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DOI: <https://doi.org/10.1021/jp310090u>

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ZORA URL: <https://doi.org/10.5167/uzh-84257>

Journal Article

Accepted Version

Originally published at:

Thøgersen, Jan; Réhault, Julien; Odelius, Michael; Ogden, Tom; Jena, Naresh K; Jensen, Svend J Knak; Keiding, Søren R; Helbing, Jan (2013). Hydration dynamics of aqueous nitrate. *Journal of Physical Chemistry B*, 117(12):3376-3388.

DOI: <https://doi.org/10.1021/jp310090u>

Hydration Dynamics of Aqueous Nitrate

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Abstract:

Aqueous nitrate, NO_3^- (aq), is studied by 2D-IR, UV-IR and UV-UV time resolved spectroscopy in combination with molecular dynamics (MD) simulations with the purpose of determining the hydration dynamics around the anion. In water the D_{3h} symmetry of NO_3^- is broken and the degeneracy of the asymmetric stretch modes is lifted. This provides a very sensitive probe of the ion-water interactions. The 2D-IR measurements reveal excitation exchange between the two non-degenerate asymmetric stretch vibrations on a 300 fs timescale concomitant with fast anisotropy decay of the diagonal peak signals. The MD simulations show that this is caused by jumps of the transition dipole orientations related to fluctuations of the hydrogen bonds connecting the nitrate ion to the nearest water molecules. Reorientation of the ion, which is associated with the hydrogen bond breaking, is monitored by time resolved UV-IR- and UV-UV spectroscopy, revealing a 2 ps time constant. These time scales are very similar to those reported for isotope-labeled water, suggesting that NO_3^- (aq) has a labile hydration shell.

Keywords: solvation, hydrated ions, hydrogen bonds, rotational anisotropy, pseudo-rotation 2D-IR spectroscopy, transient absorption spectroscopy, Car-Parrinello molecular dynamics simulations.

Introduction

When ions are dissolved in water, they can interact strongly with the nearest water molecules and influence the hydrogen bond network between the water molecules. This process is called hydration and the water molecules are said to form hydration shells around the ions. Hydrated ions are omnipresent on Earth. From hydration of salts in the oceans to ion transport through cellular membranes, the success of a vast number of chemical and biological processes relies decisively on the mechanisms of hydration. Decades of intensive research have greatly advanced the understanding of how ions are hydrated,¹⁻³ yet fundamental questions remain unanswered. Central to the ongoing studies is to what extent the hydration shells are stable with long lasting ion-water hydrogen bonds, or labile with frequent exchanges of molecules, and how these processes determine the angular and translational mobility of the ions.

In neat liquid water the molecules are interconnected by a network of hydrogen bonds linking a hydrogen atom of one molecule to an oxygen atom of another. The hydrogen bonds are 10-100 times weaker than the covalent bonds and each water molecule may form hydrogen bonds with up to four neighboring water molecules. At room temperature the hydrogen bond lengths and angles fluctuate strongly on a sub-picosecond timescale,^{4,5} but these fluctuations result in only modest changes of the local water structure. However, sometimes a hydrogen bond is broken and a bond to a new partner is formed concomitant with the reorientation of the water molecule.⁶ The water molecules reorient on average every 2.5 ps,⁷ and the hydrogen bond thus fluctuate several times before reorientation occurs. Recent models developed from molecular dynamics simulation⁸ indicate that the reorientation mainly occurs in sudden large angle jumps involving the concerted cleaving and formation of hydrogen bonds. Direct verification of this picture is difficult to obtain from experiments in neat water,⁹ but the models are able to reconcile

seemingly contradictory reorientation time constants derived from NMR, IR and Neutron scattering data.¹⁰ On the other hand, experimental evidence that a jump mechanism is responsible for the reorientation of water in the solvation shell of ions has been presented recently.^{11,12}

The hydrogen bond network changes considerably when ions are dissolved in water. Although it is still debated whether the bulk water network is affected or not,¹²⁻¹⁸ several studies of hydrated ions indicate that the nearby water molecules form quasi stable hydration shells with ion-water hydrogen bonds lasting much longer than the hydrogen bonds of neat water.^{18,19} In the case of $\text{Br}^- (\text{aq})$ ²⁰ the hydrogen shell was found to remain intact for 5 ps, while the hydration shells of $\text{BF}_4^- (\text{aq})$ ¹⁹, and $\text{ClO}_4^- (\text{aq})$ ¹⁸ exchange water molecules on average 7 ps. Polarization sensitive two-dimensional infrared (2D-IR) spectroscopy in combination with Car-Parrinello MD modeling suggest that the water molecules that are hydrogen bonded to the ion reorient in sudden large angle jumps involving the concerted breaking of a hydrogen bond to the ion and the formation of a hydrogen bond to a water molecules in the second hydration shell.¹¹ In between the angular jumps the water molecules slowly reorient with the ion in an intact ion-water shell.

The present study concerns the hydration dynamics of aqueous nitrate, $\text{NO}_3^- (\text{aq})$. In contrast to ions like BF_4^- ,¹⁹ and $\text{ClO}_4^- (\text{aq})$,¹⁸ there is no distinct fingerprint for hydrogen bonding to $\text{NO}_3^- (\text{aq})$ in the OH stretch spectrum of water, which is therefore not a suitable probe. Nevertheless, aqueous nitrate is an excellent molecule for studying ionic hydration shell dynamics, because its spectroscopic properties depend strongly on the hydrogen bonding interactions with the surrounding water, which lift the degeneracy of the two asymmetric stretch vibrations²¹. Hence, contrary to most previous experiments, which measure the dynamics of the ion-water hydrogen bonds by vibrational spectroscopy of the water molecules, the present

experiments probe the spectral signatures of the ion. This yields information of the hydrogen bonds connected to the molecular anion with little, if any, background signal from bulk solvent.

Previously, lineshape analysis of polarized Raman spectra and their temperature dependence have been used to derive picosecond reorientation time constants for several molecular ions in water including nitrate.²² Here we use 2D-IR and pump-probe spectroscopy to investigate the fluctuations in the ion-water hydrogen bonds, the intra-molecular coupling of the vibrational modes of nitrate and the reorientation of the nitrate molecules. A similar approach was recently taken by Hochstrasser and co-workers, who investigated the water-induced relaxation of Guanidinium (D_3 -symmetry).²³ Compared to their study, the nitrate ion offers the advantage of a much larger solvent-induced splitting of the degenerate vibrations, already in pure water (see below), which also allows us to better differentiate between excitation transfer and normal mode mixing processes.

The quality of our 2D-IR data deteriorates for timescales longer than 1 ps, due to a background signal from scattered light and due to the short lifetime of the vibrational states. Reorientation dynamics occurring on longer time scales are therefore monitored by time resolved spectroscopy involving excitation of the electronically excited states of nitrate. Thus UV pump–UV probe and UV pump–IR probe transient absorption spectroscopy may follow the nitrate–water interactions for up to 10 ps. The experimental findings are interpreted with the aid of both classical molecular dynamics (MD) simulations and Car–Parrinello molecular dynamics (CPMD) simulations. The combination of time resolved 2D-IR, UV-UV and UV-IR spectroscopy together with molecular dynamics simulations gives a comprehensive and consistent description of the hydration of aqueous nitrate.

Experimental section

Measurements involving UV excitation were carried out at the University of Aarhus, while the 2D-IR data were taken at the University of Zürich. Both laboratories utilized a 1 kHz amplified Ti:Sapphire laser system emitting 100 fs pulses at 800 nm with an energy of about 1 mJ.

The two-dimensional infrared (2D-IR) spectrometer in Zürich used a beam of mid-infrared femtosecond pulses from an optical parametric amplifier (OPA). Two reflections off a wedged CaF₂ window produced the probe and reference beams. Pairs of collinear pump pulses were generated by a fast-scanning interferometer with automated phase determination.²⁴ These passed a computer-controlled waveplate, which was rotated at regular intervals (approximately 20 seconds), switching the pump polarization between parallel and perpendicular relative to the polarization of the probe pulses for the quasi-simultaneous recording of spectra of different polarization conditions. Probe and reference beams were then spectrally dispersed in a monochromator and detected by a double array detector on a single shot basis as a function of coherence time t_1 between the pump pulses and the waiting time t_2 between the second pump pulse and the probe pulse. Fourier transform along t_1 directly yielded absorptive 2D-IR spectra.

Two types of transient absorption measurements were performed in Aarhus, UV pump-UV probe and UV pump-IR probe. Both measurements utilized 200 nm pump pulses of 200 fs duration generated by frequency quadrupling the 800 nm laser pulses in three consecutive frequency doubling and sum-frequency mixing BBO crystals. The pump pulses were sent through a scanning delay line and a $\lambda/2$ waveplate before they were focused through the sample by a concave mirror. The beam of 195 nm UV probe pulses was generated by frequency upconverting the output of an OPA and split into a signal and a reference beam. The signal beam was focused

on the sample by a concave mirror thus probing the sample inside the area defined by the pump beam, while the reference beam bypassed the sample. Signal and reference pulses were subsequently detected by matched photodiodes. The anisotropy data were derived from individual transient absorption measurements recorded with parallel (ΔA_{\parallel}) and perpendicular (ΔA_{\perp}) pump and probe polarizations.

For UV pump-IR probe spectroscopy, the IR probe pulses covering the entire asymmetric stretch absorption band of NO_3^- (see Fig. 1) were generated by difference frequency mixing the output pulses from an OPA. The probe beam was divided into signal and reference beams by a silicon beam splitter. The signal beam probed the sample inside the volume excited by the UV pump pulse, whereas the reference beam passed through the sample above the pump beam. The polarization of the ultraviolet pump beam was fixed at 45° with respect to that of the infrared probe pulse. An analyzing polarizer interchangeably selected the infrared signal and reference components parallel and perpendicular to the polarization of the pump beam allowing the absorption changes induced parallel (ΔA_{\parallel}) and perpendicular (ΔA_{\perp}) to the pump pulse polarization to be measured by a spectrometer equipped with a double array detector. Each measurement consisted of many consecutive pump-probe delay scans. The analyzing polarizer was rotated between each scan for quasi-simultaneous recording of ΔA_{\parallel} and ΔA_{\perp} .

The samples used in Aarhus consisted of a constantly flowing film of 14 mM (UV pump-UV probe) or 0.1 M (UV pump-IR probe) aqueous KNO_3 solution suspended between two parallel, 50 μm thick titanium wires separated by 4 mm.²⁵ For the 2D-IR experiments 1 M aqueous solution of NaNO_3 was sandwiched between two 2 mm thick CaF_2 windows with a 6 μm spacer. The peak absorption of the asymmetric stretch band of NO_3^- was approximately 0.3 on a broad

water background of roughly the same size. Control experiments were carried out without spacer, yielding samples of approximately 0.5 μm thickness, and in D_2O . No significant differences were observed. Furthermore, the band shape of the background-corrected asymmetric stretch bands of NO_3^- (aq) in the FTIR spectrum was identical at concentrations of 0.1 M and 1 M.

Theoretical section

Molecular dynamics simulations were performed to elucidate the connection between hydrogen bond fluctuations and spectral dynamics, the splitting of the asymmetric N-O stretch band and the reorientation of the NO_3^- (aq) transition moments. First, in order to investigate the concentration dependence and finite size effects, classical MD simulations at 298 K were performed for several nanoseconds on an electrolyte solution of K^+ and NO_3^- in 55 and 550 water molecules, corresponding to 0.1 M and 1.0 M concentrations, at densities of 1.0 and 1.05522 g/cm^3 .²⁶ The force field parameters for water and the potassium cation were as provided by the MDynamix database,²⁷ the force field for the nitrate ion was built from intermolecular and intra-molecular potentials from references 28 and 29. The size-dependence was studied by comparing the solute-solvent radial distribution functions, which shows small differences between the small and large classical MD simulations. Having confirmed that the small system was large enough to describe the solvation structure, Car-Parrinello MD (CPMD) simulation results were initialized from the small classical MD simulation.^{30,31} Both the classical and the CPMD simulation results were used in the analysis of the experimental data.

In the CPMD simulations, all water molecules were deuterated to ensure a small spectral overlap between the electronic and nuclear degrees of freedom. A time-step of 1 fs was used in combination with a fictitious energy mass of 500 a. u. and a Nose-Hoover thermostat (2500 cm^{-1})

coupling to the OD vibrations. After 20 ps equilibration initialized from the classical MD, the production run was 18.9 ps at 300 K. An analogous CPMD simulation at 350 K was equilibrated for 20 ps, after a period of elevated temperature. We used the BLYP functional, norm-conserving Troullier-Martins pseudo-potentials (using a semi-core description with maximum angular momentum and local p component for potassium) and a plane wave expansion of the Kohn-Sham orbitals with truncation at an energy cut-off of 70 Ry.

Quantum chemical calculations on small clusters cut out in a 5 Å radius around the NO_3^- ion were used to sample vibrational spectra along the classical MD and the CPMD simulation trajectories. In Gaussian09³² partial optimization and vibrational analysis on the BLYP/aug-cc-pVDZ level of approximation were performed on NO_3^- in a solvent environment at frozen geometry for 180 configurations along each trajectory. (The basis set convergence of the results was confirmed by calculations with aug-cc-pVTZ, which reproduce the relative changes in the vibrations for the different configurations within a few cm^{-1}). For the vibrational modes, frequencies and dipole derivatives were used to simulate the IR spectrum and to study the orientational fluctuations in the molecular frame. The fluctuations in the vibrational modes were similar in the classical and the CPMD simulations and we only display the CPMD results. An ad-hoc constant shift was applied to calculated IR frequencies in the comparison to experiment.

Spectroscopy

Before presenting the experimental results we give a brief overview of the spectroscopy of the nitrate ion. The isolated nitrate ion is planar (D_{3h}) with the excess negative charge delocalized over the molecule, and, accordingly, the two asymmetric stretch vibrations are degenerate.

Hydration breaks the molecular symmetry and lifts the degeneracy^{21,33} giving rise to the two absorption bands near 1350 cm⁻¹ and 1400 cm⁻¹ shown in Fig. 1.

Conceptually, the mechanism for the splitting of the asymmetrical stretch vibrations can be formulated in terms of the individual N-O groups. Through intermolecular interactions, primarily hydrogen bonding of the oxygen atoms in NO₃⁻ but also long-range electrostatic fields, the N-O groups experience variations in the shape of the stretching potentials. The three N-O stretches couple to give rise to a symmetric stretch and two asymmetric stretch normal modes, with small contributions from in-plane bending vibrations. These normal modes are sensitive to the differences in the N-O potentials. Thus, the symmetric stretch mode (near 1100 cm⁻¹) is no longer fully symmetric and the asymmetric stretch modes become non-degenerate. They may, however, still be pictured as orthogonal linear combinations of the degenerate gas-phase modes sketched in Fig. 1, and their transition dipole moments lie in the molecular plane and are perpendicular to each other.

To illustrate the sensitivity of the asymmetric N-O stretch band of NO₃⁻ to hydrogen bonding interactions, we have used electron structure calculations at the level MP2/6-311+G(d,p) to study the frequency splitting of the asymmetric stretch absorption band pertaining to the complexes of NO₃⁻ and a single uncomplicated molecule/atom, X, from the first two periods of the periodic table. The calculated splitting is then related to ΔG° . The trend of the data in Fig. 2 shows that the frequency splitting increases steadily from nearly 0 cm⁻¹ for molecules without a permanent dipole moment, like N₂, to 150 cm⁻¹ for strongly hydrogen bonding species like HF.

Previously, the splitting of the asymmetric stretch vibrations of NO₃⁻ has been calculated for clusters of nitrate with one or two water molecules³⁴, which underestimated the experimental

value for the bulk solvent. It was speculated that in aqueous solution transient charge localization may lead to strong, highly asymmetric H-bonding involving mainly one N-O group. Indeed, recent MD simulations studying the structure and dynamics in the hydration shell of NO_3^- show that there is instantaneously a substantial asymmetry in the hydration both in structure and in dynamics.³⁵ The asymmetry in hydrogen bonding is correlated with transient elongations of individual N-O bonds associated with a charge redistribution in the NO_3^- ion.³⁶ Size selective infrared multiple-photon photo-dissociation spectra of $\text{NO}_3^-(\text{H}_2\text{O})_n$ clusters provide beautiful experimental demonstration of the effect of symmetry in the hydration of the nitrate ion.³⁷ They show a strong splitting of the asymmetric stretch mode for $n=1,2$ and 5 , but only a single resonance for $n=3$, when all three oxygen atoms appear to form similar hydrogen bonds. These results suggest that transient fluctuations of the strength or geometry of the hydrogen bonds between the nitrate anions and the water molecules in the hydration shell will be reflected very sensitively in the linear and non-linear infrared spectra of the asymmetric stretch vibrations.

Results

2D-IR spectroscopy

The 2D-IR data was recorded by exciting and probing the fundamental asymmetric stretch transitions of aqueous nitrate with infrared laser pulses covering the entire absorption band. The measurements were taken with parallel and perpendicular polarizations of the pump and probe pulses. Fig. 3 shows the 2D-IR spectra of $\text{NO}_3^-(\text{aq})$ for a number of time delays between excitation and probing of the asymmetric stretch vibrations. Data recorded with parallel (ΔA_{\parallel}) and perpendicular (ΔA_{\perp}) polarization of pump and probe beams are shown in the first two columns. The magic angle signals, ΔA_{MA} , are calculated from these data using $\Delta A_{\text{MA}} = \Delta A_{\parallel} +$

$2\Delta A_{\perp}$ and plotted in the third column. They show the intensity changes free from re-orientation effects. All spectra are normalized to the minimum negative signal for better comparison. The negative components of the diagonal signals at excitation wavenumbers 1340 cm^{-1} and 1400 cm^{-1} reflect the asymmetric stretch absorption maxima of the spectrum in Fig. 1. These signals are due to the pump-induced bleaching of ground state nitrate ions and stimulated emission from the first excited states of the asymmetric stretch vibrations. Because of band overlap the corresponding red-shifted positive excited state absorption signal is only visible for the lower frequency band. The diagonal signals are slightly tilted along the diagonal due to an inhomogeneous frequency distribution. However, the tilt is modest, in particular for the lower frequency band, indicating strong homogeneous broadening contributions. The diagonal signal intensities decrease with time due to vibrational relaxation of the first excited state. In Fig. 4a we show this decay for the magic angle data of the lower frequency diagonal peak. Both the bleach/stimulated emission and the excited state absorption signals decrease exponentially with a time constant of 550 fs.

Having established the vibrational relaxation time of the asymmetric stretch, we turn to the anisotropy of the 2D-IR spectra. Visual inspection of the first two columns of Fig. 3 reveals a clear distinction between the data recorded with parallel and perpendicular pump and probe pulse polarizations. Hence, the transient absorption data are anisotropic. In Fig. 4b we plot the anisotropy $R(t)$, defined by:

$$R(t) = \frac{\Delta A_{\parallel} - \Delta A_{\perp}}{\Delta A_{\parallel} + 2\Delta A_{\perp}} \quad (1)$$

for the excited state absorption component of the lower frequency diagonal signal of the 2D-IR spectra. It very quickly decreases from $R(0) \approx 0.4$ to $R(1 \text{ ps}) \approx 0.1$, compatible with an

exponential decay with a 300 fs time constant. The residual anisotropy subsequently decays on a much slower timescale that is not fully captured by the 2D-IR experiments, because the scattering background begins to dominate the signal at delays much longer than the vibrational lifetime. At $t = 0$, the diagonal peak anisotropy approaches the theoretical maximum ($R = 0.4$) for non-degenerate transitions, for which $R(t) = 2/5 P_2(\cos\theta)$. Here, P_2 is the second order Legendre polynomial and θ is the angle between the excited and probed transition dipole moments.

The difference between the spectra recorded with parallel and perpendicular pump and probe pulse polarizations also point to the presence of cross peaks near the off-diagonal corners of the grey squares in the first row of Fig. 3 ($\omega_1 = 1340 \text{ cm}^{-1}$ and $\omega_3 = 1400 \text{ cm}^{-1}$ and $\omega_1 = 1400 \text{ cm}^{-1}$ and $\omega_3 = 1340 \text{ cm}^{-1}$). They are partially hidden under the dominant diagonal peaks. At waiting time zero, these cross peaks indicate a coupling of the two asymmetric stretch vibrations. Subsequently, with increasing waiting time between excitation and probing, these cross peak signals gain relative strengths with respect to the diagonal peaks. In order to extract the underlying kinetics, we have integrated the magic angle 2D-IR spectra in two regions with relatively little spectral overlap near $\omega_3 = 1420 \text{ cm}^{-1}$. These are marked by green and purple rectangles in the spectrum recorded after 500 fs in the third column of Fig. 3 (bottom). The normalized difference of these two signals is a good approximation for the relative growth of the cross peaks on the background of the overall signal decay.³⁸ It is plotted in Fig. 4c and can be fitted by a single exponential function with a 300 fs rise time, which is indistinguishable from the anisotropy decay time of the diagonal peaks. A more precise determination of the kinetic time constants requires the fitting of the full 2D-IR spectra. Because of the strong correlation between

the two asymmetric stretch frequencies (see below), established fitting procedures³⁹ would, however, also be only an approximation.

In order to isolate the cross peaks, the dominant diagonal contributions can be eliminated by computing weighted difference 2D-IR spectra⁴⁰ ($\Delta A_{WD} = \kappa \Delta A_{\perp} - \Delta A_{\parallel}$), using $\kappa = (1 - R(t))/(1 + R(t))$, where $R(t)$ is the diagonal peak anisotropy. These difference spectra are shown in the rightmost column of Fig. 3. The solid black lines reproduce the nodal line between bleach and excited state absorption components of the lower right off-diagonal signal at zero waiting time. It is tilted towards the anti-diagonal, pointing to an anti-correlation of the frequencies of the two asymmetric stretch modes. The anti-diagonal tilt of the cross peaks can still be seen in the 300 fs spectrum, but it has decayed after about 500 fs. Because of the uncertainties in subtracting parallel and perpendicular spectra, a more quantitative analysis of the tilting angles is not possible. However, the upper left off-diagonal signals appears to be less tilted, which could indicate different broadening of the two asymmetric stretch transitions.

UV pump – IR probe spectroscopy

We now address the reorientation dynamics of NO_3^- (aq) by UV- IR transient absorption spectroscopy, which complements the 2D – IR results on the picosecond time scale. In the UV pump – IR probe measurements, the rotational anisotropy measurements are initiated by exciting the $\pi\pi^*$ transition of the NO_3^- ions with linearly polarized laser pulses having a wavelength of 200 nm and a duration of 200 fs. Previous studies have shown that excitation of nitrate at 200 nm leads to partial photodissociation, while approximately half of the excited nitrate molecules return to the electronic ground state and relax to the vibrational ground state.⁴¹ The photoproducts are irrelevant to the present work. Since the transition dipole moment of the $\pi\pi^*$

excitation lies in the plane of the ions ($X^1A'_1 \rightarrow ^1E'$ in D_{3h} symmetry),⁴² the plane of the unexcited nitrate ions remaining in the electronic ground state is predominantly perpendicular to the excitation pulse polarization. With time, molecular reorientation results in an isotropic distribution of the molecular planes. This reorientation is monitored by probing the asymmetric stretch transition of NO_3^- with 100 fs linearly polarized infrared laser pulses covering the entire absorption band of the asymmetric stretch vibration. The rotational anisotropy is derived from Eq. 1 by probing the induced change in the absorption measured with the probe polarization parallel (ΔA_{\parallel}) and perpendicular (ΔA_{\perp}) to the pump pulse polarization. Because the transition dipole moments pertaining to the $\pi\pi^*$ transition and the asymmetric stretch transitions both lie in the plane of the ions, only out of plane molecular rotations contribute to $R(t)$. The amplitude of $R(t)$ is therefore a measure of the fraction of out of plane rotations relative to in-plane rotations and has a maximum theoretical value of $R = 0.1$. Key to applying this experimental approach to elucidate the hydrated ion dynamics is that it selectively measures the rotation of electronic ground state $\text{NO}_3^-(\text{aq})$ ions, which are *unaffected* by the excitation pulse. That is, it measures the reorientation of NO_3^- in thermal equilibrium with the bulk water network.

The time dependent photoinduced change in the infrared absorption $\Delta A(\nu, t)$ pertaining to the asymmetric stretch vibration of the aqueous nitrate solution is depicted in the contour plot in Figure 5. The data set is recorded with the pump beam polarization parallel to the probe polarization. A similar data set is recorded with perpendicular pump and probe beam polarizations (not shown). The absorption dynamics have been described in detail in a previous publication⁴³ and are only summarized here in order to explain how the reorientation dynamics are extracted from the data. The transient absorption data are dominated by a strong negative absorption change (blue colors) with minima at 1340 cm^{-1} and 1400 cm^{-1} reflecting the

asymmetric stretch absorption band in Fig. 1. The negative absorption change is caused by pump induced depletion of ground state nitrate population. The absorption partially recovers as half of the excited nitrate ions return to the ground state during the first 10 ps. The vibrational relaxation of the nitrate ions associated with their return to the ground state is seen as the positive induced absorption (red and yellow colors) on the low frequency side of the asymmetric stretch absorption band. Due to the anharmonic ground state potential of nitrate this induced absorption propagates towards higher frequencies as the excited molecules relax to the vibrational ground state in 2-3 ps. The induced absorption pertaining directly to the nitrate ion is superimposed on a background covering the entire spectrum. At low frequencies the initial dynamics of the background is obscured by the absorption associated with vibrationally relaxing nitrate molecules, whereas the transient absorption signal at high frequencies reveals an induced absorption decaying on a 2 ps timescale. A similar absorption background was observed in a previous photolysis experiments on aqueous nitrite and shown to replicate the absorption change observed when weakening the hydrogen bonds by heating the nitrite solution.⁴⁴ Accordingly we ascribe the absorption background observed in aqueous nitrate to heated water molecules in vicinity of the photo-excitation site.

An accurate analysis of the reorientation dynamics of the nitrate ions necessitate the separation of the absorption dynamics of ground state nitrate ions from the absorption of vibrationally excited nitrate ions and the thermally induced absorption changes of water. Here, this is done by Singular Value Decomposition (SVD) of the data set. Despite the fact that the timescales for the absorption of the three contributions are similar, SVD convincingly separates three components. The weighted spectra of the three dominating components are depicted in Fig.

6 a-c. The two curves in each figure represent the data obtained with the pump beam polarization parallel respectively perpendicular to the probe beam polarization.

Together the three components derived by SVD account for 97 % of the measured transient absorption dynamics thus giving a close to complete description of the data. Several important conclusions may be inferred from this analysis. The spectral component shown in Fig.6a dominates the contributions shown in Fig. 6b and Fig. 6c by an order of magnitude. The shape of the SVD spectra in Fig.6a derived from the measurements obtained with the two different polarizations is identical, but the (absolute) amplitude of the spectrum associated with parallel pump and probe beam polarizations is slightly larger than that obtained for perpendicular pump and probe polarizations. This is distinctly different from the spectra of the minor contributors depicted in Fig. 6b and Fig. 6c, for which the shape as well as the amplitude of the spectra obtained for different pump and probe polarizations are identical. Accordingly, only the dominating spectral component is anisotropic. Since the time dependence of each SVD component is inherently the same at each frequency and the two minor SVD components are isotropic, the anisotropy is frequency independent.

In general the spectra derived by SVD represent the orthonormal basis of the spectral components rather than the spectra of the molecular species themselves. Nevertheless, comparing the dominating SVD component to the asymmetric stretch absorption band (see Fig. 6d) reveals a close similarity between the two spectra. The similarity agrees with the fact that the (negative) transient absorption resulting from photoinduced depletion of ground state nitrate ions dominates the data in Fig. 5. Hence, the dominating SVD spectrum is assigned to ground state nitrate ions and will be used as such in the following derivation of the rotational anisotropy of the aqueous nitrate ions. The assignment of the spectra in Fig.6b and Fig. 6c is less certain.

Guided by the red shift of the absorption relative the spectrum of ground state hydrated nitrate, the spectrum in Fig. 6b may reflect the absorption associated to vibrationally relaxing nitrate molecules, while the spectrum in Fig. 6c is tentatively ascribed to absorption changes induced by the local temperature increase in the sample when the NO_3^- ions are excited at 200 nm. The SVD analysis shown in Fig. 6 is performed on delays between 0.5 and 44 ps. Still, the SVD analysis is rather insensitive to the chosen time interval. Thus, the SVD results remain virtually unaffected when changing the time interval for the analysis from 0.5-5 ps to 0.5-44 ps. Only when using a time interval as short as 0.5-2ps do we observe a slight change in the spectral component assigned to vibrationally relaxing nitrate ion. This change is to be expected as the spectrum of vibrationally relaxing nitrate changes within the first two picoseconds.

The rotational anisotropy, $R(t)$, is calculated by inserting the two polarization components of the SVD data assigned to the induced change in nitrate's ground state absorption into Eq. (1). To improve the statistics $R(t)$ is averaged over all probe frequencies (remember, the anisotropy is frequency independent). The resulting rotational anisotropy data presented in Fig. 7a is well approximated by a single exponential decay convoluted with the experimental response time. The initial anisotropy is $R(t = 0) = 0.05 \pm 0.01$ and the decay time is $\tau = 1.7 \pm 0.3$ ps. The initial anisotropy of $R(t=0) = 0.05 \pm 0.01$ is somewhat lower than the maximum value of $R=0.1$ expected for the applied pump-probe geometry. The origin of this discrepancy is unclear, but may be due to minor inaccuracies of the SVD analysis. The polarization sensitive UV pump-UV probe transient absorption measurements presented next confirms the $\tau = 1.7 \pm 0.3$ ps reorientation time.

UV pump-UV probe spectroscopy

The UV pump–UV probe rotational anisotropy measurements are initiated by exciting the $\pi\pi^*$ transition of the NO_3^- ions with linearly polarized laser pulses having a wavelength of 200 nm. The subsequent 195 nm probe pulses address the same $\pi\pi^*$ transition and measure the reorientation of the nitrate molecules remaining in the ground state as they relax to an isotropic distribution. Since the transition dipole moments pertaining to the $\pi\pi^*$ transition lie in the plane of the ions, only out of plane molecular rotations contribute to $R(t)$. Similar to the UV pump–IR probe measurements the amplitude of $R(t)$ therefore has a theoretical maximum value of $R = 0.1$. Fig. 7 b shows the rotational anisotropy measured at 195 nm following the excitation of hydrated nitrate ion at 200 nm. The measured anisotropy is well approximated by a single exponential decaying function having an initial value of $R(t=0) = 0.1 \pm 0.01$ and a time constant of $\tau = 1.7 \pm 0.3$ ps. This reorientation time is in full agreement with the $\tau = 1.7 \pm 0.3$ ps derived from the UV pump – IR probe measurements and furthermore agrees with the 2 ± 0.2 ps rotation time derived from line-width analysis of Raman spectra of aqueous nitrate by Perelygin et al. ²². Consequently, there is compelling experimental evidence that the hydrated nitrate ions perform out-of-plane rotations on a 2 ps timescale. The initial value of the rotational anisotropy of $R(t=0) = 0.1$ agrees with the theoretical maximum value expected for molecules with transition dipole moments defining a plane.

MD simulations

Classical MD and CPMD simulations were carried out in order to obtain information about the reorientation of the vibrational transition dipole moments for direct comparison with the experimental data. Since aqueous nitrate has already been the subject of a number of MD and QM/MM simulations in the past, we begin with a discussion of the solvation structure around the

solute in order to motivate our approach. In Fig. 8, the radial distribution functions (RDFs) of the atoms in nitrate (N,O) and the atoms in water (O_w, H_w) are presented.

The size dependence of the solvation structure in the classical MD simulation is weak, and there is only moderate temperature dependence in the CPMD simulations. However, the hydration shell of NO_3^- is much less structured in the CPMD simulations than in the classical MD simulations. In CPMD, the hydration shell structure around NO_3^- is weak with rapid fluctuations in the hydrogen bond network. This is in qualitative agreement with earlier *ab initio* quantum mechanical charge field MD studies.^{35,45} For the N- O_w and O- O_w RDFs, there is a good agreement between our CPMD results and previous QM/MM results⁴⁵ obtained with the QM-B3LYP approximation. However, there are significant differences between CPMD and QM/MM methods for the RDFs involving H_w and in the water structure outside the first hydration shell. The consistent force field in CPMD provides a balanced description of solvent and solute, but it is known to predict a too structured hydrogen bond network in water. However, the local structure in the hydration of the nitrate ions predicted by CPMD is in agreement with QM/MM and deemed more trustworthy than the classical MD. Our classical MD simulations were based on a standard point charge model, which clearly gives too strong hydration of the nitrate ion. In a follow-up study, we will analyze the structure and dynamics of prolonged CPMD simulations for comparison to other classical MD force fields, including that derived in the QM/MM study.⁴⁵

We also made a comparison of the reorientational dynamics between the CPMD and classical MD simulations. Fig. 9 presents the time-correlation function (TCFs) of the 2nd order Legendre polynomial, $\langle P_2[u(t_0+t), u(t_0)] \rangle \sim R(t)$ (equation 1) of the N-O vector in the nitrate ion and that of O-H vectors in the water molecules. Time scales for the reorientation processes are derived by fitting the TCFs in Fig. 9 to single exponential decays in the interval 1-4 ps. In the

classical simulation, the $P_2(u)$ TCF shows that orientational motion of the nitrate ion ($\tau = 3.225 \pm 0.005$ ps) is slower than for the water molecules ($\tau = 2.794 \pm 0.005$ ps). In CPMD however, the relationship is reversed at both temperatures, and the water molecules undergo very slow reorientation at 300 K. Thus, for the CPMD simulations at 300 K (350 K) the time constant of the correlation function of the nitrate ion is $\tau = 3.1 \pm 0.2$ ps ($\tau = 1.5 \pm 0.2$ ps) and for the water molecules it is $\tau = 6.71 \pm 0.05$ ps ($\tau = 2.30 \pm 0.05$ ps). The uncertainty of the correlation times of nitrate from the CPMD simulation is largely due to the limited sampling. A ratio of two between the correlation times of nitrate and water is significant and too large to be explained by the deuteration used in the CPMD simulations. When comparing to the experimental results it is important to remember that quantum effects strongly affect the reorientational correlations time in water.⁴⁶ The slow dynamics in CPMD simulations is usually handled by an ad hoc temperature increase, which is why we also performed a CPMD simulation at 350 K.

For comparison to the vibrational probes, we calculate harmonic frequencies in quantum chemical calculations on clusters cut out from the periodic simulation models. A full theoretical treatment of the non-linear spectroscopy would require a quantum mechanical treatment involving overtones²¹ and very expensive long trajectories. Our current objective is to identify the relation between the molecular interactions and the spectral response. Hence, we simply sample static spectra over the MD trajectories. While the present CPMD simulation is too short to calculate accurate correlation functions, the sampling is sufficient to obtain a qualitative picture of the mechanism for the 2D-IR experiment. The absorption bands of the stretching modes were calculated by sampling static vibrational spectra over configurations along the MD trajectory. The spectrum resulting from the CPMD simulation, shown in Fig. 10a satisfactorily reproduces the frequency as well as the width and frequency splitting of the experimental

asymmetric stretch band. In the lower part of Fig. 8a, a similar analysis of the configurations in the classical MD simulation reveals a splitting which is over-estimated. Accordingly, the differences in the solvation structure noticed in the RDFs in Fig. 8 around the nitrate ion are reflected in the IR spectrum. This shows that the combination of the reorientation dynamics and the IR spectrum gives constraints on the solvation of the nitrate ion. Fig. 10b shows the instantaneous frequencies as a function of time. In agreement with the 2D-IR data, there is clear anti-correlation between the high- and low-frequencies in the sampled asymmetric stretches. A strong red-shift in one of the asymmetric stretches is for the same configuration associated with a reduced red-shift in the other stretching frequency. The distribution of the symmetric stretch frequency is narrower, but the variations are also correlated with the asymmetric stretching modes. The variations in frequency are related to the hydration of the N-O groups. Hence, we can use the splitting to characterize the degree of asymmetry in the hydration shell.

Fig. 11 a) shows the orientation of the transition dipole moment relative to a given N-O bond for the high and low frequency components of the asymmetric stretch absorption band. In the graph we can follow how the angle between a given N-O vector and the transition moments changes on a sub-picosecond timescale. The Car-Parrinello MD simulations also reveal how the frequency fluctuations of the asymmetric stretch vibrations correlate with the reorientations of the vibrational transition moment. The weak correlation between the angular changes in Fig. 11 a) and the frequency fluctuations in Fig. 10 b) suggests that the angular jumps occur predominantly when the frequency splitting is small, i.e. when the solvation shell is nearly symmetric. This indicates that the anisotropy decay observed in 2D-IR experiments with a time constant of 300 fs can be related to reorientations of the transition dipole moments, which do not

require the physical rotation of the NO_3^- anion in the laboratory frame. Hence, we will refer to these dynamics as pseudo-rotations in the molecular frame.

Discussion

To summarize the results, the reorientation of NO_3^- in water has been investigated by three complementary ultrafast spectroscopic probes. In the all-infrared 2D-IR experiments we both excite and probe the asymmetric stretch transition dipoles. The initial anisotropy close to 0.4 is a signature of a non-degenerate transition, and directly confirms that the degeneracy of the asymmetric stretch vibration of the molecule is lifted in aqueous solution. As a result, both in-plane as well as out-of-plane reorientation may be detected. Within the first few hundred femtoseconds after excitation, the transition dipole moments of the asymmetric stretch vibrations partially lose the memory of their original orientation. The anisotropy decays to a residual value of 0.1, characteristic of a plane-degenerate transition. Consistent with this observation, the CPMD simulations report a large number of orientation-flips within a few picoseconds of the IR transition dipole moments in the molecular frame (see Fig. 11). Accordingly, the 300 fs time-constant in our experiments should be assigned to an in-plane pseudo-rotation of the transition dipole moments that does not require the reorientation of the ion itself. Physical reorientation (tumbling of the molecular plane) takes picoseconds. This motion is probed by the anisotropy decay in the UV-UV and UV-IR experiments, where we excite a plane-polarized electronic transition and are only sensitive to the re-orientation of the molecular plane. We thus confirm an earlier analysis of the symmetric stretch Raman band of nitrate²², which had produced a remarkably similar time constant (2 ± 0.3 ps).

The fast anisotropy decay of the diagonal peaks in the 2D-IR spectrum takes place on the same timescale as the redistribution of vibrational excitation between the two asymmetric stretch modes of NO_3^- (aq), which can be monitored by the relative growth of the cross peak signals with a 300 fs time constant (Figs. 3 and 4). Usually, for non-degenerate transitions, the transfer of excitation from one vibrational mode to another does not lead to a significant reorientation of the corresponding transition dipole moments. In fact, the anisotropy of cross peaks arising from excitation transfer may often even be used to obtain information on molecular structure⁴⁷. For a plane-degenerate excited state, on the other hand, coherent linear combinations of the two transitions may be excited, which can lead to initial anisotropy values different from 0.4^{48, 49}. Anisotropy decay to a value of 0.1 is then a signature of the formation of a statistical mixture of the two transitions after dephasing⁵⁰.

Aqueous nitrate is an intermediate case, where the solvent-induced splitting of the two asymmetric stretch transitions is sufficiently strong to produce two separate absorption bands. The 2D-spectra also separate diagonal and cross-peak contributions which overlap and can cause unusual anisotropy values in conventional pump-probe data. Despite the clear distinction between excitation and probing of lower and higher frequency modes in our experiments, we observe in-plane randomization of the transition dipole moment orientations on an ultrafast timescale. Excitation transfer alone, as for example between different water molecules with overlapping OH stretch spectra⁵¹, is not sufficient to account for this observation. For illustration consider the schematic picture of solvated nitrate ions in Fig.12. The splitting of the asymmetric stretch band in Fig. 12a arises from the elongation of one N-O bond due to (stronger) hydrogen bonding, essentially stabilizing one of the resonance structures of NO_3^- . The corresponding transition dipole moments are then oriented in the molecular frame as indicated by the blue

(higher frequency) and red (lower frequency) solid arrows. In this configuration, transfer of excitation from one asymmetric stretch mode to the other would lead to cross peak growth and diagonal peak decay, but it would not affect the anisotropy of the diagonal peaks. The same is true for the configuration sketched in Fig. 12b, in which one nitrogen atom is more weakly hydrogen bonded than the other two. Fluctuations of the hydration shell, on the other hand, may rapidly stabilize different ‘resonance structures’ of the nitrate ion, leading to a re-orientation (Fig. 12c and d) of the transition dipole moments, or there may even be an exchange of frequency ordering (transition between configurations a and b). As the normal mode coordinates change, the initial excitation is projected onto a new basis. Population redistribution and anisotropy decay thus have a common cause and take place concomitantly.

In a study of guanidinium in water and water-glycerol mixtures, Hochstrasser and coworkers have recently analyzed in detail the effects of excitation versus energy transfer on the anisotropy of 2D-IR and dispersed photon echo signals of a quasi-degenerate mode²³. They concluded that in their case fast anisotropy decay is primarily caused by excitation transfer, whereas reorientation of the transition dipole moments is slower. For guanidinium, however, a weak band splitting, which allows one to distinguish cross peak and diagonal peak contributions, can only be observed in water-glycerol mixtures but not in pure water. It is thus possible that the interaction with glycerol stabilizes an asymmetric charge distributions in guanidinium for a longer time than water can support a given hydration shell around nitrate. The 2D-IR model spectra presented in ref. 23 on the other hand clearly illustrate that the diagonal peak anisotropy cannot decay by excitation transfer alone.

Note that transitions between configurations like those shown in Fig.12 require only modest changes in the strength of the hydrogen bonds between water molecules and the ion. Indeed,

already previous MD simulations of the structural dynamics of nitrate's hydration shell showed a substantial asymmetry in both the structure and dynamics of the hydrogen bonds³⁵. This asymmetry is correlated with transient elongations of individual N-O bonds associated with charge redistribution in the NO_3^- ion.³⁶ The charge redistribution dynamics were accompanied by rapid (0.2-0.9 ps) fluctuations of the hydrogen bonds. In our Car-Parrinello simulations we find that the lower frequency asymmetric stretch transition dipole moment is always confined to an angle section of approximately $\pm 30^\circ$ perpendicular to the shortest N-O bond, as shown in Fig. 12e and f. The idealized configurations in Fig. 12b-d can be regarded as a qualitative representation of this result. Aligning all computed NO_3^- structures along the longest N-O bond yields a somewhat wider transition dipole angle distribution (not shown). A single long intramolecular bond as sketched in Fig. 12a thus seems to be less frequent. Indeed, most structures are characterized by three distinct bond lengths, which lead to the observed distribution of angles. The N-O bond length reflects the strength of hydrogen bonding to that N-O group. Hence, in terms of the solvation, the observed asymmetry means that the situation with an enhanced hydrogen bonding at two of the N-O groups is more frequent than a single N-O group being strongly hydrogen bonded. Hence, the distribution of transition moments taken along the shortest N-O will be narrower than the distribution along any one of the longer bonds. For reorientation within the cone in Fig. 12f the longer two bonds exchange their role, whereas the large jumps in transition dipole orientation seen in Fig. 11 are always accompanied by the lengthening of the formerly shortest N-O bond and the shortening of a previously longer bond. These jumps may therefore be associated with charge redistribution in the ion as a result of changes in the hydration shell structure. The anisotropy decay and excitation transfer in the 2D-

IR spectra of the asymmetric stretch vibrations thus directly report on the timescale of hydrogen bond fluctuations between water and the ion.

In a systematic analysis of nitrate's asymmetric stretch band Hynes and co-workers²¹ have treated the ion-bath interaction as forces in normal mode coordinates of undistorted NO_3^- . Already when restricting the calculations to only the two anti-symmetric stretch vibrations it was possible to reproduce the observed band splitting.¹ Other normal modes, in particular force components along the symmetric stretch coordinate, needed to be included in the analysis in order to quantitatively match the shape and asymmetry of the experimental spectrum²¹. Consideration of only the asymmetric stretch vibrations necessarily results in a perfect anticorrelation of the two asymmetric stretch frequencies. Force fluctuations along the symmetric stretch coordinate, on the other hand, seem to be mainly responsible for (inhomogeneous) broadening and tend to shift both asymmetric stretch transitions. Indeed, the small tilt of the cross peaks in our 2D-IR spectra indicate only a relatively weak anticorrelation, in agreement with the frequency trajectories of our CPMD simulations. Further reduction of the tilt of diagonal and cross peaks is certainly due to homogenous broadening contributions. These may arise not only from fast frequency fluctuations, but directly from the rapid re-orientation of the transition dipole moments. Because of this complexity the detailed analysis of the 2D-IR line shapes is left for future work. Here we have concentrated on the direct implications of our observations for our picture of the water-ion interactions. On the 300 fs timescale discussed so far, nitrate does not rotate, as the time-constant of rotational diffusion in the UV-IR and UV-UV measurements is approximately 2 ps. Hence the hydrogen bonds fluctuate several times before the ion reorients. It is unlikely that reorientation with a 2 ps time constant is possible without an exchange of

¹Their parameters for a two-dimensional potential may even be 'abused' to predict a ratio of diagonal/off-diagonal anharmonicities (3:1) that is in qualitative agreement with the ratio of diagonal over cross peak intensities in our 2D-IR spectra.

hydrogen bonding partners, so nitrate cannot be dragging a longer-lived hydration shell. The CPMD simulations confirm that hydrogen bonds are indeed broken and re-formed with new partners. Interestingly, very similar time constants for hydrogen bond fluctuations of sub ps^{4,5} and rotational diffusion of 2.5 ps have been reported for isotope labeled water.⁷ Despite its negative charge, nitrate does not seem to strongly perturb its local environment. However, because NO_3^- does not distort the water spectrum like, for example, BF_4^- or ClO_4^- , we cannot directly probe the reorientation of the water molecules surrounding the ion. The spectroscopy of the asymmetric stretch vibrations and Car-Parrinello MD simulations nevertheless suggest that the hydration ‘shell’ of nitrate is by no means a static entity – in fact it appears to be just as labile as neat water.

In agreement with the experimental results the CPMD simulations find that the reorientation of the nitrate ion may be faster than the surrounding solvent molecules. Comparison of the results from the CPMD simulations and the classical MD simulation show that the ratio of the reorientational correlation times of solute and solvent depends on the solvation structure. In CPMD the fast motion of the nitrate molecules is associated with a weakly structured first solvation shell.

Conclusion

In summary, the anisotropy decay in 2D-IR and transient absorption spectra of aqueous nitrate was analyzed with the help of Car-Parrinello molecular dynamics simulations. Fluctuations of the solvating water molecules provoke frequent symmetry-breaking N-O bond length changes. The instantaneous normal mode coordinates of the asymmetric stretch vibrations, whose degeneracy is lifted by these asymmetric distortions in water, are therefore continuously

changing. This could be observed as a rapid re-orientation of the transition dipole moments concomitant with excitation redistribution between the two asymmetric stretch modes, revealing a 300 fs time constant. Key to the experimental distinction between mode re-orientation and simple excitation hopping was the large splitting of the asymmetric stretch band and clear distinction between diagonal and cross peak signals in the 2D-IR spectra. Non-linear spectroscopy can thus clearly identify the characteristic time for hydrogen bond fluctuations and – via the anisotropy decay of the pump-probe data – the persistence time of ion-water hydrogen bonds, which is approximately five times longer. As both time scales are very similar to those reported for isotope-labeled water, we inferred that the water molecules surrounding the nitrate ion do not form a very stable hydration shell. As a consequence, the theoretical description of NO_3^- (aq) is expected to depend critically on the quality of the water model, and we found indeed that the structural details in the vicinity of the ion differ between our CPMD results and previous MD and QM/MM approaches. However, the rapid reorientation of the vibrational dipole transition moments caused by symmetry breaking fluctuations in the hydrogen bond network around the nitrate ion is a general finding not specific to the CPMD simulation model but also present for the classical force field. The CPMD simulations show that the reorientation of the nitrate anion is faster than that of the water molecules, which is an interesting result, since earlier IR studies have shown that the reorientation of the water molecules are slowed down in the vicinity of anions.^{1, 3, 12, 52} Insight from this study shows how we can reconcile these data. In a weakly structured solvation shell, a small anion can reorient without dragging its first solvation shell with it. Hence, the anion will rotate while the water molecules in the cage around it will remain influenced by the electrostatic field of the ion. A speculative mechanism would be that water molecules temporarily reorient when the hydrogen bond to the anion breaks, but that there

is a high probability for reforming a hydrogen bond to the same anion, although to a different oxygen atom. We will in future studies perform detailed analysis on prolonged simulations to test this hypothesis. In this context a comparison of the CPMD with other simulation models such as *ab initio* QM/MM approaches and classical polarizable force fields will be interesting. Here we have used theory primarily to interpret our experimental observables, but we did not attempt to compute spectra or correlation functions for a quantitative comparison. Given the high sensitivity of the asymmetric stretch vibrations to solvation despite the relatively weak molecular interactions, this may be a critical albeit difficult test for our ability to describe and understand the very details of hydration.

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Acknowledgments:

M. Odelius thanks the Swedish Research Council, Carl Tryggers Foundation and Magnus Bergvall Foundation for financial support. The MD simulations were made possible through

generous allocations of computer time through SNIC at the Swedish National Supercomputer Center (NSC) and High Performance Computing Center North (HPC2N), Sweden. S. R. Keiding thanks the Danish Council for Independent Research and the Carlsberg Foundation for financial support.

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Figure captions:

Figure 1. Infrared absorption spectrum of NO_3^- (aq) (black line). Green and red lines represent a fit with two Gaussian functions. The arrows in the molecular pictures to the right indicate the atomic displacement vectors for the degenerate asymmetric stretch vibrations.

Figure 2. Calculated frequency splitting of the asymmetric stretch vibrations of nitrate, induced by the binding to the solvent molecule. The calculations are done using Gaussian09 at the level MP2/G-311+G(d,p) on the reaction $\text{NO}_3^- + \text{X} \rightarrow [\text{NO}_3\text{X}]^-$.

Figure 3. 2D-IR spectra of NO_3^- (aq) at different waiting times $t_2=0$ fs, 100 fs, 300 fs and 500 fs (from top to bottom). First and second columns show, respectively, spectra recorded with parallel (ΔA_{\parallel}) and perpendicular (ΔA_{\perp}) polarization of pump and probe pulses. The third column represents the magic angle signals ($\Delta A_{\text{MA}} = \Delta A_{\parallel} + 2\Delta A_{\perp}$). In the right column the diagonal signals are suppressed to highlight the cross peaks in the weighted difference spectra ($\Delta A_{\text{WD}} = \kappa \Delta A_{\perp} - \Delta A_{\parallel}$) with $\kappa = 3$ (0 fs), 2.48 fs (100 fs), 1.8 (300 fs), 1.6 (500 fs). The vertical axis is the excitation frequency, while the horizontal axis is the probe pulse frequency. Negative signals (bleach and stimulated emission) are blue, while positive signals (excited state absorption) are red. All spectra are normalized to the minimum negative signal for better comparison. For better orientation, the corners of the grey squares in the first row show the centers of the diagonal and off-diagonal signals. Black lines in the last row indicate the tilt (nodal line between bleach and excited state absorption) of the bottom right cross peak at short delays. The green and purple rectangles are the spectral regions used to calculate the signal of Fig. 4c.

Figure 4. a) Magic angle signal of the lower frequency diagonal peak at $\omega_1 = 1340 \text{ cm}^{-1}$ and $\omega_3 = 1320 \text{ cm}^{-1}$ (red circles, excited state absorption component) and 1360 cm^{-1} (blue squares, ground

state bleach/stimulated emission component). Lines show single exponential fits with a time constant of 550 fs. b) Anisotropy decay of the lower frequency diagonal peak (at $\omega_1=1340\text{ cm}^{-1}$, $\omega_3=1320\text{ cm}^{-1}$) with exponential fit with a 300 fs time constant. c) Cross peak growth given by $(s_1-s_2)/s_2$, where s_1 and s_2 are the magic angle 2D-IR signals integrated in the spectral regions marked by the green and purple rectangles in Fig. 3. The solid line is an exponential fit with a 300 fs time constant.

Figure. 5. Contour plot of the time dependent change in the infrared absorption $\Delta A(\nu, t)$ of the aqueous nitrate solution induced by the 200 nm pump pulses. Blue colors indicate negative changes in the induced absorption, while red and yellow colors represent positive absorption changes. The data are recorded with the pump beam polarization parallel to the probe polarization. The dashed lines indicate the two local absorption minima associated with ground state nitrate.

Figure. 6. The spectra of the three dominating components from the SVD analysis of the measured transient absorption dynamics presented in Fig. 5. Note the different scales on the vertical axes. The component depicted in Fig. a) is an order of magnitude large than the components in b) and c). The two curves in each figure represent the data obtained with the pump beam polarization parallel (black) and perpendicular (red) to the probe beam polarization. Only the curves in a) exhibit anisotropy. d) compares the dominant a) component (dots) to the FTIR spectrum of the steady-state spectrum of ground state nitrate measured by FTIR (line).

Figure. 7 a) The rotational anisotropy, $R(t)$, derived from the SVD analysis of the UV pump – IR probe transient absorption data. b) The rotational anisotropy measured at 195 nm when aqueous nitrate is excited at 200 nm in the UV pump – UV probe measurements. Both data sets are fitted

with a single exponential function with a time constant of 1.7 ± 0.3 ps convoluted with the experimental response time (line).

Figure 8 The radial distribution functions for the solvation structure around nitrate. Results from the classical MD simulations with either 55 (green lines) or 550 (dashed lines) water molecules are compared with results from CPMD simulations at 300 K (blue lines) and 350 K (red lines).

Figure 9. The reorientational motion of nitrate (solid lines) and water (dashed lines) is presented in terms of the time correlation function (TFCs) for the 2nd order Legendre polynomial, $P_2(u)$, of the N-O vector and the O-H vector in water. Results from the classical MD simulation (green lines) are compared with results from CPMD simulations at 300 K (blue lines) and 350 K (red lines).

Figure 10 a) The calculated absorption spectrum of the asymmetric stretch extracted from the MD simulations. The calculated spectrum is compared to the experimentally determined absorption spectrum of Fig. 1. Frequency distributions are displayed as bar diagrams, without considering transition strengths. A simulated spectrum based on classical MD is also included. The graphs in b) show the instantaneous frequencies as a function of time.

Figure 11. a) Time-dependent orientation of the asymmetric stretch transition dipole moments (modeled as dipole derivatives of the vibrational modes) relative to an N-O bond direction in a molecule-fixed frame. Red and blue colors indicate lower and higher frequency transitions. b) The frequency fluctuations in the asymmetric stretching modes presented in Fig. 10 b) are plotted again to highlight the correlation with the reorientational dynamics in Fig. 11 a).

Figure 12. a-d) A schematic representation of different solvated ion configurations. Transitions between these configurations caused by solvation shell fluctuations can explain the observed cross peak growth and simultaneous decay of the diagonal peak anisotropy due to reorientation of their transition dipole moments. Dipole moment orientations are indicated by red and blue arrows which correspond to lower and higher frequency transitions, respectively. e) Transition dipole orientations of the lower frequency asymmetric stretch transition from our sampling of the CPMD simulation in the molecular frame after aligning all molecules along the shortest N-O bond, pointing in the x -direction. f) Histogram of the angle distribution.

Figures:

Fig. 1

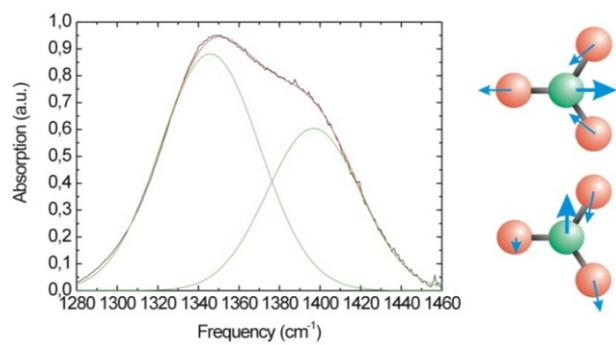


Fig. 2

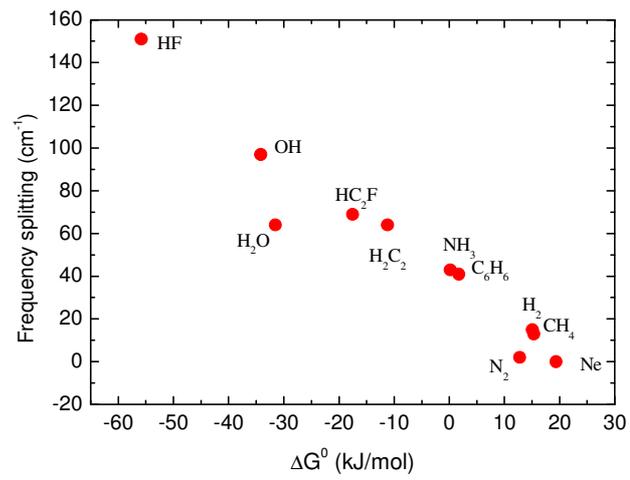


Fig. 3

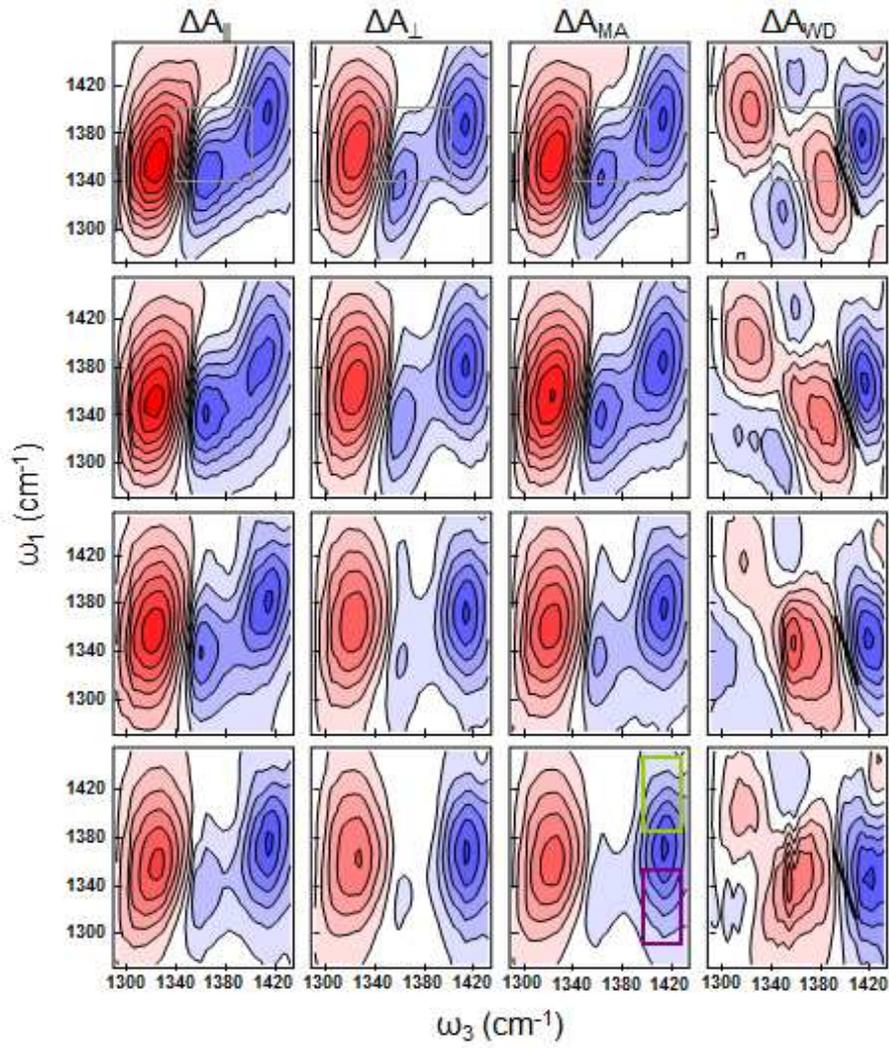


Fig. 4

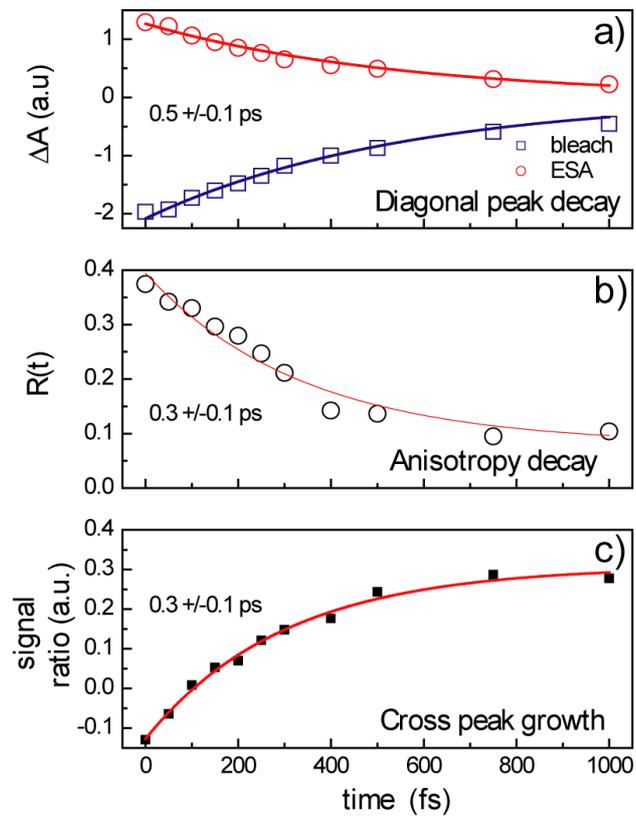


Fig. 5

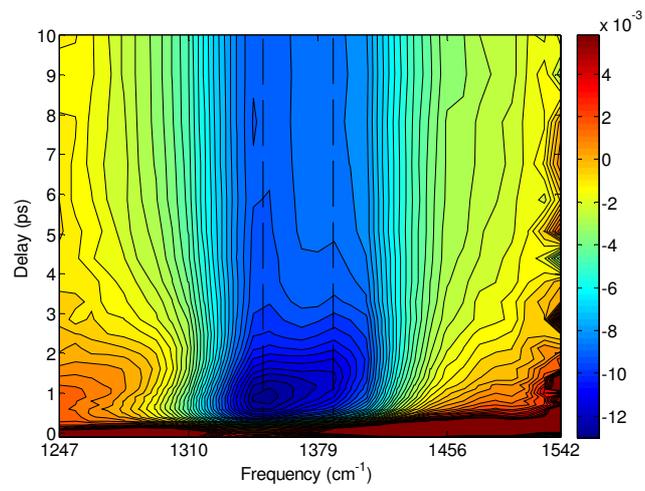


Fig. 6

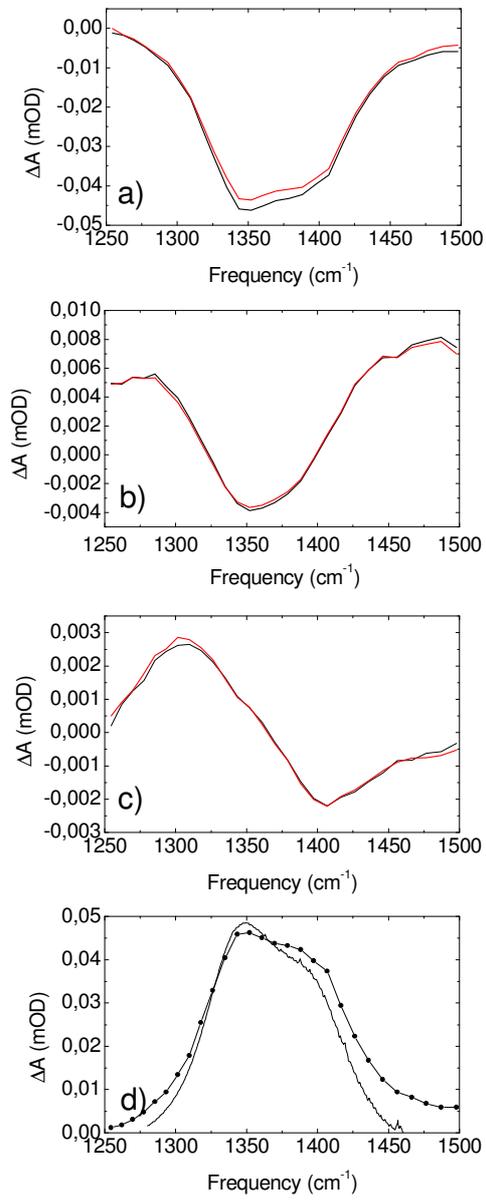


Fig. 7.

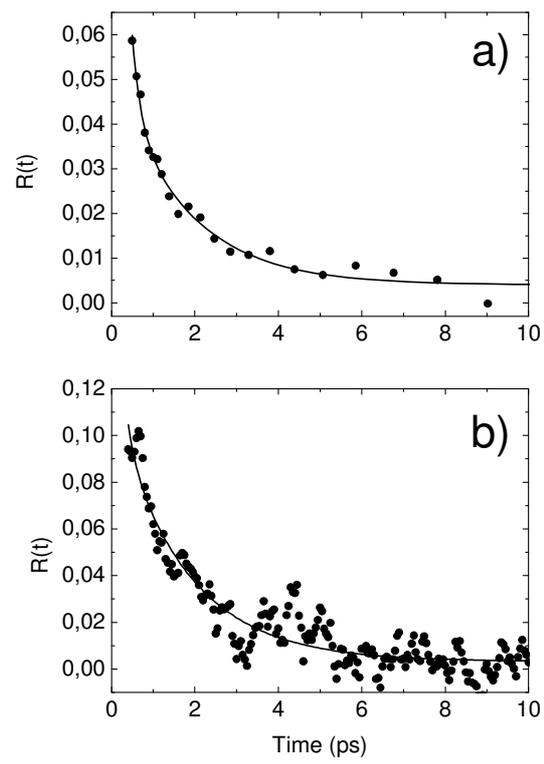


Fig. 8

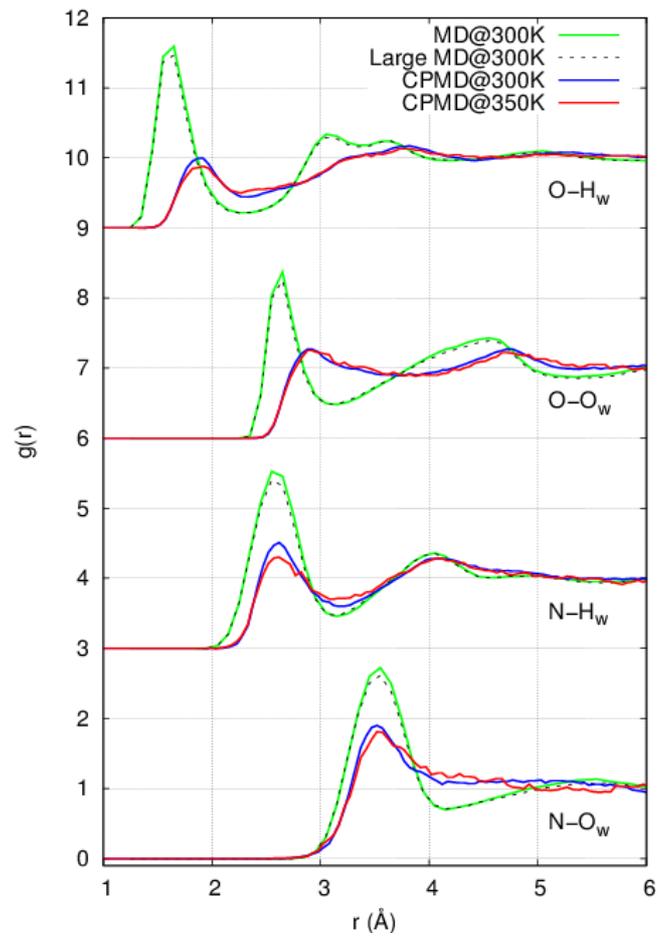


Fig. 9

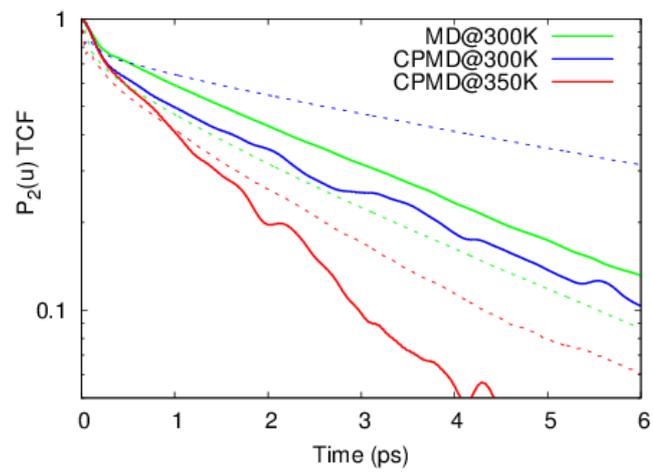


Fig. 10

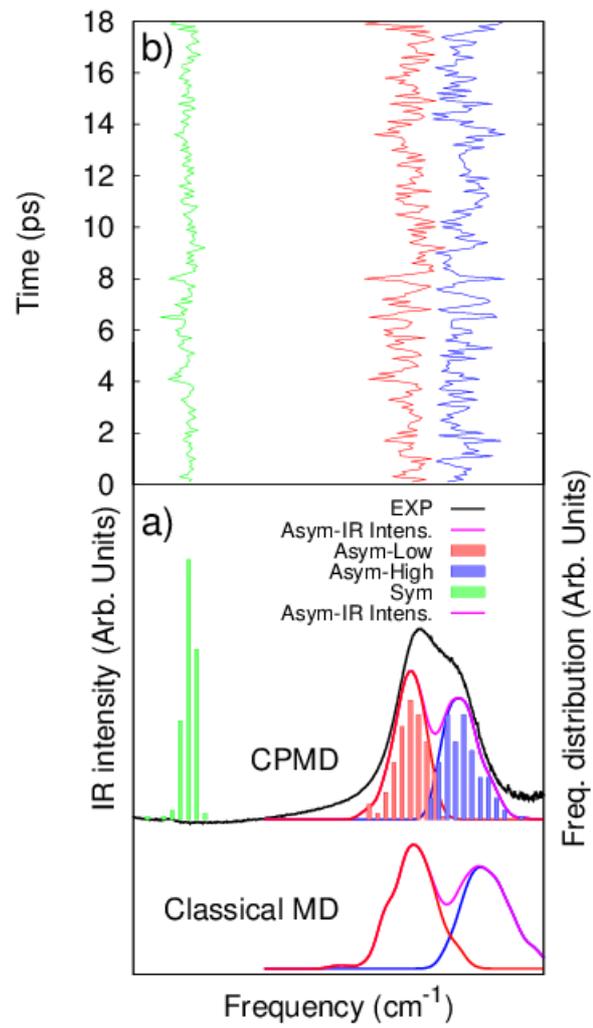


Fig. 11

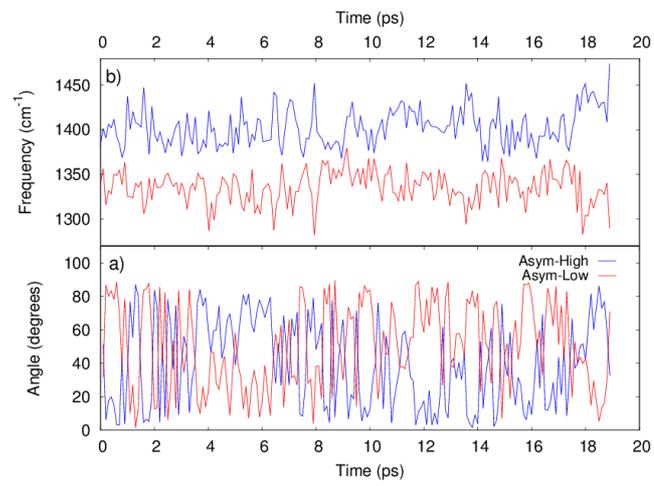
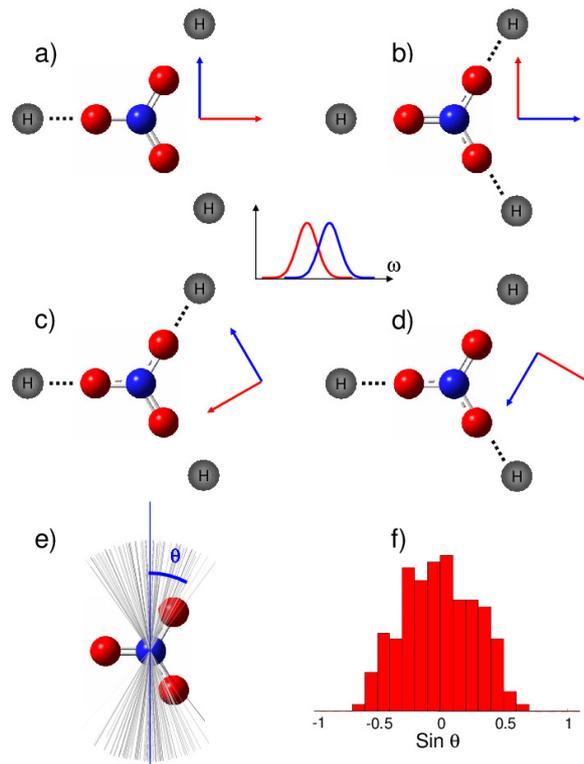


Fig. 12.



TOC graphics

