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Low Dose X-Ray Speckle Visibility Spectroscopy Reveals Nanoscale Dynamics in Radiation Sensitive Ionic Liquids

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X-ray radiation damage provides a serious bottleneck for investigating microsecond to second dynamics on nanometer length scales employing x-ray photon correlation spectroscopy. This limitation hinders the investigation of real time dynamics in most soft matter and biological materials which can tolerate only x-ray doses of kGy and below. Here, we show that this bottleneck can be overcome by low dose x-ray speckle visibility spectroscopy. Employing x-ray doses of 22–438 kGy and analyzing the sparse speckle pattern of count rates as low as 6.7×10^{-3} per pixel, we follow the slow nanoscale dynamics of an ionic liquid (IL) at the glass transition. At the prepeak of nanoscale order in the IL, we observe complex dynamics upon approaching the glass transition temperature T_G with a freezing in of the alpha relaxation and a multitude of millisecond local relaxations existing well below T_G . We identify this fast relaxation as being responsible for the increasing development of nanoscale order observed in ILs at temperatures below T_G .

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Introduction.—Soft matter, such as, for example, supercooled liquids, can display relatively slow dynamics on atomic or nanometer length scales. Measuring nanoscale dynamics on timescales of seconds to milliseconds constitutes a considerable experimental challenge. It can be addressed directly by x-ray photon correlation spectroscopy (XPCS) experiments employing coherent x-ray beams [1–7] by tracing fluctuations in x-ray speckle patterns. However, the highly intense x-ray beams of third-generation storage rings can also be the cause of considerable radiation damage to the samples. Atomic-scale XPCS experiments use x-ray doses of MGy to GGy, which can lead to beam-induced dynamics even in hard condensed matter samples [8]. Soft and biological matter samples are much more sensitive to radiation damage, rendering XPCS experiments with MGy x-ray doses impossible. Overcoming this bottleneck of radiation damage is even more important for the upcoming diffraction limited storage rings (DLSRs), which provide an increase in x-ray brilliance of up to 2 orders of magnitude [2,9,10]. The remedy for beam damage is frequent sample replacement as can be realized using, e.g., flow cells [11] or changing the exposure spot on the sample. However, low dose XPCS experiments often show very noisy correlation functions which do not yield conclusive insights into the dynamics or did not extend beyond feasibility studies [12,13].

Here, we demonstrate that x-ray speckle visibility experiments with nanometer spatial resolution can be performed with very low noise levels at radiation doses as low as a few

kGy. The concept consists in spreading the dose needed to obtain a correlation function over the entire sample volume by measuring at every spot on the sample the visibility of a speckle pattern as a function of the exposure time. Mitigating the absorbed dose in this way is done at the expense of signal strength; the collected speckle patterns are sparse with only 10^{-2} photons per pixel and possibly even less. We show that the speckle visibility correlation function can nevertheless be extracted by the proper assignment of photon probabilities using a sufficiently large number of images.

Ionic liquids (ILs) display a structural heterogeneity on the nanoscale [14,15] evidenced by a characteristic prepeak in diffraction experiments at Q positions around $Q = 2\text{--}3 \text{ nm}^{-1}$. The intensity of this prepeak increases with a decreasing temperature with the intriguing finding that it does not saturate at T_G , but instead this nanoscale correlation is found to increase even at temperatures well below T_G [16]. Neutron spin echo (NSE) experiments revealed that this prepeak is accompanied by complex heterogeneous dynamics with multiple relaxation channels in the picosecond to nanosecond regime at temperatures well above T_G [16]. However, the dynamics at and below the glass transition is occurring on timescales too slow for NSE experiments and has thus so far not been investigated. Applying our scheme, we are able to investigate the nanoscale dynamics of an imidazolium-based ionic liquid (C8mimCl, C₁₂H₂₃ClN₂) around its glass transition temperature.

The nanoscale order quickly decays under the intense x-ray radiation, which requires one to optimize the signal to noise ratio (SNR) in such experiments. In XPCS, an isochronous sequence of N speckle patterns is recorded at the same spot on the sample and used to calculate the intensity autocorrelation function for every pixel according to

$$g_2(i) = \frac{1}{N-i} \sum_{n=1}^{N-i} I(n)I(n+i) / \langle I \rangle^2 = 1 + \beta(Q) |f(Q, \tau)|^2. \quad (1)$$

Here i denotes the temporal separation between two frames of the recorded time series, $f(Q, \tau)$ is the intermediate scattering function (ISF), and $\beta(Q)$ is the (Q -dependent) x-ray speckle contrast. The maximum available scattering intensity k_{\max} per pixel before the damage threshold is reached is thus distributed onto N images of intensity k_{\max}/N each. The SNR in XPCS experiments is [17] [see details in Supplemental Material [18], Eq. (24)]

$$\text{SNR} \sim \frac{\beta k_{\max}}{N} \frac{\sqrt{N_{\text{pairs}} N_{\text{pix}} N_{\text{rep}}}}{\sqrt{2(\beta + 1)}}, \quad (2)$$

where N_{pairs} denotes the number of image pairs available for correlating at this specific time delay, N_{pix} the number of pixels, and N_{rep} the number of repeats of this series. Maximizing the SNR requires one to choose N as small as possible while still being able to cover the full time interval $(N-1)\tau$ of interest. Assuming typical values of $N_{\text{pix}} \sim 10^6$, $k_{\max} = 10^{-2}$, $N = 100$, and $\beta = 0.1$, we find that the SNR is on the order of one, even for $N_{\text{rep}} \sim 1000$. Because experiments usually require the systematic change of additional parameters such as the temperature or sample composition, increasing N_{rep} by orders of magnitude one quickly runs into practical problems.

The SNR can be maximized by utilizing the full available scattering intensity k_{\max} for retrieving dynamic information from just a single image by x-ray speckle visibility spectroscopy (XSVS). In this case, single images are taken with different exposure times t_e at different sample positions. The XSVS correlation function is then determined by analyzing the contrast of the smeared out speckle patterns as a function of exposure time t_e given by [19]

$$\beta(Q, t_e) = \frac{\beta_0}{t_e} \int_0^{t_e} 2 \left(1 - \frac{\tau}{t_e} \right) |f(Q, \tau)|^2 d\tau, \quad (3)$$

which yields access to the temporally averaged ISF. Reference [19] shows examples for retrieving the ISF from Eq. (3). The SNR of XSVS is [20]

$$\text{SNR} = \beta k_{\max} \sqrt{\frac{N_{\text{pix}} N_{\text{rep}}}{2(\beta + 1)}} \quad (4)$$

[Supplemental Material [18], Eq. (18)], which can be considerably larger than the one of XPCS, depending on the choice of the number of images and delay times. XSVS also allows easy scanning of a nonlinear time window, as it depends only on the adjustable exposure time. While the temporal averaging of the ISF represents a loss of information compared to XPCS, we point out that for low dose experiments XSVS may be the only way to obtain any dynamic information at all.

Theory of low intensity x-ray speckle visibility spectroscopy.—Working with low radiation doses result in speckle patterns of very low intensities on the order of 10^{-2} photons per pixel and below. The speckle contrast from such sparse images can be retrieved by analyzing the photon statistics. The probability of detecting k photons during an exposure time t_e is given by the negative binominal distribution [21]

$$P(k, t_e) = \frac{\Gamma(k + M(t_e))}{\Gamma(M(t_e))\Gamma(k + 1)} \left(1 + \frac{M(t_e)}{\bar{k}} \right)^{-k} \times \left(1 + \frac{\bar{k}}{M(t_e)} \right)^{-M(t_e)}, \quad (5)$$

where \bar{k} is the average intensity per pixel and $M(t_e)$ is the number of modes in a single speckle pattern which is connected to the speckle contrast via $\beta(t_e) = 1/M(t_e)$. Its value can be determined, for instance, from the ratio of zero- vs one-photon events $R_{0,1}(t_e)$ (see Supplemental Material [18])

$$\beta(t_e) = P(0, t_e)/P(1, t_e) - 1/\bar{k} = R_{0,1}(t_e) - 1/\bar{k}. \quad (6)$$

Experiment.—The experiment has been performed at the P10 coherence beam line at PETRA III at DESY. A photon energy of 13 keV has been used with a beam size of $4 \times 3 \mu\text{m}^2$ FWHM. The estimated flux of the partially coherent beam is 5×10^{10} photons/s. A single 50 ms exposure yielded on average an intensity of 6.7×10^{-3} photons per pixel. Temperatures of $T = 190\text{--}235$ K have been achieved by a liquid nitrogen cryostat. Photons have been detected by the EIGER 4M detector [22] (pixel size of $75 \times 75 \mu\text{m}^2$) of which 8.08×10^5 pixels have been used for the dynamic analysis. To reduce beam damage, we systematically scanned the sample through the x-ray beam by steps of $10 \mu\text{m}$, taking a single exposure for each exposure time t_e and sample position. For every temperature, we have measured ten exposure times varying between 50 and 1000 ms. For every exposure time, 1000 speckle patterns have been measured.

We used the ionic liquid 1-octyl-3-methylimidazolium chloride ($\text{C}_{12}\text{H}_{23}\text{ClN}_2$) with $T_G = 214$ K [23]. The mass

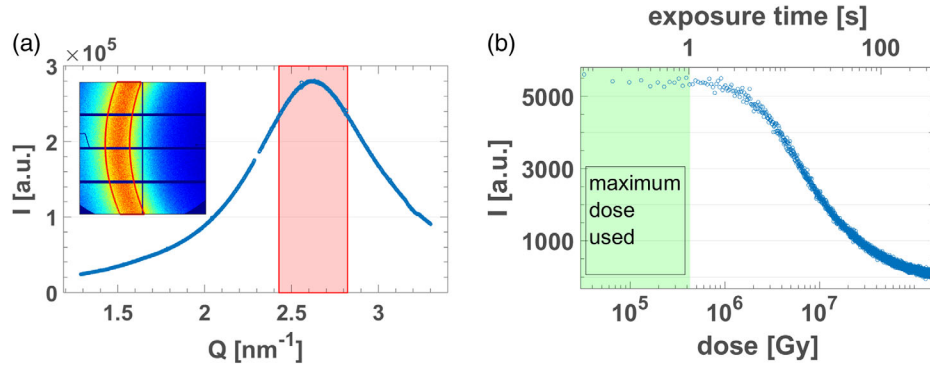


FIG. 1. (a) X-ray diffraction pattern from the supercooled ionic liquid C8mimCl associated with the nanoscale order. The inset shows the diffraction pattern as recorded on the area detector. The red box indicates the Q range used for the XSVS analysis. (b) The integrated x-ray signal from the area of the red box in (a) plotted as a function of the absorbed x-ray dose or exposure time. The green area indicates the range of doses used in the experiment.

absorption coefficient of the IL sample at 13 keV photon energy is $5 \text{ cm}^2/\text{g}$. The sample was filled into 1 mm diameter quartz glass capillaries ($100 \mu\text{m}$ wall thickness). The maximum speckle contrast in our experimental configuration is 0.1.

Results.—Averaging 1000 diffraction patterns with 1 s exposure each yields the well-known prepeak in the structure factor of ionic liquids with the maximum of the correlation located at $Q = 2.6 \text{ nm}^{-1}$ [Fig. 1(a)]. In crystallography, the intensities and widths of Bragg peaks are sensitive markers of the loss of order due to beam damage, helping to monitor even tiny radiation-induced structural changes [24]. The situation is different for disordered samples such as liquids and glasses with no long-range order present, providing only broad diffraction features. Radiation damage manifests itself by more or less abrupt changes to the scattering intensities as, for example, observed for SAXS experiments from protein solutions [25,26].

For determining the radiation damage threshold, we continuously exposed the sample on a fixed spot and acquired frames with an exposure time of 75 ms at a temperature of $T = 205 \text{ K}$, i.e., below the glass transition temperature. Using the estimated x-ray flux and calculated absorption properties of the ionic liquid, we can convert the exposure time into the corresponding x-ray dose. To calculate the dose, we used the estimated flux of 5×10^{10} photons per second on the sample. For the dose calculation, we assume that the total number of absorbed photons spreads over a volume of $7.6 \times 10.2 \times 800 \mu\text{m}^3$, which corresponds to 3σ of the Gaussian beam profile. Integrating the scattering signal over the Q range indicated in Fig. 1(a) yields the scattering power as a function of the absorbed x-ray dose [Fig. 1(b)]. A loss in scattered intensity of the nanoscale structure is detected starting at a dose of 2 MGy. The dose at which the intensity decreases by a factor of 2 is 7.15 MGy at a real space resolution of $2\pi/Q \sim 2.4 \text{ nm}$.

In protein crystallography, the maximum tolerable dose depends on the resolution aimed for and is typically around

$10 \text{ MGy}/\text{\AA}$ for cryogenically cooled protein crystals [24]. In contrast, for our IL the ratio of dose and resolution is more than one order of magnitude smaller with $0.3 \text{ MGy}/\text{\AA}$ pointing to the radiation sensitivity of the IL. In the dynamic experiment, we used doses ranging from 22 kGy to a maximum value of 438 kGy. However, dynamic properties may be affected at doses below the threshold of structural damage [8]. Experiments with a comparable dose have shown that the energy spreads over a large volume of the sample, and the expected heating is of the order of 10 K [27].

The dynamics of the sample can be inferred from the change of the photon probabilities as a function of exposure time t_e . There are two contributions to the change of photon probabilities in x-ray speckle visibility: The first one is the trivial scaling of the average intensity per pixel $\bar{k} = rt_e$ with r denoting the incoming photon rate. The second factor is the change in mode number $M(t_e)$, which is associated with the dynamics of the sample. In Fig. 2 we plot the difference between the measured zero- (blue) and one-photon probabilities (red) and the calculated ones assuming only a simple scaling of the average intensity ($T = 200 \text{ K}$). Both curves in Fig. 2 deviate from zero and change with t_e , indicating that the number of modes $M(t_e)$ is changing during the exposure. Specifically, we note that the number of detected zero-photon events is decreasing with an increasing exposure time and the number of one-photon events is increasing, in agreement with an increasing mode number due to the dynamics of the sample. We observe a similar behavior for the two- and three-photon events.

To determine the correlation function, we calculate the ratio of the zero- and one-photon events and the averaged photon probability \bar{k} for all speckle images using the scattering signal within the area marked by the red lines in Fig. 1. From these values, we can obtain the speckle contrast β using Eq. (6). This contrast value is calculated for the 1000 recorded images for each exposure time t_e and for each temperature T . The error bar is determined from the variance of the 1000 contrast values. Two examples for the

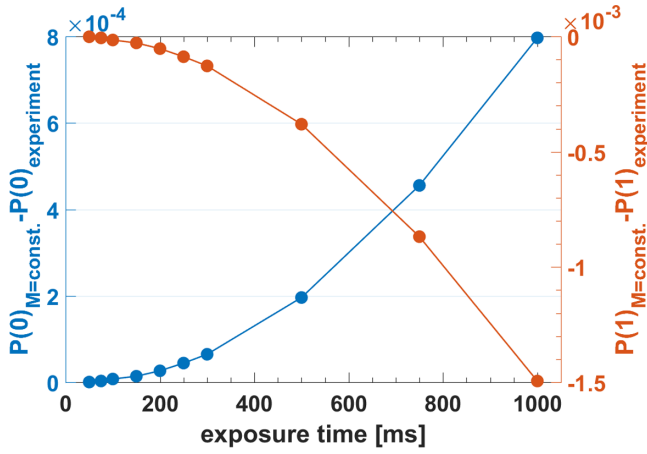


FIG. 2. Blue data: The difference between the actual measured zero-photon probability and the zero-photon probability calculated assuming no dynamics but a simple increase of the mean count rate according to $k = rt_e$. The decreasing number of zero-photon events indicates the decreasing speckle contrast as a function of the exposure time caused by the sample dynamics. Red data: The same plot as in blue but here for the one-photon probabilities.

contrast β calculated from the ratio of zero- and one-photon events for temperatures of $T = 220$ and 200 K are shown in Fig. 3. Both contrasts drop as a function of the exposure time, with the 200 K data providing a much slower loss of speckle visibility, indicating slower dynamics at lower temperatures. The correlation function at $T = 220$ K approaches a contrast value of zero at large values of the exposure time, indicating that the speckle patterns are fully decorrelated at exposure times beyond 1 s. At lower temperatures, however, the dynamics is partially frozen on the timescales of seconds and beyond, as the correlation functions do not approach the value of zero. It is also apparent that there are several relaxation mechanisms involved in the low temperature dynamics of the ionic

liquid. At $T = 200$ K, for example, in the glassy phase, we observe a fast relaxation with timescales below 100 ms, a second relaxation with timescales of 400 ms, and a frozen-in component. We label the three observed processes by $P1$, $P2$, and $P3$, and the solid lines shown in Fig. 3 represent fits using an exponential function and a fixed Kohlrausch-Williams-Watts stretching exponent of 0.7 for all three processes [28]. Unfortunately, the limited time window prevents a self-consistent refinement of the three different processes over the whole temperature range. We note, however, that multiple relaxation mechanisms have also been observed in the picosecond to nanosecond regime in the liquid phase at temperatures well above T_G [16].

The complete picture of nanoscale dynamics at the glass transition can be inferred from Fig. 4(a), showing the speckle contrast over the whole measured temperature interval around T_G . The appearance of a frozen-in component is apparent for timescales of 1000 ms and at temperatures of 210 K and below [see also the contour plot in Fig. 4(b)]. This is where the system falls out of equilibrium and the slowest relaxation gets arrested. This relaxation is attributed to an overall (α -type) relaxation of the nanoscale order in the IL [29]. However, we still observe dynamics on timescales of 100 – 200 ms in the glassy phase. As this motion does not fully decorrelate the speckle contrast, it must be local in nature. However, the fact that it still contributes a rather large component to the decorrelation of the dynamic structure factor at T_G indicates that the associated motion has a considerable spatial extension. The microscopic origin of this mode is not clear, but one can relate it to the delicate balance of electrostatic and van der Waals interactions between the ions and alkyl chains in the IL. This still allows for relaxation mechanisms on the nanoscale which do not freeze out completely below T_G . It is tempting to associate the observed increasing nanoscale order below T_G with the presence of this fast localized motion [29].

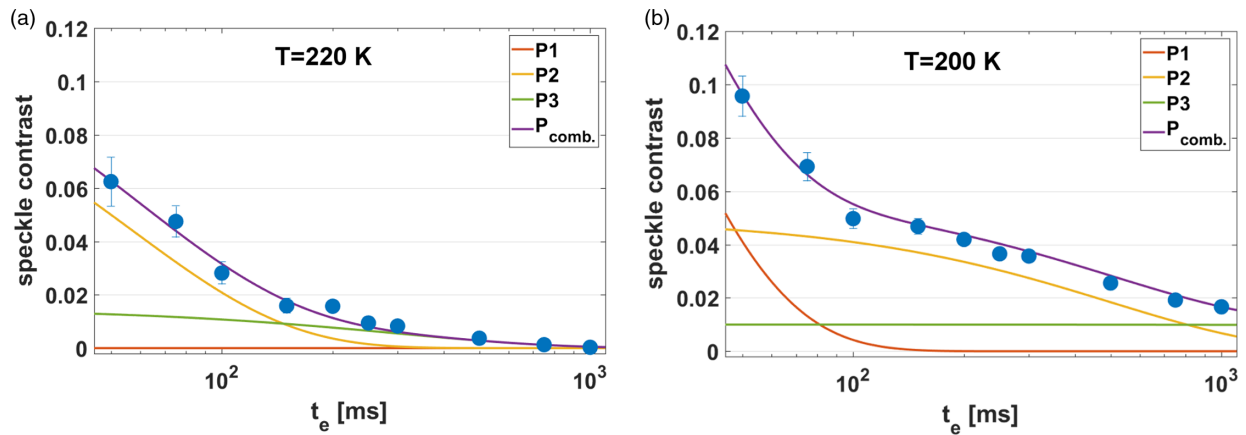


FIG. 3. Speckle contrast of the ionic liquid as a function of the exposure time for temperatures of 220 and 200 K and $Q = 2.6 \text{ nm}^{-1}$, respectively. The solid lines are assignments to three different processes labeled $P1$, $P2$, and $P3$. The calorimetric glass transition temperature is 214 K.

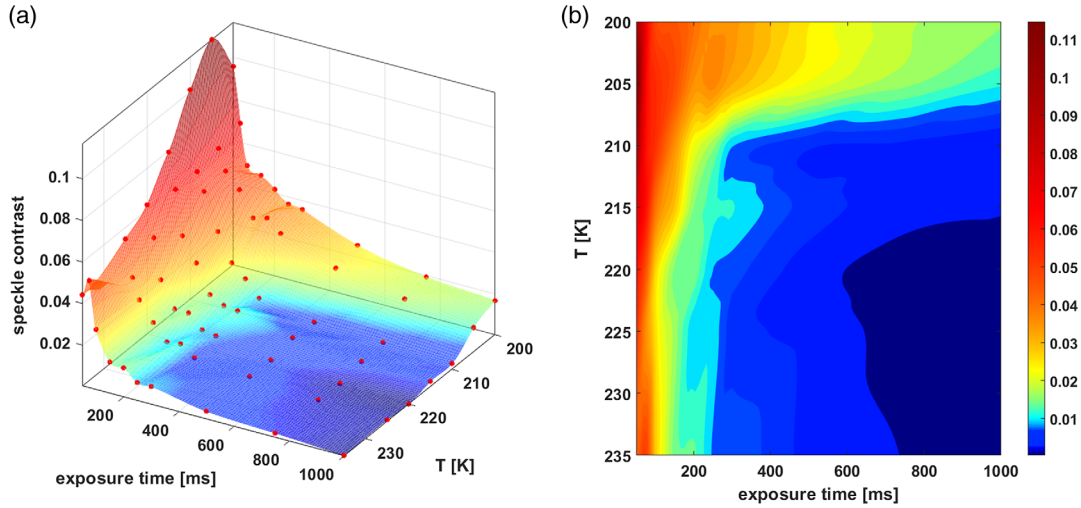


FIG. 4. (a) Speckle contrast as a function of the temperature and delay time for the ionic liquid C8mimCl at $Q = 2.6 \text{ nm}^{-1}$. The dynamics is progressively slowing down upon approaching T_G with a separation in a fast and slow frozen-in dynamics visible at the lowest temperatures. (b) Contour plot of the pattern shown in (a).

The speckle contrast at the largest available delay time of 1 s is a measure of the arrested component P_3 of the dynamics. The temperature dependence of this component is plotted in Fig. 5(a), showing a sudden increase at a temperature which we associate with the glass transition. The maximum value of the measured frozen-in contrast is 0.017, which, when normalized by the maximum contrast of 0.1, yields a nonergodicity parameter [30,31] of $f_c = 0.17$ at 200 K [Fig. 5(a)]. Connecting this to the Debye-Waller factor via $f_c = \exp(-Q^2 u^2/3)$ we calculate the localization length characterizing the range of motion associated with this fast mode. We find a value of $\sqrt{u^2} \approx 0.85 \text{ nm}$, which is smaller than the nanostructure correlation length of $L_{\text{cor}} \approx 2\pi/Q = 2.4 \text{ nm}$. In fact, this value is closer to typical charge alternation correlation lengths

in the IL structure which appear at distances between 0.6 and 0.9 nm. Thus, this fast motion would be in the charge alternation domains and not be related to the apolar domains.

As we cannot disentangle the three processes over the whole temperature range, we chose a different way to obtain further insight into the origin of the here observed dynamics. We evaluate the first moment of the nonfrozen components of the correlation functions representing a mean decorrelation time for each temperature. These deduced time constants as a function of the temperature are shown in Fig. 5(b). Close to T_G , we can disentangle frozen-in and still active motions. The fast averaged motion shows an activation energy of $(25 \pm 5) \text{ kJ/mol}$, which is in the range of values reported for different relaxation processes in the liquid phase

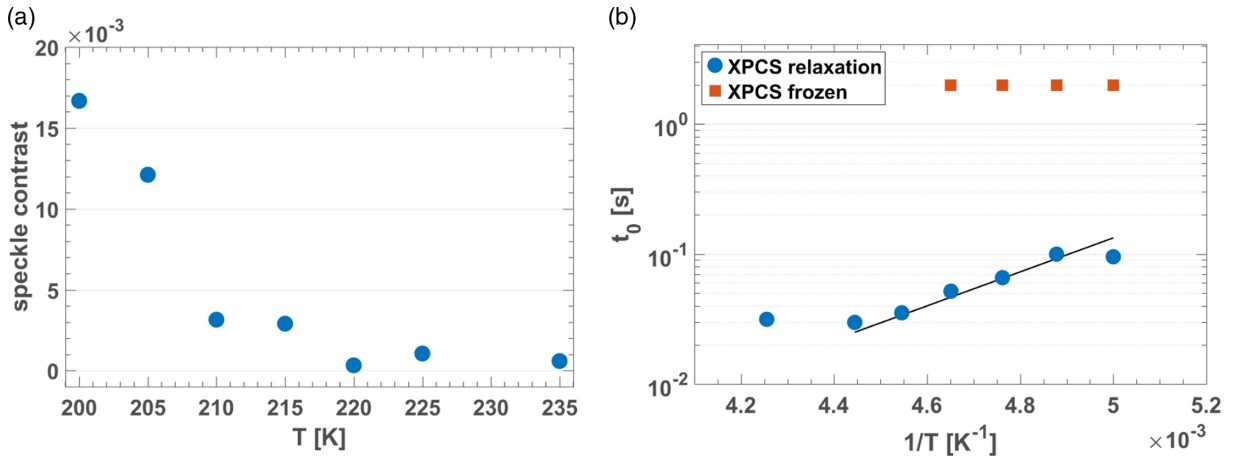


FIG. 5. (a) The x-ray speckle contrast at $t_e = 1 \text{ s}$ exposure time as a function of the temperature. (b) Temperature dependence of the averaged decorrelation time of the nonfrozen part in the IL (blue dots). The red squares represent the appearance of the frozen-in components (i.e., dynamics slower than 1 s). The black line represents an Arrhenius fit with an activation energy of 25 kJ/mol.

such as ionic diffusion processes, nanostructural relaxation, and local reorientation of the alkyl chains and the imidazolium rings [29].

In conclusion, by spreading the dose over a large sample area and evaluating single photon events, we demonstrate that XSVS experiments can be performed at x-ray doses significantly smaller than necessary for typical XPCS experiments. Using this method, we observe the nanoscale dynamics of an ionic liquid below the glass transition. The relaxation in the glassy phase is not completely frozen. Instead, we find different processes which are still active in the glassy phase on timescales of 100 ms and below besides a frozen-in α -type relaxation. This fast dynamic is associated with the peculiar nanostructure of the ionic liquids, which still allows for motion below the calorimetric glass transition. It also becomes apparent that for a deeper understanding of the glass transition a larger range of timescales needs to be probed, which becomes available with the new DLSRs.

Finally, we note that this approach is an important step forward to measure dynamics in biological samples on nanometer length scales. While the dose used at the shortest exposure of 22 kGy is still higher than the tolerable dose of most proteins, some larger proteins can endure doses of up to 8 kGy [32], which is certainly within the range of the presented approach. With larger scattering intensities at smaller Q values, we see no principle obstacle to perform XSVS experiments with doses of 1 kGy and below.

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