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¹ Fluctuating Diffusivity Emerges even in Binary Gas Mixtures

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Abstract

Diffusivity in some soft matter and biological systems changes with time, called the fluctuating diffusivity. In this work, we propose a novel origin for fluctuating diffusivity based on stochastic simulations of binary gas mixtures. In this system, the fraction of one component is significantly small, and the mass of the minor component molecule is different from that of the major component. The minor component exhibits fluctuating diffusivity when its mass is sufficiently smaller than that of the major component. We elucidate that this fluctuating diffusivity is caused by the time scale separation between the relaxation of the velocity direction and the speed of the minor component molecule.

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5 I. INTRODUCTION

⁶ Brownian motion is widely observed in soft matter systems, and standard Brownian mo-⁷ tion is described by a simple stochastic process known as the Wiener process[1, 2]. In this ⁸ process, the mean square displacement (MSD) increases linearly with time and is accompa-⁹ nied by a Gaussian displacement distribution[3, 4]. Although this simple Brownian motion ¹⁰ is fully understood, a new type of Brownian motion has been recently reported; although ¹¹ the MSD is proportional to time, the displacement distribution deviates from the Gaussian ¹² distribution[5–8]. This motion is known as Brownian (or Fickian) yet non-Gaussian diffu-¹³ sion and cannot be described by the simple Wiener process. This process can be successfully ¹⁴ described by the Langevin equation with the time-dependent fluctuating diffusivity[8].

$$\frac{d\mathbf{R}(t)}{dt} = \sqrt{2D(t)}\boldsymbol{\xi}(t),\tag{1}$$

¹⁵ where $\mathbf{R}(t)$ denotes the position of the Brownian particle, D(t) denotes the fluctuating ¹⁶ diffusivity, and $\boldsymbol{\xi}(t)$ is Gaussian white noise. The fluctuating diffusivity obeys a stochastic ¹⁷ process independent of $\mathbf{R}(t)$. The first and second order statistical moments of $\boldsymbol{\xi}(t)$ are ¹⁸ given as $\langle \boldsymbol{\xi}(t) \rangle = \mathbf{0}$ and $\langle \boldsymbol{\xi}(t) \boldsymbol{\xi}(t') \rangle = \mathbf{I} \delta(t-t')$, where $\langle \dots \rangle$ represents the statistical average ¹⁹ and \mathbf{I} is the unit tensor.

The origins of the fluctuating diffusivity in soft matter and biological systems can be classified into two categories[9]. The first origin is a spatially and/or temporally heterogeneous environment[10, 11]. For instance, particles in supercooled liquids (glass formers)[12–14], colloidal suspensions [7, 15, 16], biological systems [5, 6, 17, 18], and active matter [19, 20] exhibit fluctuating diffusivities, owing to their heterogeneous environments. The second origin is the fluctuation in the conformational degrees of freedom. That is, the diffusivtive can fluctuate depending on the fluctuations of the conformation or orientation of a molecule[8, 21, 22]. Examples include the center of mass of an entangled polymer[8] and rod-like particle solution[22].

Here, one question may arise: are there only two origins of fluctuating diffusivity? In this study, we demonstrate that the third origin of fluctuating diffusivity exists by investigating simple gas systems, i.e., binary gas mixtures comprising hard spheres with different masses, in which the fraction of one component is sufficiently small. These systems do not possess a heterogeneous environment nor conformational degrees of freedom, which are known to be the origins of fluctuating diffusivity. The gas molecules are assumed to be spherical and ³⁵ do not have any internal degrees of freedom. They are randomly distributed in space, and ³⁶ there is no spatial correlation. Even in such systems, the fluctuating diffusivity causing ³⁷ Brownian yet non-Gaussian diffusion emerges under specific conditions. We elucidate that ³⁸ the observed fluctuating diffusivity originates from the separation of time scales of two ³⁹ relaxation processes of the minor component; the velocity direction relaxation and speed ⁴⁰ relaxation.

41 II. SYSTEM

The dynamics of a single molecule A in another gas molecule B is investigated as a model 43 of binary gas mixtures, where the fraction of molecules of gas A is sufficiently small. The 44 molecules A and B have different masses, m_A and m_B , and sizes σ_A and σ_B , respectively. 45 The system is in equilibrium with inverse temperature β , and the number density of molecule 46 B is ρ . Molecule A moves ballistically until it collides with molecule B. Molecule A instan-47 taneously changes its velocity by collision based on the conventional hard-sphere interaction 48 [23, 24] as follows:

$$\boldsymbol{v}_{A}^{\prime} = \boldsymbol{v}_{A} - \frac{2m_{B}}{m_{B} + m_{A}} \left(\boldsymbol{v}_{A} - \boldsymbol{v}_{B} \right) \cdot \hat{\boldsymbol{r}}_{AB} \hat{\boldsymbol{r}}_{AB}.$$
(2)

⁴⁹ Here, \boldsymbol{v}_A' is the velocity of molecule A after collision, \boldsymbol{v}_A and \boldsymbol{v}_B are the velocities of molecules ⁵⁰ A and B before collision, respectively, and $\hat{\boldsymbol{r}}_{AB}$ is the unit vector connecting the centers of ⁵¹ molecules A and B. Here, it should be mentioned that this collision protocol is not crucial ⁵² for the following results; similar data will be obtained for other interaction potentials such ⁵³ as the Weeks-Chandler-Andersen potential.

In gas systems, the dynamics of a molecule can be approximately described as a Marko-⁵⁵ vian stochastic process because the dynamic correlations are weak[23, 25, 26]. Therefore, ⁵⁶ we employ the kinetic Monte Carlo (KMC) method [27, 28] to simulate the dynamics of ⁵⁷ molecule A. Collision statistics are required for implementing the KMC method. In hard-⁵⁸ sphere gas, the probability density of molecule A colliding with molecule B with v_B at \hat{r} ⁵⁹ and time interval s for a given v_A becomes

$$P(\boldsymbol{v}_{B}, \hat{\boldsymbol{r}}_{AB}, s | \boldsymbol{v}_{A})$$

$$= \rho \sigma^{2}(\boldsymbol{v}_{B} - \boldsymbol{v}_{A}) \cdot \hat{\boldsymbol{r}}_{AB} \left(\frac{\beta m_{B}}{2\pi}\right)^{3/2} \exp\left(-\frac{\beta m_{B} \boldsymbol{v}_{B}^{2}}{2}\right)$$

$$\times \exp\left[-F(\boldsymbol{v}_{A})s\right] \Theta\left[(\boldsymbol{v}_{A} - \boldsymbol{v}_{B}) \cdot \hat{\boldsymbol{r}}_{AB}\right].$$
(3)

Here, $\sigma = (\sigma_A + \sigma_B)/2$, $F(\boldsymbol{v}_A)$ is the average collision frequency of molecule A with velocity \boldsymbol{v}_A , and $\Theta(x)$ is the Heaviside step function (collision does not occur for $(\boldsymbol{v}_A - \boldsymbol{v}_B) \cdot \hat{\boldsymbol{r}}_{AB} < 0$). Here, we emphasize that Eq. (3) does not depend on the spatial position nor time; the statistics depend only on the velocity of molecule A. The explicit expression of $F(\boldsymbol{v}_A)$, derivation of Eq. (3), and numerical scheme are explained in Appendix A and B. The dynamics of molecule A can be characterized only by the mass ratio $\mu = m_A/m_B$. We employ dimensionless units by setting $m_B = 1$, $\beta^{-1} = 1$, and $1/\rho\sigma^2 = 1$.

67 III. NUMERICAL RESULTS

Figure 1 shows the MSD $\langle \Delta \mathbf{R}^2(\Delta t) \rangle$, where $\Delta \mathbf{R}(\Delta t) = \mathbf{R}(\Delta t) - \mathbf{R}(0)$ and Δt denotes the time lag. For comparison, we have included the prediction by the Enskog theory[26, 29]:

$$\left\langle \Delta \mathbf{R}^2(\Delta t) \right\rangle = \frac{3\tau_c^2}{2\mu} \left[-1 + \frac{2\Delta t}{\tau_c} + e^{-2\Delta t/\tau_c} \right],\tag{4}$$

 $_{70}$ where τ_c is the crossover time from ballistic to diffusive regions defined as follows:

$$\tau_c = \sqrt{9\mu(\mu+1)/32\pi}.$$
 (5)

⁷¹ The results obtained from the KMC simulations exhibit simple ballistic and diffusive be-⁷² haviors in the simulated μ range, and these results are almost perfectly reproduced by the ⁷³ Enskog theory. We naively expect that the dynamics of molecule A is simple Brownian ⁷⁶ motion with constant diffusivity for any μ .

⁷⁶ However, the dynamics of molecule A is not simple Brownian motion for small μ . Figure 2 ⁷⁷ shows the trajectories of molecule A for sufficiently large and small mass ratios $\mu = 10^2$ and ⁷⁸ 10^{-4} . The observation time is $T = 10^6 \tau_c$, and the trajectories are mapped onto the xy⁷⁹ plane. The colors express the magnitude of the scaled temporal displacement for a time lag ⁸⁰ $\Delta t = 10\tau_c$. For $\mu = 10^2$, the fast (red) and slow (blue) areas are homogeneously distributed; ⁸¹ this is consistent with simple Brownian motion. By contrast, for $\mu = 10^{-4}$, large clusters ⁸² of fast and slow areas are clearly observed. This implies that the dynamics of molecule A⁸⁴ results with typical mass ratios, $\mu = 10^2$ and 10^{-4} , as the representative cases of simple ⁸⁵ Brownian motion and non-trivial diffusion, respectively. Data for other mass ratios are ⁸⁶ summarized in Appendix C.



FIG. 1. Mean square displacements (MSDs) of the molecule A for several mass ratios μ . The symbols are the KMC simulation data, and the black solid curves represent the prediction by the Enskog theory (Eq. (4)).

To examine whether the dynamics of molecule A is Gaussian, we calculate the self-part of the van Hove correlation functions, which is defined as $G_s(\Delta X, \Delta t) = \langle \delta[\Delta X - (X(t + \Delta t) - X(t))] \rangle$, where X(t) is the position of molecule A in the x direction at time t. Figure 3 shows $G_s(\Delta X, \Delta t)$ for various Δt . For $\mu = 10^2$, $G_s(\Delta X, \Delta t)$ is Gaussian within the simulated Δt range. In contrast, for $\mu = 10^{-4}$, $G_s(\Delta X, \Delta t)$ deviates from the Gaussian distribution within an intermediate time lag, $10^1 \leq \Delta t/\tau_c \leq 10^4$. This deviation disappears for a sufficiently large time lag $\Delta t/\tau_c \gtrsim 10^5$. Therefore, Brownian yet non-Gaussian diffusion appears for $\mu = 10^{-4}$ at the intermediate time scale. This behavior is commonly observed for $\mu < 1$ as shown in Fig. C.1 in Appendix. The non-Gaussian behavior can be also observed in T the non-Gaussian parameter (NGP) shown in Fig. C.2 in Appendix. The NGP exhibits **90** non-negligible peaks for $\mu < 1$.

To analyze the non-Gaussian behavior in detail, we calculate the ergodicity breaking (EB) 101 parameter[8, 30] defined as follows:

$$EB(\Delta t, T) = \frac{\left\langle \left[\overline{\delta^2}(\Delta t, T) \right]^2 \right\rangle}{\left\langle \overline{\delta^2}(\Delta t, T) \right\rangle^2} - 1.$$
(6)

¹⁰² Here, $\overline{\delta^2}(\Delta t, T)$ denotes the time-averaged MSD for the time lag Δt and finite observation



FIG. 2. Typical trajectories of the molecule A during $0 \le t \le 10^6 \tau_c$ for $(a)\mu = 10^2$ and $(b)10^{-4}$ from the KMC simulation. The trajectories are mapped onto the xy plane. The colors represent the reduced temporal displacement $|\mathbf{R}(t + \Delta t) - \mathbf{R}(t)| / \sqrt{\langle \Delta \mathbf{R}^2(\Delta t) \rangle}$ with $\Delta t = 10\tau_c$.

103 time T:

$$\overline{\delta^2}(\Delta t, T) = \frac{1}{T - \Delta t} \int_0^{T - \Delta t} [\mathbf{R}(t + \Delta t) - \mathbf{R}(t)]^2 dt.$$
(7)

The dependence of the EB parameter on Δt was theoretically proven to be weak when $T \gg \Delta t$ [8]. Therefore, we set $\Delta t/\tau_c = 10$ and calculate the EB parameter as a function of T for $T/\tau_c \ge 10^2$. Figure 4 displays the observation time dependence of the EB parameter,



FIG. 3. Self-part of the van Hove correlation functions of the molecule A for different time lags Δt from the KMC simulation. (a) $\mu = 10^2$ and (b) 10^{-4} . For convenience, the displacement is normalized using the root MSD $\sqrt{\langle \Delta X^2(\Delta t) \rangle}$. The solid black curves represent the Gaussian distribution.

¹⁰⁷ which simply exhibits a decay $\text{EB} \propto T^{-1}$ in the entire T range for $\mu = 10^2$. This implies ¹⁰⁸ that the dynamics of molecule A follows a Gaussian process. In contrast, for $\mu = 10^{-4}$, ¹⁰⁹ the EB parameter exhibits a shoulder before the Gaussian decay $\text{EB} \propto T^{-1}$. This is also ¹¹⁰ observed for other sufficiently small mass ratios, $\mu \ll 1$, as shown in Fig. C.3 in Appendix. ¹¹¹ The existence of this shoulder can be attributed to the fluctuating diffusivity [8], and the ¹¹² characteristic crossover time $\tau_{\rm EB}$ from the shoulder to the EB $\propto T^{-1}$ decay can be interpreted ¹¹³ as the relaxation time of the fluctuating diffusivity[8]. The crossover time $\tau_{\rm EB}$ for $\mu = 10^{-4}$ is ¹¹⁴ estimated from the two curve fittings EB $\propto T^{-\alpha}$ where $0 < \alpha < 1$ for short T and EB $\propto T^{-1}$ ¹¹⁵ for long T regions. The obtained $\tau_{\rm EB}$ for $\mu = 10^{-4}$ is approximately equal to the time scale ¹¹⁶ at which the van Hove correlation function becomes Gaussian.



FIG. 4. Ergodicity breaking (EB) parameters corresponding to $\mu = 10^2$ and 10^{-4} from the KMC simulation. The time lag is set as $\Delta t = 10\tau_c$. The dotted lines indicate the fitting results to the power-laws EB $\propto T^{-\alpha}$ and EB $\propto T^{-1}$.

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119 IV. ORIGIN OF THE FLUCTUATING DIFFUSIVITY

When μ is sufficiently small, i.e., $\mu \ll 1$, the velocity of molecule A is significantly larger than that of molecule B, i.e., $|\boldsymbol{v}_A| \gg |\boldsymbol{v}_B|$. Under such a condition, the motion of molecule A is similar to that in a matrix of immobile obstacles such as Lorentz gases[23, 31– 34]. The speed of molecule A is nearly unchanged by a few collisions, whereas the velocity direction is randomized. Therefore, we expect that the relaxation times of the speed and velocity direction of molecule A will be considerably different if μ is small. We calculate the correlation functions corresponding to the velocity direction $C_d(\Delta t)$ and speed $C_s(\Delta t)$:

$$C_d(\Delta t) = \left\langle \frac{\mathbf{V}(\Delta t)}{|\mathbf{V}(\Delta t)|} \cdot \frac{\mathbf{V}(0)}{|\mathbf{V}(0)|} \right\rangle,\tag{8}$$

$$C_s(\Delta t) = \frac{\langle |\mathbf{V}(\Delta t)| |\mathbf{V}(0)| \rangle - \langle |\mathbf{V}| \rangle^2}{\langle |\mathbf{V}|^2 \rangle - \langle |\mathbf{V}| \rangle^2}.$$
(9)

Figure 5 displays $C_d(\Delta t)$ and $C_s(\Delta t)$ obtained from the KMC simulations. The figure clearly reveals that the relaxation of $C_s(\Delta t)$ (filled red symbols) is significantly slower than that 122 of $C_d(\Delta t)$ for $\mu = 10^{-4}$ (open red symbols). This behavior is commonly observed if μ is sufficiently small as shown in Figs. C.4 and C.5. The relaxation times of the direction τ_d and 124 speed τ_s can be estimated from $C_d(\Delta t)$ and $C_s(\Delta t)$, respectively. The estimates scaled by 125 τ_c (Eq. (5)) are summarized in Fig. C.6. For $\mu = 10^{-4}$, τ_d is found to be comparable to τ_c , whereas τ_s is much longer than τ_c . In addition, τ_s is of the same order as $\tau_{\rm EB}$, which strongly 127 implies that the relaxation of the fluctuating diffusivity in the binary gas mixtures is related to that of the speed of the molecule A. Here, it should be emphasized that such a timescale separation between the velocity direction and speed is not present without ballistic motion. Thus, the mechanism of the fluctuating diffusivity observed for purely diffusive motions in 131 some heterogeneous environments[5, 7, 17, 35] is different from that in our system.



FIG. 5. Correlation functions of the velocity direction $C_d(\Delta t)$ and speed $C_s(\Delta t)$ of molecule A (Eqs. (8) and (9)) for $\mu = 10^{-4}$ and 10^2 from the KMC simulation.

Based on the above results, we propose a possible scenario for the emergence of fluctuating 132 ¹³³ diffusivity in our binary gas mixture with $\mu \ll 1$. At the intermediate time scale $\tau_d \lesssim T \lesssim \tau_s$, molecule A diffuses because its velocity direction changes randomly. The speed of molecule 134 A remains approximately constant, $|\mathbf{v}_A(t)| \approx v_A$, and thus the diffusion coefficient can be 135 described by a function of constant as $D(t) = D(v_A)$. At the long timescale $T \gtrsim \tau_s$, D(t)136 starts to fluctuate temporarily owing to the fluctuations of $|v_A(t)|$. At the very long time 137 scale $T \gg \tau_s$, the fluctuation of the diffusivity is smeared out and the Gaussian normal 138 ¹³⁹ diffusion with the effective diffusion coefficient $D_{\text{eff}} = \langle D \rangle$ is observed. Therefore, the origin 140 of the fluctuating diffusivity in our system is the separation of the relaxation timescales of ¹⁴¹ the velocity direction and speed. This scenario also explains the clusters observed in Fig. 2; ¹⁴² they reflect the persistence of the molecule A speed within the timescale τ_s .

To validate the proposed scenario, we theoretically calculate the van-Hove correlation function of the molecule A with $\mu \ll 1$. At the intermediate timescale $\tau_d \lesssim T \lesssim \tau_s$, the dynamics of the molecule A can be virtually described as a mobile particle in dilute fixed spherical obstacles. Then the diffusion coefficient is calculated as $D(|\boldsymbol{v}_A|) = |\boldsymbol{v}_A|/3\pi$ [23]. The probability density of the displacement of the molecule A under a given speed $v_A = |\boldsymbol{v}_A|$ is Gaussian:

$$P(\Delta X; \Delta t | v_A) = \frac{1}{\sqrt{4\pi D(v_A)\Delta t}} \exp\left(-\frac{\Delta X^2}{4D(v_A)\Delta t}\right).$$
 (10)

In equilibrium, v_A obeys the Maxwell-Boltzmann distribution: $P_{\rm MB}(v_A) = 4\pi v_A^2 (2\pi)^{-3/2} \exp(-v_A^2/2)$. By taking the equilibrium average of Eq. (10) with respect to v_A , we have the van-Hove correlation function $G_s(\Delta X, \Delta t)$ at the intermediate timescale $\tau_d \leq \Delta t \leq \tau_s$:

$$G_s(\Delta X; \Delta t) = \int_0^\infty dv_A P(\Delta X; \Delta t | v_A) P_{\rm MB}(v_A).$$
(11)

¹⁵² We numerically calculate Eq. (11) and show the result in Fig. 6. The theoretical prediction ¹⁵³ by Eq. (11) reasonably agrees with the KMC simulation result. This result supports our ¹⁵⁴ scenario on the fluctuating diffusivity; the fluctuating diffusivity in our system originates ¹⁵⁵ from the separation of the relaxation timescales between the velocity direction and the speed. ¹⁵⁶ The tail of $G_s(\Delta X; \Delta t)$ from the Gaussian distribution has been observed in several systems. ¹⁵⁷ The tail in Eq. (11) can be approximately calculated using the saddle point method:

$$G_s(\Delta X; \Delta t) = \sqrt{\frac{3}{4\pi}} \frac{|\Delta X|}{\Delta t} \exp\left[-3\left(\frac{3\Delta X^2}{8\sqrt{2}\Delta t}\right)^{\frac{2}{3}}\right] \qquad \text{(for } \Delta X \gg 1\text{)}.$$
 (12)

¹⁵⁸ Thus we find that the tail is not the exponential nor the stretched Gaussian distributions, ¹⁵⁹ which are often observed in glass-forming liquids[12, 13, 36, 37] or some biological systems[6, ¹⁶⁰ 10, 17, 19, 20].

161 V. RELATION TO OTHER SYSTEMS

The motion of molecule A with $\mu \ll 1$ can be considered to be similar to that in the Lorenz gas model[31], which has been widely investigated as a dynamic model for light gas molecules in spatially fixed obstacles [23, 32–34]. In the Lorentz gas model, only the velocity direction changes and the speed remains unchanged at any timescale. Thus, the mechanism



FIG. 6. Theoretical prediction of the scaled self-part of the van-Hove correlation function $G_s(\Delta X, \Delta t)$ of the molecule A (ochre curve). For comparison, the KMC simulation results with $\mu = 10^{-4}$ for different time lags (Fig. 3(b)) are shown with symbols and the Gaussian distribution is displayed with the black curve.

that causes the fluctuating diffusivity observed in our system cannot be realized in Lorentzgas systems.

177 VI. CONCLUSION

¹⁷⁸ In this study, we identified a novel origin of fluctuating diffusivity, which is neither en-¹⁷⁹ vironmental heterogeneity nor conformational degrees of freedom. Fluctuating diffusivity ¹⁸⁰ emerges in simple binary gas mixtures with mass and fraction contrasts when the mass ¹⁸¹ of the minor component molecule is sufficiently small in comparison to that of the major ¹⁸² component. We showed that fluctuating diffusivity originates from the timescale separation ¹⁸³ between the relaxation times of the velocity direction and the speed of the minor component ¹⁸⁴ molecule. Our findings open a new modeling path for fluctuating diffusivity. They will also ¹⁸⁵ shed light on the kinetic behavior of gas systems from a new aspect. We hope that the ¹⁸⁶ predicted non-Gaussian behavior and fluctuating diffusivity will be experimentally observed ¹⁸⁷ in the future.

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192 Appendix A: Collision statistics

The kinetic Monte Carlo (KMC) method requires collision statistics as inputs. In the present case, the probability density of a collision for molecule A is required, which can be derived based on the gas kinetic theory[23, 39, 40]. We employ the following assumptions on our system:

- 197 1. The dynamics of molecule A obeys a Markovian stochastic process.
- 198 2. Molecule B is homogeneously distributed in space.

¹⁹⁹ From assumptions 1 and 2, the collision rate at which molecule A with velocity v_A collides ²⁰⁰ with molecule B with velocity v_B can be expressed as follows:

$$\rho\sigma^2(\boldsymbol{v}_A - \boldsymbol{v}_B) \cdot \hat{\boldsymbol{r}}_{AB} \Theta[(\boldsymbol{v}_A - \boldsymbol{v}_B) \cdot \hat{\boldsymbol{r}}_{AB}], \qquad (A1)$$

where $\Theta(x)$ denotes the Heaviside step function (no collision occurs when $(\boldsymbol{v}_A - \boldsymbol{v}_B) \cdot \hat{\boldsymbol{r}}_{AB} < 0$). The required probability density for collision $P(\boldsymbol{v}_B, \hat{\boldsymbol{r}}_{AB}, s | \boldsymbol{v}_A)$ can be decomposed into the product of three factors. The first factor is the cumulative waiting-time distribution of molecule A with velocity \boldsymbol{v}_A . Owing to the Markovian nature of the dynamics, this factor becomes an exponential distribution. The second factor is the probability density of the velocity of the colliding molecule B, which is the Maxwell-Boltzmann velocity distribution. ²⁰⁷ The third factor is the collision rate, given by Eq. (A1). For the calculation of the first ²⁰⁸ factor, the probability density of the waiting time s is required:

$$P(s|\boldsymbol{v}_A) = F(\boldsymbol{v}_A)e^{-F(\boldsymbol{v}_A)s}.$$
(A2)

209 Here, $F(\boldsymbol{v}_A)$ is the average collision frequency and is expressed as follows:

$$F(\boldsymbol{v}_{A}) = \int d\boldsymbol{v}_{B} d\hat{\boldsymbol{r}}_{AB} \rho \sigma^{2}(\boldsymbol{v}_{B} - \boldsymbol{v}_{A}) \cdot \hat{\boldsymbol{r}}_{AB} \Theta((\boldsymbol{v}_{B} - \boldsymbol{v}_{A}) \cdot \hat{\boldsymbol{r}}_{AB}) P_{\text{MB}}(\boldsymbol{v}_{B}; m_{B})$$

$$= \frac{\rho \pi \sigma^{2}}{\sqrt{\alpha}} \left[\left(\sqrt{\alpha} |\boldsymbol{v}_{A}| + \frac{1}{2\sqrt{\alpha} |\boldsymbol{v}_{A}|} \right) \operatorname{erf}(\sqrt{\alpha} |\boldsymbol{v}_{A}|) + \frac{1}{\sqrt{\pi}} \exp\left(-\alpha |\boldsymbol{v}_{A}|^{2}\right) \right],$$
(A3)

²¹⁰ where $\alpha = \beta m_B/2$. The first factor is the probability of no collisions occurring during time ²¹¹ s, which is calculated as follows [41]:

$$\Psi(s|\boldsymbol{v}_A) = \int_s^\infty ds' P(s'|\boldsymbol{v}_A) = e^{-F(\boldsymbol{v}_A)s}.$$
 (A4)

²¹² The second factor is simply expressed as

$$P_{\rm MB}(\boldsymbol{v}_B; m_B) = \left(\frac{\beta m_B}{2\pi}\right)^{3/2} \exp\left(-\frac{\beta m_B \boldsymbol{v}_B^2}{2}\right). \tag{A5}$$

²¹³ The probability density $P(\boldsymbol{v}_B, \hat{\boldsymbol{r}}_{AB}, s | \boldsymbol{v}_A)$ can be expressed as

$$P(\boldsymbol{v}_B, \hat{\boldsymbol{r}}_{AB}, s | \boldsymbol{v}_A) = \Psi(s | \boldsymbol{v}_A) P_{\text{MB}}(\boldsymbol{v}_B; m_B) \rho \sigma^2 (\boldsymbol{v}_A - \boldsymbol{v}_B) \cdot \hat{\boldsymbol{r}}_{AB} \Theta[(\boldsymbol{v}_A - \boldsymbol{v}_B) \cdot \hat{\boldsymbol{r}}_{AB}].$$
(A6)

²¹⁴ Equations (A6), (A4), and (A5), give Eq. (3) in the main text.

215 Appendix B: Numerical scheme for KMC simulation

The collision-based dynamics of molecule A can be simulated using the KMC method[27, 217 28] with Eq. (A6) as the input. The initial velocity of molecule A is sampled based on 218 Maxwell-Boltzmann distribution. The probability density of the initial velocity represented 219 in dimensionless units is

$$P_{\rm MB}(\boldsymbol{v}_A;\boldsymbol{\mu}) = \left(\frac{\boldsymbol{\mu}}{2\pi}\right)^{3/2} \exp\left(-\frac{\boldsymbol{\mu}\boldsymbol{v}_A^2}{2}\right). \tag{B1}$$

²²⁰ where μ is the mass ratio m_A/m_B , the same as in the main text. Since Eq. (B1) is a Gaussian ²²¹ distribution, \boldsymbol{v}_A can be sampled using the Box-Muller method[?].

For the time evolution of the system, sampling of the stochastic variables v_B , \hat{r}_{AB} , and s_{223} are required. However, the simultaneous sampling of these variables is technically difficult.

²²⁴ Therefore, we decompose the probability density $P(\boldsymbol{v}_B, \hat{\boldsymbol{r}}_{AB}, s | \boldsymbol{v}_A)$ into several conditional ²²⁵ probability densities as follows:

$$P(\boldsymbol{v}_B, \hat{\boldsymbol{r}}_{AB}, s | \boldsymbol{v}_A) = P(\hat{\boldsymbol{r}}_{AB} | \boldsymbol{v}_B, s, \boldsymbol{v}_A) P(\boldsymbol{v}_B | s, \boldsymbol{v}_A) P(s | \boldsymbol{v}_A),$$
(B2)

²²⁶ where $P(\hat{\boldsymbol{r}}_{AB}|\boldsymbol{v}_B,s,\boldsymbol{v}_A)$, $P(\boldsymbol{v}_B|s,\boldsymbol{v}_A)$, and $P(s|\boldsymbol{v}_A)$ are defined as follows

$$P(s|\boldsymbol{v}_A) = \int d\boldsymbol{v}_B d\hat{\boldsymbol{r}}_{AB} P(\boldsymbol{v}_B, \hat{\boldsymbol{r}}_{AB}, s|\boldsymbol{v}_A) = F(\boldsymbol{v}_A)e^{-F(\boldsymbol{v}_A)s},$$
(B3)

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$$P(\boldsymbol{v}_B|s, \boldsymbol{v}_A) = \int d\hat{\boldsymbol{r}}_{AB} \, \frac{P(\boldsymbol{v}_B, \hat{\boldsymbol{r}}_{AB}, s|\boldsymbol{v}_A)}{P(s|\boldsymbol{v}_A)} = \frac{\rho \pi \sigma^2 |\boldsymbol{v}_A - \boldsymbol{v}_B| P_{\rm MB}(\boldsymbol{v}_B; m_B)}{F(\boldsymbol{v}_A)}, \qquad (B4)$$

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$$P(\hat{\boldsymbol{r}}_{AB}|\boldsymbol{v}_B,s,\boldsymbol{v}_A) = \frac{P(\boldsymbol{v}_B,\hat{\boldsymbol{r}}_{AB},s|\boldsymbol{v}_A)}{P(\boldsymbol{v}_B|s,\boldsymbol{v}_A)P(s|\boldsymbol{v}_A)} = \frac{1}{\pi} \frac{\boldsymbol{v}_A - \boldsymbol{v}_B}{|\boldsymbol{v}_A - \boldsymbol{v}_B|} \cdot \hat{\boldsymbol{r}}_{AB} \Theta[(\boldsymbol{v}_A - \boldsymbol{v}_B) \cdot \hat{\boldsymbol{r}}_{AB}].$$
(B5)

229 $F(\boldsymbol{v}_A)$ in dimensionless units becomes

$$F(\boldsymbol{v}_A) = \pi \left(|\boldsymbol{v}_A| + 1/|\boldsymbol{v}_A| \right) \operatorname{erf}\left(|\boldsymbol{v}_A|/\sqrt{2} \right) + \sqrt{2\pi} \exp\left(-|\boldsymbol{v}_A|^2/2\right).$$
(B6)

²³⁰ Based on these decomposed probability densities, s, v_B , and \hat{r}_{AB} can be sampled sequen-²³¹ tially. s can be sampled using the inversion method[?] with Eqs. (B3) and (B6), respec-²³² tively.

Equation (B4) can be rewritten with the relative velocity, $v_r = v_B - v_A$. Without loss and generality, the relative velocity can be expressed by spherical coordinates according to $v_r = v_r \cos \phi \sin \theta e_x + v_r \sin \phi \sin \theta e_y + v_r \cos \theta e_z$. Here, e_x, e_y and e_z are orthonormal basis vectors and e_z is set to $e_z = v_A/|v_A|$. Subsequently, Eq. (B4) is reduced to

$$P(v_r, \theta, \phi | s, \boldsymbol{v}_A) = \frac{1}{4(2\pi)^{3/2} F(\boldsymbol{v}_A)} v_r^3 \sin \theta \exp\left[-(v_r^2/2 + |\boldsymbol{v}_A|^2/2 + |\boldsymbol{v}_A|v_r \cos \theta)\right].$$
(B7)

²³⁷ Because ϕ is not included in Eq. (B7), ϕ can be sampled from the uniform distribution. The ²³⁸ conditional probability density of v_r is obtained by integrating Eq. (B7) over θ and ϕ as ²³⁹ follows:

$$P(v_r|s, \boldsymbol{v}_A) = \int d\theta d\phi P(v_r, \theta, \phi|s, \boldsymbol{v}_A)$$

$$= \left[\frac{\pi}{(2\pi)^{3/2} |\boldsymbol{v}_A| F(\boldsymbol{v}_A)} \exp\left(-\frac{|\boldsymbol{v}_A|^2}{2}\right)\right] v_r^2 \exp\left(-\frac{v_r^2}{2}\right) \sinh\left(|\boldsymbol{v}_A| v_r\right).$$
(B8)

²⁴⁰ v_r can be sampled using the rejection method[?] with Eq. (B8). The conditional probability ²⁴¹ density of θ is:

$$P(\theta|v_r, \phi, s, \boldsymbol{v}_A) = \int d\phi \, \frac{P(v_r, \theta, \phi|s, \boldsymbol{v}_A)}{P(v_r|\tau, \boldsymbol{v}_A)} = \left[\frac{|\boldsymbol{v}_A|v_r}{2\sinh(|\boldsymbol{v}_A|v_r)}\right] \sin\theta \exp\left(-v_r|\boldsymbol{v}_A|\cos\theta\right).$$
(B9)

²⁴² Subsequently, θ can be sampled using the inversion method. v_B is obtained from sampled ²⁴³ v_r , θ , and ϕ .

In a similar manner, Eq. (B5) can be simplified using spherical coordinates. Without loss of generality, $\hat{\boldsymbol{r}}_{AB}$ can be expressed as $\hat{\boldsymbol{r}}_{AB} = \cos \phi' \sin \theta' \boldsymbol{e}_{x'} + \sin \phi' \sin \theta' \boldsymbol{e}_{y'} + \cos \theta' \boldsymbol{e}_{z'}$. Here, $\boldsymbol{e}_{x'}, \boldsymbol{e}_{y'}, \text{ and } \boldsymbol{e}_{z'}$ are orthogonal basis vectors and $\boldsymbol{e}_{z'}$ is set to $\boldsymbol{e}_{z'} = -\boldsymbol{v}_r/|\boldsymbol{v}_r|$. Subsequently, 247 Eq. (B5) can be expressed as:

$$P(\theta', \phi' | \boldsymbol{v}_B, s, \boldsymbol{v}_A) = \frac{1}{\pi} \cos \theta' \sin \theta' \Theta(\cos \theta').$$
(B10)

²⁴⁸ Equation (B10) does not depend on ϕ' . Therefore, ϕ' can be sampled from a uniform ²⁴⁹ distribution, and θ' can be sampled using the inversion method with Eq. (B10). \hat{r}_{AB} can be ²⁵⁰ constructed from θ' and ϕ' .

251 Appendix C: Additional Simulation Data

In the main text, we showed the representative simulation data only with mass ratios $\mu =$ ²⁵³ 10⁻⁴ and 10². In this Appendix, we show the results with different mass ratios $10^{-4} \leq \mu \leq$ ²⁵⁴ 10². The self-part of the van-Hove correlation functions of the molecule A with $\mu = (a)10^{0}$, ²⁵⁶ (b)10⁻¹, (c)10⁻², and (d)10⁻³ are displayed in Fig. C.1. The non-Gaussian parameters ²⁵⁸ against time lag with various μ are shown in Fig. C.2. Fig. C.3 displays the EB parameters ²⁶⁹ with various μ . Figs. C.4 and C.5 show the time-correlation functions of the direction and ²⁶¹ the speed of the molecule A. From the data in Figs. C.3-C.5, we estimate the characteristic ²⁶² timescales for EB, direction, and speed. The characteristic timescale for EB can be estimated ²⁶³ as the crossover time, as explained in the main text. The characteristic time scales for the ²⁶⁴ direction and time are estimated as

$$\tau_{\gamma} = \frac{\int_{0}^{\infty} d\Delta t \,\Delta t C_{\gamma}(\Delta t)}{\int_{0}^{\infty} d\Delta t \, C_{\gamma}(\Delta t)},\tag{C1}$$

265 with $\gamma = d, s$. These estimates are displayed in Fig. C.6.



FIG. C.1. The self-part of the van Hove correlation function of molecule A for (a) $\mu = 10^{0}$, (b) 10^{-1} , (c) 10^{-2} , and (d) 10^{-3} from the KMC simulation. The displacement is normalized using the root mean square displacement $\sqrt{\langle \Delta X^{2}(\Delta t) \rangle}$. The solid curves represent the Gaussian distribution.



FIG. C.2. The non-Gaussian parameter of the molecule A defined as $3\langle\Delta \mathbf{R}^4(\Delta t)\rangle/5\langle\Delta \mathbf{R}^2(\Delta t)\rangle^2 - 1$ with various mass ratios from the KMC simulation.



FIG. C.3. The ergodicity breaking (EB) parameter of molecule A for various μ from the KMC simulation. The dotted lines represent the curve fittings according to the power laws EB $\propto T^{-\alpha}$ and EB $\propto T^{-1}$ at the short and long-time regions. The shoulder and the crossover behavior can be observed only for $\mu = 10^{-3}$ and 10^{-4} .



FIG. C.4. Direction correlation function of molecule A, $C_d(\Delta t)$, for various mass ratios μ , from the KMC simulation.



FIG. C.5. The speed correlation function of the molecule A, $C_s(\Delta t)$, for various mass ratios μ , from the KMC simulation.



FIG. C.6. The direction and speed relaxation times τ_d and τ_s , and the crossover time $\tau_{\rm EB}$ from the KMC simulation data in Figs. C.3-C.5. $\tau_{\rm EB}$ is estimated only for $\mu = 10^3$ and 10^4 .

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