the four remaining decades using an Arrhenius fit \( \tau (T) \propto \exp (E/T) \), where \( E = 2.67 \) in three dimensions (2.97 in two dimensions), and locate \( T_o \). Importantly, the Arrhenius extrapolation is a safe choice as it, at worst, underestimate the relaxation times.

Computation of relaxation spectra. The computation of relaxation spectra \( \gamma (\omega) \) first requires to differentiate the correlation function with respect to the logarithm of time. We use first-order finite-difference approximation. Namely, if the configurations are stored at logarithmically spaced times \( \{ \tilde{t}_k \}_{k=1,...,N} \), we have

\[
\frac{d \gamma (\omega)}{d \log \tilde{t}_k} = \frac{\gamma (\tilde{t}_k) - \gamma (\tilde{t}_{k-1})}{\log (\tilde{t}_k) - \log (\tilde{t}_{k-1})}
\]
The integral in equation (1) is then evaluated as

\[ \chi''(\omega) = -\sum_{k} \frac{dF_i(t_k)}{dt} \omega \frac{\partial}{\partial \omega} \log \left( \frac{t_k}{t_{k-1}} \right) . \]  

(9)

We use the bond-breaking correlation function \( C_g \) instead of \( F_i \) in two dimensions. In the Supplementary Information, we discuss errors that arise from computing the spectrum when \( F_i \) does not decay to zero. We also discuss issues related to statistical noise and a comparison with direct Fourier transforms.

**Trap model.** We consider traps with energy levels \( E \geq 0 \) drawn from the exponential power distribution

\[ \rho(E) = \frac{\alpha}{E_0 \Gamma(1/\alpha)} e^{-E/E_0} \gamma , \]

(10)

where \( \gamma \) is the gamma function, and take \( E_0 = 1 \) in the following. We assume that the dynamics at temperature \( T \) is thermally activated. The energy \( E \) of a trap is renewed after a Poisson-distributed timescale of mean \( \langle \tau(E) \rangle = e^{\alpha T}. \) The equilibrium energy distribution at temperature \( T \) is

\[ P_{eq}(T, E) = \frac{\rho(E) e^{\alpha T}}{Z(T)} , \]

(11)

where \( Z(T) = \int_0^\infty dE \rho(E) e^{\alpha T}. \)

We monitor the relaxation dynamics by computing the average persistence function \( p(t) \). In the absence of dynamic facilitation, the persistence can be directly computed as

\[ p(t) = \int_0^\infty dE P_{eq}(T, E) e^{-t/\langle \tau(E) \rangle} . \]

(12)

In the absence of dynamic facilitation, the average persistence is evaluated using Mathematica (NIntegrate and WorkingPrecision of 30). We then calculate the relaxation spectrum \( \chi''(\omega) \) by following the procedure described previously, replacing \( F_i(t) \) with persistence \( p(t) \). We compute persistence \( p(t) \) over a time interval large enough to observe full decorrelation, namely, \( [10^{-10}, 10^{10}] \), for \( \alpha = 1.1 \) and \( T = 0.629 \), and minimize errors in the relaxation spectrum.

**Simulations of the facilitated trap model.** We consider a system composed of \( N \) traps. We initialize the system with an equilibrium condition by sampling the traps’ energies directly from the equilibrium distribution \( P_{eq}(T, E) \). Since the cumulative probability distribution of energies \( C_{eq} \) cannot be explicitly computed, we use Mathematica to evaluate it, and to numerically construct the reciprocal function \( E = C_{eq}^{-1} \). For each of the \( N \) traps, we generate \( X \) uniformly distributed in \([0, 1]\), and assign it an energy \( E = E(X) \). This procedure generates an initial condition in equilibrium. Each trap is assigned a renewal time exponentially distributed, with mean \( e^{\alpha T} \). We initialize persistence \( p(t=0) \) of all the traps to one.

The dynamics proceeds as follows. First, we identify trap \( i \) with the smallest renewal time \( \tau_{int} \) which will relax first. We update all the other traps by subtracting \( \tau_{int} \) to their renewal time \( \tau_i \). When the trap \( i \) relaxes, its persistence is set to zero, that is, \( p_i \rightarrow 0 \), and we give it a new energy value sampled from \( \rho(E) \), as well as a new renewal time, as described above.

This relaxation event then affects all the other traps. We attempt to displace their energy by a random amount \( \delta E \) (different for each trap) uniformly distributed in \([-\delta E, \delta E] \). The scaling with \( N \) ensures that the resulting dynamics is independent of \( N \). We then accept or reject this attempt to leave the equilibrium probability distribution \( P_{eq} \) unchanged. To this end, we introduce an effective potential \( V = -T \log P_{eq} \) and compute the change in effective potential \( \delta V = T(E^{eq} - E) - \delta E \). When \( \delta V < 0 \), the change in energy is accepted; otherwise, it is accepted with probability \( \exp(-\delta V/T) \). When accepted, we pick a new renewal time exponentially distributed with average \( e^{\alpha T} \). When the move is completed, we again determine which of the traps is the next one to relax, and proceed as before.

We measure the average persistence \( p(t) = \langle \sum p_i(t)/N \rangle \), where the brackets indicate average over independent runs, and where the sum runs over all the traps. We simulate the dynamics of the model until the total persistence is equal to zero.

**Data availability**

Source data are provided with this paper. The data that support the findings of this study are available from the corresponding author upon reasonable request.

**Code availability**

The codes used in this study are available from the corresponding author upon reasonable request.

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**Author contributions**

B.G., C.S. and I.B. designed the research. B.G. and C.S. carried out the simulations. B.G., C.S. and I.B. analysed the data and wrote the paper.

**Competing interests**

The authors declare no competing interests.

**Additional information**

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