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Density functional theory based molecular dynamics study of solution composition effects on the solvation shell of metal ions

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We present an ab initio molecular dynamics study of the alkali metal ions Li+, Na+, K+ and Cs+, and of the alkaline earth metal ions Mg²⁺ and Ca²⁺ in both pure water and electrolyte solutions containing the counterions Cl⁻ and SO₄^{2−}. Simulations were conducted using different density functional theory methods (PBE, BLYP and revPBE), with and without the inclusion of dispersion interactions (-D3). Analysis of the ion-water structure and interaction strength, water exchange between the first and second hydration shell, and hydrogen bond network and low-frequency reorientation dynamics around the metal ions have been used to characterise the influence of solution composition on the ionic solvation shell. Counterions affect the properties of the hydration shell not only when they are directly coordinate to the metal ion, but also when they are at the second coordination shell. Chlorine ions reduce the sodium hydration shell and expand the calcium hydration shell by stabilizing under-coordinated hydrated $Na(H_2O)_5^+$ complexes and over-coordinated $Ca(H_2O)_7^{2+}$. The same behaviour is observed in CaSO₄(aq), where Ca²⁺ and SO₄²⁻ form almost exclusively solvent-shared ion pairs. Water exchange between the first and second hydration shell around Ca2+ in CaSO4(aq) is drastically decelerated compared with the simulations of the hydrated metal ion (single Ca2+, no counterions). Velocity autocorrelation function analysis, used to probe the strength of the local ion-water interaction, shows a smoother decay of Mg2+ in MgCl2(aq), which is a clear indication of a looser inter-hexahedral vibration in the presence of chlorine ions located in the second coordination shell of Mg2+. The hydrogen bond statistics and orientational dynamics in the ionic solvation shell show that the influence to the water-water network cannot only be ascribed to the specific cation-water interaction, but also to the subtle interplay between the level of hydration of the ions, and the interactions between ions, especially those of opposite charge. As many reactive processes involving solvated metal ions occur in environments that are far from pure water but rich in ions, this computational study shows how the solution composition can result in significant differences in behaviour and function of the ionic solvation

Introduction

A fundamental understanding of the processes controlling the hydration of ions is of importance to low-temperature geochemistry and biological systems. For example, in aqueous environments the processes of nucleation and growth of ionic crystals such as calcite (CaCO₃) or magnesite (MgCO₃), are controlled by the dynamics of hydration—dehydration around the cation.^{1–3} In the hydration model proposed by Frank and

Wen to explain deviations of ionic solution behaviour relative to pure water, the insertion of an ion in solution radially aligns water molecules in concentric regions characterised by different levels of interaction between water and ion.⁴ The water molecules in the innermost region, also known as the first solvation shell (FSS), are tightly bound to the centric ion and exhibit translational and rotational entropy loss compared with pure water.⁵ Several molecular dynamics (MD) studies of hydrated metal ions (single cation, no counterions) have been conducted to characterize the structure and dynamics of the ionic FSS, with the results confirming what anticipated by the Frank and Wen hydration model.⁶⁻⁹

lons in aqueous electrolyte solutions are present as free ions only in dilute solutions. Otherwise, cations and anions form contact or solvent-shared ion pairs. The structure and dynamics of the ionic solvation shell can, therefore, be significantly affected by the subtle interplay between the level of hydration of the ions, and the interactions between ions, especially those of opposite charge. In more general terms, the composition of the solution, including the balance between ion-solvent and ion-ion interactions, could result in significant

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d. School of Environmental Sciences, University of East Anglia, Norwich, NR4 7TJ, UK Electronic Supplementary Information (ESI) available: details of simulated systems, additional analysis of MD simulations (ion pairing, dynamics of the hydration shell, velocity autocorrelation function, hydrogen bond statistics, protocol for the calculation of time correlation functions). See DOI: 10.1039/x0xx00000x

differences in behaviour and function of the FSS. Consequently, an accurate characterization of the effect of solution composition on the structure and dynamics of the ionic solvation shell is important to understand and control reactive processes involving ions in solution.

The MD simulation technique has been extensively used to gain insights into the elusive molecular-level processes controlling the properties of aqueous electrolyte solutions. 11,12 However, the force field used to describe the ion-water interaction is crucial to obtain an accurate characterization of structure and dynamics in electrolyte solutions. 13,14 An important contribution to this field was the development of ab initio MD (AIMD), where forces are derived from the electronic structure, 15,16 usually in the framework of density functional theory (DFT),17 which provides the capability of studying nonadditivity effects in the dynamics of ions solvation shells. AIMD simulations, using the Car-Parrinello and Born-Oppenheimer schemes, of cations 18-23 and anions 24,25, have been mostly conducted on an isolated ion in pure liquid water, with no counterions in solution. Electrolyte solutions, on the other hand, have been subject to only a few AIMD studies, including the characterization of the dissociation of the NaCl in water,²⁶ the cooperative ionic effects are of the Na⁺ and Cl⁻ on the hydrogen bonding network,²⁷ and the influence of a static electric fields.28

Here, we report an AIMD study of the of the alkali metal ions Li⁺, Na⁺, K⁺ and Cs⁺, and of the alkaline earth metal ions Mg²⁺ and Ca²⁺, in pure water and electrolyte solutions containing the counterions Cl⁻ and SO₄²⁻, in order to characterize composition and concentration effects on the structure and dynamics of the ionic solvation shell. We have chosen the above combination of cations and anions because they are commonly found in nature, including in the composition of the sea and groundwater.²⁹ Salt concentrations ranging from 0.9 to 3.8 mol.kg⁻¹ were considered to quantify the effect of solution composition on the strength of ion—water interaction and to characterise the structure and low-frequency dynamics of the FSS.

After having introduced details of the simulations, we next report on the effect of the solution composition on the structure and dynamics of the ionic solvation shell. These results are rationalised in terms of the mechanism of water exchange around cations and the strength of ion—water interaction. The distribution of hydrogen bonds and the reorientation dynamics of water molecules, in the bulk solution and the first solvation shell, are finally presented.

Computational details

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Ab initio (Born-Oppenheimer) MD simulations were conducted with the electronic structure code CP2K/Quickstep code (4.1).³⁰ CP2K implements DFT based on a hybrid Gaussian plane wave. The PBE³¹ generalized-gradient approximations for the exchange and correlation terms were used, together with the Grimme's –D3 dispersion correction, to provide a more accurate description of the structure of liquid water.^{32,33} Simulations were also conducted using the BLYP^{34,35} and

revPBE³⁶ functionals, with and without the -D3 terms. The Goedecker-Teter-Hutter (GTH) pseudopotentials were used to describe the core–valence interactions. All atomic species were represented using a double-zeta valence polarized (DZVP) basis set. The plane wave kinetic energy cut off ($E_{\rm cut}$) was set to 1000 Ry. The k-sampling was restricted to the Γ point of the Brillouin zone. The tolerance for the wave function optimization was set to 10^{-6} au that allows for a 1 fs time steps with reasonable energy conservation. AIMD simulations were carried out in the canonical (NVT) ensemble (constant number of particles N, volume V, temperature T) using a Nosé-Hoover chain thermostat to maintain the average temperature at 300 K. Periodic boundary conditions were applied throughout.

Simulation protocol. First, we have conducted classical MD simulation of 64 water molecules in the isothermal-isobaric (NPT) ensemble (constant number of particles N, pressure P = 1 atm, temperature T = 300 K) for 1 ns to generate an equilibrated aqueous solution. The last configuration was used to generate solutions of hydrated metal (M) ions (single M, no counterions, by replacing one H₂O with one ion (M: Li⁺, Na⁺, K⁺, Mg²⁺, and Ca²⁺). Aqueous electrolyte solutions with concentrations ranging from 0.9 to 3.8 mol.kg⁻¹ were generated by randomly replacing water molecules with cations, M, and anions, X (Cl⁻ or SO₄²⁻) ions, by making sure that the initial configuration did not contain contact ion pairs. Classical MD simulations used the pairwise Lennard-Jones (LJ) potential model developed by Zeron et al.38 (denoted as Madrid-2019 forcefield) for Li+, Na+, K+, Mg2+, Ca2+, Cl-, and SO₄²⁻ in aqueous solution.³⁸ In Madrid-2019, the water molecules are represented by the TIP4P/2005 model,³⁹ and the monovalent and divalent ions are modelled using charges of 0.85 and 1.7, respectively (in electron units). This potential model allows a very accurate description of the densities of the solutions up to high concentrations.³⁸ The Madrid-2019 has not been developed for Cs+ and to generate equilibrated solutions of CsCl we used the LJ potential model from the Amber database.40 For each system, we conducted 6 ns of classical MD (NPT, T = 300 K and P = 1 atm) to equilibrate the cell volume and the last configuration was used as the starting point of 50 ps of AIMD. Details of the solutions considered in this study (number of ions and water molecules, cell lengths of the NVT simulations, solution concentrations) are reported in Table S1 of Electronic Supporting Information (ESI).41 For comparison purposes, we have also conducted AIMD simulations of pure liquid water (64 H₂O, 1 g cm⁻³) and of Cl⁻ in 63 H₂O molecules. The procedure of first conducting classical MD (NPT) simulations to equilibrate the volume of the simulation box followed by AIMD (NVT) simulations has been routinely adopted to study the properties of hydrated ions. 18,20,22,42 We did not attempt to run AIMD in the NPT ensemble, which could determine the "correct" system density at zero pressure, because depending on the DFT method (functional and dispersion correction) equilibrium density of water can change significantly.⁴³ Instead, we chose to fix the system size to match the one obtained using the classical MD simulation of these systems. NPT trajectories must be long enough to assure statistical convergence, and the convergence This article is licensed under a Creative Commons Attribution 3.0 Unported Licence.

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of pressure for DFT plane wave MD simulations in the NPT ensemble requires a significantly higher basis set cutoff due to grid sensitivity issues when the volume of the simulation cell can fluctuate. 44,45

Results and discussion

Solvation structure

Ion-water radial distribution function. Information regarding the structural properties of the coordination sphere of the alkali metal ions (M = Li⁺, Na⁺, K⁺ and Cs⁺) and alkaline earth metal ions ($M = Mg^{2+}$ and Ca^{2+}) in pure liquid water and in the presence of chlorine (Cl⁻) and sulphate (SO₄²⁻) counterions, can be deducted from the radial distribution functions (RDFs) of the M-O pairs, $g_{MO}(r)$, in **Fig. 1**, which represent the probability, relative to a random distribution, of finding a water oxygen atom at a distance r from the metal ion. Comparison of the hydration structure of the cations (isolated ion, no counterions) with the results obtained from other AIMD simulations and experimental measurements (Table S2 of ESI) shows that the PBE-D3 functional with the hybrid Gaussian (DZVP) plane wave (E_{cut} = 1000 Ry) basis set gives an accurate description of the position and amplitude of the M-O RDF first peak as well as the average number of water molecules in the first hydration shell. For example, our results for Na⁺ and K⁺ are in good agreement with recent AIMD simulations conducted with the strongly constrained and appropriately normed (SCAN) meta-GGA functional for complex condensed phase systems.¹⁹ A comparison of the ionwater RDFs obtained from AIMD (PBE-D3) and classical MD (Madrid-2019) are also reported in Fig. S1 of ESI. Fig. 1 shows that the average metal-water distance of the first coordination shell increases when moving down the alkali and alkali earth group. The lithium, magnesium, calcium and, to a lesser extent, sodium ions are characterized by a more rigid and well defined first solvation structure. As to the hydration structure of the K⁺ and Cs⁺, the intensities of the first peaks are lower and their profiles are much less defined. In aqueous electrolyte solutions, the presence of the counterions in solution increases the intensity and decreases the full-widthhalf-maximum of the first peak of Li⁺ and Mg²⁺; no significant effect is observed for the $g_{MO}(r)$ profiles of Na⁺ and Ca²⁺. On the other hand, the chlorine ions broaden the RDFs of the K+ and Cs+ cations, indicating a perturbation of their coordination shells and a more labile hydration structure.

Ion-water coordination shell distribution and ion pairing. We report in Fig. 2 the probability distribution of the number of water molecules in the first coordination shell of the metal ions, obtained from the analysis of the AIMD trajectories of hydrated ions and of the 1.9 kg mol⁻¹ aqueous electrolyte solutions. For the hydrated ions the average size of the FSS, as well as the number of accessible hydration states, increases as we move down the periodic table. The tetra-hydrated lithium complex [Li(H2O)4]+, and the hexahydrated magnesium and calcium complexes $[Mg(H_2O)_6]^{2+}$ and $[Ca(H_2O)_6]^{2+}$ were the only species detected for these ions. On the other hand,

hydrated Na⁺, K⁺ and Cs⁺ can access multiple hydration states: the caesium ion, for example, can coordinate of the coordinate of and nine water molecules. In aqueous electrolyte solutions, metal ions are not purely hydrated, but they tend to form

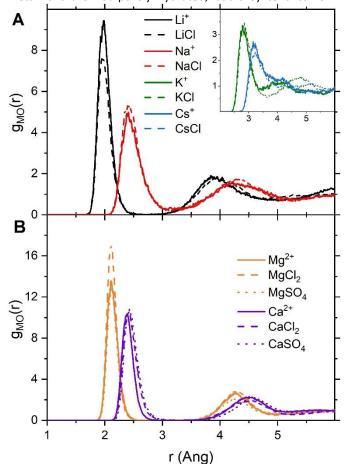


Figure 1. Ion-water radial distribution functions, $g_{MO}(r)$, obtained from AIMD (PBE-D3) simulations of hydrated metal ions (single Mn+, no counterions) and of aqueous electrolyte solution (1.9 mol.kg⁻¹). (A) Solutions containing Li⁺, Na⁺, K⁺, Cs⁺. (B) Solutions containing Mg²⁺ and Ca²⁺.

contact ion pairs (CIPs), solvent-shared ion pairs (SSHIPs), and solvent-separated ion pairs (SSIPs) with the counterions in solution. The percentages of CIPs, SSHIPs and SSIPs in the 1.9 mol.kg solutions are listed in Table 1, while the results for the more dilute (0.9 mol.kg⁻¹) and concentrated (3.8 kg.mol⁻¹) solutions are reported in Table S3 of ESI. For the 1.9 mol.kg solutions, Cs+ has a higher tendency of forming CIPs with Cl-(~73%) than Na+ and K+, which prefer solvent-shared or solvated-separated ions pairing. The caesium-chloride RDFs (Figure S2 of ESI) shows that the presence of an intense peak centred at 3.6 Å, which is roughly the sum of the ionic radii of Cs⁺ and Cl⁻ in water.⁴⁶ AIMD simulations of a larger simulation box, containing 8 Cs+ and 8 Cl- in 710 water molecules (17 ps), show similar ion pairing (80.0% CIP, 16.7% SSHIP and 3.3% SSIP) and Cs-Cl RDF (3.6 Å) results (Figure S2 of ESI). A recent EXAFS study of Cs-Cl ion pairing in dilute and concentrated caesium chloride solutions reported Cs-Cl distances of approximately 3.5 Å,47 confirming the conclusions obtained in this study. For 1.9 mol kg-1 NaCl(aq) and KCl(aq), the metalchloride RDFs have a very broad profile between 3 and 4 Å,

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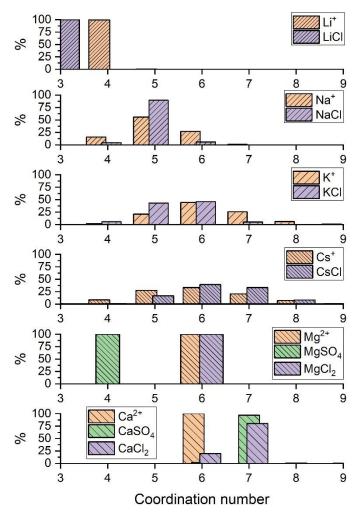


Figure 2. Probability distributions of the number of water molecules in the first coordination shells of the metal ions Li⁺, Na⁺, K⁺, Cs⁺, Mg²⁺ and Ca²⁺. Results obtained from the analysis of AIMD (PBE-D3) trajectories of hydrated metal ions (single Mⁿ⁺, no counterions) and of aqueous electrolyte solution (1.9 mol.kg⁻¹).

confirming the tendency to form separated ion pairs. For KCl(aq), CIPs are observed in the more dilute 0.9 mol kg-1 solution (36% CIP). In the most concentrated solutions, 3.8 mol kg⁻¹ both Na⁺ and K⁺ form CIPs, 43.9 % in NaCl(aq) and 56% in KCl(aq). The tendency, reported in this study, of Cs+ to form CIPs with Cl⁻ contrasts with classical MD simulations of M-Cl (M = Na, K, and Cs) in solution, which reported dissociation rate constants (k) following the trend $k_{NaCl} < k_{CsCl}.^{48}$ One source of this incongruency could be the accuracy of the interatomic potential, since calculations of the dissociation of NaCl in water from AIMD simulations reported a remarkable difference in the free energy profiles between the ab initio and classical potentials. 49,50 This stresses the importance of the inclusion of many-body effects in the interaction potentials used to simulate electrolyte solutions.¹² Other factors could also affect the population of contact and solvent shared IPs other than the absolute strength of the ion-chloride interaction such as the relative dynamics of exchange of water and chlorine around the ions. The analysis of the number of ligand exchange around Na+, K+ and Cs+ ions shows, in fact, a competitive exchange between the Cl⁻ and H₂O, whereas for

K⁺ and Cs⁺ the frequency of water exchange is visignificantly faster than chlorine (**Table S4** in ESI). DOI: 10.1039/D0CP01957G

For LiCl(aq), the analysis of the ion-water coordination shell distribution shows that in the presence of Cl-, the number of water molecules coordinated to Li* reduces from four to three as a result of the formation of Li*···Cl contact ion pairs (Table 1). In 1.9 mol.kg⁻¹ NaCl(aq), Na⁺ and Cl⁻ form solvent-shared or solvent-separated IP+. Despite not being coordinated to Na+, chlorine ions induce a significant contraction of the sodium hydration shell: in NaCl(aq) the sodium ions are in the fivecoordinated state for 90% of the simulation period, compared to approximately 50% of the simulation period of hydrated Na⁺ (isolated Na⁺, no counterions). Consequently, counterions such as Cl⁻ can stabilize under-coordinated metal-water complexes even when they are not part of the first coordination shell of Na⁺. The Cs⁺ and Cl⁻ ions are, for most of the simulation period, present as CIPs (72%) but due to the high flexibility of caesium first hydration shell, the coordination number distribution is only marginally affected by the presence of chlorine ions (Fig. 2). In MgCl₂(aq) the hexahydrated [Mg(H₂O)₆]²⁺ is the only hydrated complex present in solution, and in MgSO₄(aq) only four water molecules are directly coordinated to Mg²⁺ because the magnesium and sulphate ions form very stable contact ion pairs (Table 1). Static DFT calculations of the free energy of formation (ΔG) of the Mg²⁺/SO₄²⁻ and Mg²⁺/Cl⁻ contact ion pairs confirm the much higher propensity of SO₄²⁻ to form stable CIPs with Mg²⁺ compared with Cl⁻: Mg(H₂O)₆²⁺ + SO₄²⁻ \rightarrow MgSO₄(H₂O)₄ + 2H₂O, Δ G = −160 kJ.mol⁻¹; (Mg(H₂O)₆²⁺ + Cl⁻ → $MgCl(H_2O)_5^+ + H_2O$, $\Delta G = -25 \text{ kJ.mol}^{-1} \text{ (wB97XD/aug-cc-pdvz)}$ level of theory in the CPCM continuum solvation model). The coordination shell distribution of the calcium ion displays a counterintuitive behaviour, with the coordination number (CN) of Ca²⁺ increasing from six in pure water to around seven in CaCl₂(aq) (Fig. 2). The chlorine ions are hardly ever coordinated to the calcium ions, with which they predominantly form SSHIPs (Table 1). Therefore, the chlorine ions located in the second coordination shell of Ca2+ stabilize the seven-coordinated $Ca(H_2O)_7^{2+}$ complex over the sixcoordinated Ca(H₂O)₆²⁺ species. The same behaviour is observed in CaSO₄(aq), where Ca²⁺ and SO₄²⁻ form almost exclusively SSHIPs (99.6%) and the epta-hydrated calcium complexes are stabilised by the sulphate ions located in the second hydration shell of metal ion.

The average CN of for the first hydration shell of Ca^{2+} has been still a controversial issue for AIMD simulations.⁵¹ Several DFT based MD simulations, using the Car-Parrinello and Born-Oppenheimer schemes, of the hydrated calcium ion (single Ca^{2+} , no counterions) reported six water molecules in the first hydration shell of Ca^{2+} ,⁵² which agrees our result (**Fig. 2**). However, experimental results from extended X-ray absorption fine structure (EXAFS) measurements in $CaCl_2$ solutions, with concentrations ranging from 0.12 to 6 mol.kg¹, resulted in CNs between 6.8 and $8.^{53,54}$ As previously discussed, the presence of counterions (Cl⁻ and SO_4^{2-}), which are normally present in experiments, stabilizes the higher coordination states of Ca^{2+} , yielding a computed CN of Ca^{2+} in $CaCl_2(aq)$ and $CaSO_4(aq)$ (6.9) in much better agreement with

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EXAFS measurements of CaCl₂ solutions. Therefore, one simple reason for the disagreement between theory and experiment could be the composition of the solution used to conduct the simulations. Other factors such as the size of the simulation box, the length of the simulation period, and the choice of the density functional method have also been considered by conducting AIMD simulations of hydrated Ca²⁺ (no counterions) in a box containing 63 and 124 water molecules using the PBE, rev-PBE and BLYP functionals with and without the -D3 dispersion correction (Table 2).

Table 1. The speciation of the cations M (Li⁺, Na⁺, K⁺, Cs⁺, Mg²⁺ and Ca²⁺) and anions X (CI-, SO₄²⁻) as a function of concentration: contact ion pair (CIP) when M and X are in direct physical contact; solvent-shared ion pairs (SSHIP) when M and X are separated by one water molecule; solvent-separated ion pairs (SSIP) when M and X are separated by at least two water molecules. Assignments made from the analysis of the M-X radial distribution functions (RDF): CIP if $r_{M-X} \le r_{M-X}^{min1}$; SSHIP if $r_{M-X}^{min1} < r_{M-X} \le r_{M-X}^{min2}$; SSIP if $r_{M-X} > r_{M-X}^{min2}$ where r_{M-X}^{min1} and r_{M-X}^{min2} are the positions of the first and second minima, respectively, of the M-X RDF. Results from AIMD (PBE-D3) simulations.

System	<i>b</i> (mol kg ⁻¹)	CIP (%)	SSHIP (%)	SSIP (%)
LiCl	1.9	85.6	14.4	0.0
NaCl	1.9	0.0	68.9	31.1
KCl	1.9	3.3	51.4	45.3
CsCl	1.9	72.8	27.2	0.0
$MgCl_2$	1.9	0.0	55.5	44.5
CaCl ₂	1.9	0.7	97.1	2.2
$MgSO_4$	1.9	100.0	0.0	0.0
$CaSO_4$	1.9	0.0	99.6	0.4

The results show that: a simulation period of 50 ps is sufficiently long to get a convergent distribution of the calcium-water coordination shell; Ca2+ in 63 and 124 water molecules yield CNs of 6.0 and 6.5; BLYP-D3 gives a CN of 6.9 and a coordination distribution that is shifted to higher coordination states compared the average CN obtained with PBE-D3 (6.0) and revPBE-D3 (6.1). Both the system size and the DFT method can, therefore, influence the coordination distribution and average coordination size. This analysis suggests that solution composition, system size, and level of theory should all be considered when comparing AIMD simulations and experimental determinations of ionic coordination numbers.

Table 2. Probability distributions of the number of water molecules Aincibe nfirst coordination shells of hydrated Ca²⁺ (no counterions) Poblained Cash Polifferent 5715 based AIMD methods. NH20/Ca2+ is the number of water molecules per calcium in the simulation cell; t_{sim} is the simulation period used to compute the coordination number (CN) distribution around Ca2+

Method	N _{H2O} / Ca ²⁺	t _{sim} (ps)				
			% 6	% 7	% 8	Avg.
PBE	63	50	100.0	0.0	0.0	6.0
PBE-D3	63	50	95.1	4.9	0.0	6.0
		100	96.6	3.4	0.0	6.0
		500	98.7	1.3	0.0	6.0
	124	200	47.6	51.6	0.8	6.5
revPBE	63	50	38.2	61.7	0.1	6.6
revPBE-D3	63	50	86.9	12.1	1.0	6.1
BLYP	63	50	32.7	17.3	0.0	6.3
BLYP-D3	63	500	15.0	80.7	4.3	6.9

Dynamics of the ionic solvation shell

The frequency of water exchange around cations is generally considered as the rate determining for the reactions involving these ions in aqueous solutions.55-58 We have characterised the dynamics of the first and second hydration shells of the alkali metal ions (Li+, Na+, K+ and Cs+) and alkaline earth metal ions (Mg²⁺ and Ca²⁺) in pure liquid water and electrolyte solutions using the "direct" method proposed by Hofer and coworkers.59 This methodology has been previously applied for, among others, the characterization of ligand exchange between coordination shells of hydrated alkaline earth metal ions and their carbonate and bicarbonate complexes, 21,22,60 as well as the quantification of the water exchange frequency around calcium sites in calcite-water interface 61,62 and hydroxyapatite nanopores. 63 In the "direct" method, the MD trajectories are analysed for water molecule movements and whenever a water molecule crosses the boundary of the cation coordination shell its path is followed; if its new position outside or inside this shell lasts for more than $\tau^* = 0.5$ ps then the event is accounted as a real exchange event (Nex). The choice of $\tau^* = 0.5$ ps is based on the average lifetime of a hydrogen bond between the solvent molecules, obtained using femtosecond spectroscopy,64 and is the standard value used in the applications of the direct method; setting τ^* to 5 ps gives only small differences to the statistics of water exchange (Table S5 of ESI). From the number of water exchange events, the MRT of the water molecules in the first and second hydration shell of the ion can be computed as MRT = (CN \times t_{sim})/ N_{ex} , where CN is the average number of water molecules in the first or second coordination shell and $t_{\rm sim}$ is the duration time of the simulation.⁴¹ A recent comparison by Wolthers and co-workers of the direct and survival function methods to characterize the calcium dehydration frequencies from MD trajectories,²² concluded that the direct methods could provide shorter mean residence times than SF, but requires shorter simulation times than SF to obtain statistically meaningful output, and is better suited to analyse shorter trajectories generated by ab initio MD simulations.

Table 3 reports the values of N_{ex} in the first and second coordination shell of the metal ions, together with the number of exchange events normalised to 10 ps, $N_{\text{ex}}/10$ ps, which

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represents a measure of the "lability" of the hydration shell of the metal. For the alkali metal ions, the first shell solvent exchange dynamics increases substantially on going from Li⁺ (MRT=11.8 ps) to the other alkali metal ions Na⁺ (1.1 ps), K⁺ (0.4 ps) and Cs⁺ (0.2 ps). The MRT of solvent around H₂O and Cl⁻ is approximately 0.5 ps, which is comparable to the values obtained for K⁺ and Cs⁺, confirming the lability of these ions. As reported in **Table 3**, AlMD simulations of hydrated Mg²⁺ (single ion, no counterions) showed no exchanges around Mg²⁺, compared with compared to ~50 of such exchanges (more facile dehydration) in 500 ps around aqueous Ca²⁺ (61.2 ps). The reactivity of the cations varies, therefore, as follow: Mg²⁺ << Ca²⁺ < Li⁺ << Na⁺ < K⁺ ≈ Cs⁺. We have rationalized this behaviour by conducting classical metadynamics MD simulations to compute the free energy as a function of the

metal—water coordination number (CN) for Li⁺, Na⁺_{cw} Mg²⁺_{ce} and Ca²⁺, from which thermodynamic information of 19 the accessible coordination states for these cations can be extracted (**Fig. S3** in ESI). Ca²⁺, Li⁺ and Na⁺ have several coordination states corresponding to a minimum on the free energy profile and for Na⁺ the transition between these states is almost barrierless. On the other hand, the only coordination state accessible at room temperature to Mg²⁺ is six and the free energy barrier to go from the six to the five coordinate state (32 kJ mol⁻¹) is about three times higher than the free energy barrier of hydrated Ca²⁺ to interchange between the six and seven coordination environments. As Mg²⁺ is strongly hydrated, water exchange is drastically retarded in its FSS.

Table 3. Number of accounted water exchange events $(N_{ex}^{H_2O})$ in the first and second coordination shells of the metal ions Li⁺, Na⁺, K⁺, Cs⁺, Mg²⁺ and Ca²⁺, with a duration of more than $\tau^* = 0.5$ ps, obtained from the analysis of AIMD (PBE-D3) simulations of hydrated metal ions (single M, no counterions), aqueous electrolyte solution, pure water and hydrated chlorine ion. Values of $N_{ex}^{H_2O}$ normalized to the number of cations in solution. Mean residence time (MRT) of water molecules in the first hydration shell computed using the relation MRT = $(t_{\text{sim}} \times \text{CN})/N_{\text{ev}}$ where t_{sim} is the period of simulation (in ps) and CN is the average ion-water coordination number of the first and second coordination shell, which was obtained from the integration of the ion-water radial distribution function up to the first and second minimum, respectively. Concentration (b) in mol.kg⁻¹.

			First Shell					Second Shell		
System	b	$t_{\sf sim}$	CN	$N_{ex}^{H_2O}$	$N_{ex}^{H_2O}/10_{ps}$	MRT (ps)	CN	$N_{ex}^{H_2O}$	$N_{ex}^{H_2O}/10_{\ m ps}$	MRT (ps)
Li ⁺	0.9	50	4.0	17.0	3.4	11.8	15.3	2066	413.2	0.4
LiCl	1.9	50	3.0	7.5	1.5	20.0	13.9	3760	752.0	0.2
Na⁺	0.9	50	5.1	233.0	46.6	1.1	20.0	1852	370.4	0.5
NaCl	0.9	50	5.1	178.0	35.6	1.4	21.4	2022	404.4	0.5
	1.9	50	5.0	195.5	39.1	1.3	20.9	3152	630.4	0.3
	3.8	50	4.7	186.5	37.3	1.3	19.3	7205	1441.0	0.1
K ⁺	0.9	50	6.2	750.0	150.0	0.4	21.1	1675	335.0	0.6
KCl	0.9	50	4.8	536.0	107.2	0.4	21.9	1852	370.4	0.6
	1.9	50	5.5	693.5	138.7	0.4	20.4	3832	766.4	0.3
	3.9	50	3.9	694.0	138.8	0.3	19.5	6964	1392.8	0.1
Cs+	0.9	50	8.3	1299.0	259.8	0.3	21.5	1556	311.2	0.7
CsCl	1.9	50	6.3	983.0	196.6	0.3	25.0	3726	745.2	0.3
Mg ²⁺	0.9	40	6.0	-	_	_	20.0	1057	264.3	0.8
$MgCl_2$	1.9	50	6.0	_	_	_	18.9	1266	253.2	0.7
$MgSO_4$	1.9	50	4.5	_	-	-	14.8	3223	644.6	0.2
Ca ²⁺	0.9	500	6.0	49.0	1.0	61.2	21.0	15694	313.9	0.7
CaCl ₂	1.9	50	6.8	32.0	6.4	10.6	19.1	2429	485.8	0.4
CaSO ₄	1.9	50	6.9	_	_	_	21.0	3121	624.2	0.3
H ₂ O	-	50	3.9	536	107.3	0.4	23.0	2316	463.3	0.5
Cl-	0.9	40	6.2	502	125.5	0.5	26.4	1702	425.5	0.6

As to the dynamics of the second hydration shell, for all ions, the MRTs is less than 1 ps and close to the MRT of solvent around H₂O (0.5 ps). This result contrasts with previous values obtained from AIMD simulations of the alkaline earth metal ions Mg²⁺, Ca²⁺ and Sr²⁺, which reported MRTs between 3.3 and 6.8 ps. These simulations, however, used smaller simulation boxes than the one used in this study and did not employ dispersion corrected DFT functionals.²¹ Regarding the effect of counterions in solution to the dynamics of exchange around the cations, in LiCl(aq) the formation of Li+/Cl- CIPs increases significantly the MRT around Li* to 20 ps. The other, very labile, alkali ions Na+, K+ and Cs+ are substantially unaffected by the presence of counter-ions in solution and the dynamics of water exchange around Na+, K+ and Cs+ remain very fast. AIMD simulations of MgCl₂(aq) and MgSO₄(aq) show no exchanges around Mg²⁺. However, notice the contrasting behaviour of the chloride and sulphate ions on the solvent exchange dynamics around Ca^{2+} . In both solutions the counterion form SSHIPs with Ca^{2+} , but in $CaCl_2(aq)$ the MRT is 10.6 ps (faster water dynamics) compared with 61.2 ps (slower water dynamics) in pure liquid water; in $CaSO_4(aq)$ there were no accounted water exchange events around the FSS of the calcium ions.

A further comment concerns the effect of the level of theory and system size on the solvent exchange dynamics around Na $^+$, K $^+$, Ca $^{2+}$ and H $_2$ O (**Table 4**). We conducted calculations using different DFT methods, with and without the -D3 correction, and simulation boxes. While the absolute number of exchanges (N $_{\rm ex}$) varies significantly, the values of the mean residence time in the solvation shell of Na $^+$ ($^-$ 1 ps), K $^+$ ($^-$ 0.4 ps) and H $_2$ O ($^-$ 0.3 ps) is only marginally affected by the density functional and number of water molecules in the

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simulation box. For Ca2+, no exchanges were accounted with the PBE functional without the -D3 dispersion correction, and the value of MRT varies from 14 ps (revPBE-D3) to (71.7 ± 16.5) ps (PBE-D3) (Table 4). Recent estimates using different interatomic potential models and techniques to characterise the dynamics of water produced give MRT values lying in the range of 60-80 ps, in good agreement with the estimate obtained with the PBE-D3 functional.²² Using the same functional (PBE-D3), the MRT value for Ca²⁺ reduces to 13.1 ps from 61.2 ps on increasing the system size, which reveals that for Ca²⁺ the MRT can be sensitive to the system size.

Caging effect in the hydration shell of ions

Towards resolving the origin of the different local ion-water interaction, the velocity-autocorrelation functions (VACFs) of the metal ions (M) were computed.65 VACF is defined as

$$VACF_{M}(t) = \frac{1}{N_{O}N_{M}} \sum_{j=1}^{N_{O}} \sum_{i=1}^{N_{M}} v_{i}(t_{j}) \cdot v_{i}(t_{j} + t)$$
 (2)

where v_i is the velocity vector of atom i, N_O and N_M are the number of time origins spaced by t and number of metal ions respectively. Fig. 3 displays the VACF profiles obtained from AIMD (PBE-D3) simulations of the hydrated cation (single Mn+, no counter-ions) and aqueous electrolyte solutions (1.9 mol.kg⁻¹). Comparison of the VACF profiles obtained from the full set of NaCl, KCl and CaSO₄ solutions are reported in Fig. S4 of ESI.

Table 4. Number of accounted water exchange events $(N_{ex}^{H_2O})$ in the first coordination shell of Na⁺, K⁺, Ca²⁺, and H₂O with a duration of more than t* = 0.5 ps. The average coordination number (CN) of first shell is given by the integration of the metal-oxygen radial distribution function up to the first minimum. The values of $N_{ex}^{H_2 O}$ are normalized to the number of metal ions. Mean residence times (MRT) are in ps.

Na+ 6	3 PBE-			· · ex	MRT
		D3 5.1	50	233.0	1.1
	revPBI	E-D3 5.6	50	277.0	1.0
1	24 PBE-	D3 5.6	20	86.0	1.3
K+ 6	3 PBE-	D3 6.2	50	750.0	0.4
	revPBI	E-D3 6.1	50	665.0	0.5
Ca ²⁺ 6	3 PBI	E 6.0	50	0.0	_
	PBE-	D3 6.0	50	4.4 ± 1.0^{a}	71.7 ± 16.5 a)
	BLY	P 6.3	50	7.0	45.0
	BLYP-	D3 6.8	50	$19.6 \pm 1.0^{a)}$	15.6 ± 2.1 a)
	revP	BE 6.6	50	15.0	22.0
	revPBI	E-D3 6.1	50	22.0	13.9
Ca ²⁺ 12	24 PBE-	D3 6.5	50	26.3 ± 4.3 a)	12.7 ± 2.1 a)
H ₂ O 6	4 PBE-	D3 3.9	50	536.0 ± 19.3 b)	0.4 ± 0.0 b)
	revPBI	E-D3 4.1	50	938.7 ± 12.1 b)	0.2 ± 0.0 b)
	BLYP-	D3 4.5	50	822.4 ± 20.6 b)	0.3 ± 0.0 b)
H ₂ O 12	24 PBE-	D3 4.2	50	592.7 ± 42.3 b)	0.4 ±0.0 b)

 $^{^{\}text{a)}}$ The standard deviations of N_{ex} around Ca^{2+} have been computed from the variation of overlapping block averages of trajectories lasting 50 ps. b) The standard deviations of Nex around H2O have been computed from the averages of the number of exchanges around each water molecule in the simulation box.

MD studies of amorphous systems have used VACF analysis to probe the strength of interaction of network formers (e.g.

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Si, Al, P) and network modifiers (e.g. Ca, Na), with other surrounding "cage".66-68 In particular, the ode una help this to negative values in VACF profiles results from the so-called cage effect for the tagged particle; it takes some time for the particle to escape from the cage formed by its surrounding neighbors.⁶⁹ The VACF is very sensitive to the local ion hydration environment and changes in the profiles of this time correlation function, such as oscillatory behaviour and position of the first minimum (t_0), can be used as descriptors of the strength of ion interaction with the surrounding water molecules and of the rigidity of the interatomic vibration.⁷⁰ Rigid intra-atomic vibrations lead to a fast-oscillatory trend for the VACFs of Mg²⁺, Li⁺ and to a lesser extent Ca²⁺ (Fig. 3). Conversely, Na+, K+ and Cs+ ions show a much broader first minimum, which is not followed by a marked oscillatory behaviour as their interaction with the surrounding cage is weaker. The VACFs of Na⁺ ($t_0 = 0.1$ ps), K⁺ ($t_0 = 0.16$ ps), and Cs⁺ $(t_0 = 0.5 \text{ ps})$ in particular, display a smooth decay (Fig. 3A), which suggests a weak caging effect by the solvent; this explains the facile kinetics of dehydration of these cations. The lithium ion shows a much sharper first minimum ($t_0 = 0.04 \text{ ps}$) followed by a marked oscillatory behavior because of the more rigid interatomic vibration between the smaller Li⁺ with the surrounding hydration cage. The comparison of the Ca2+ and Mg²⁺ VACFs (**Fig. 3B**) shows a very different oscillatory behaviour ($t_0 = 0.01$ ps for Mg²⁺ and $t_0 = 0.06$ ps for Ca²⁺), which is indicative of the strengthening of ion-interaction with the surrounding water molecules and increased rigidity of the interatomic vibrations ongoing from Ca2+ to Mg2+. The

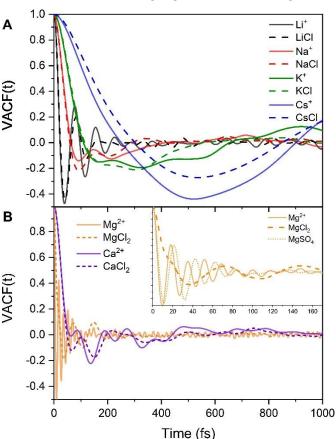


Figure 3. Velocity autocorrelation functions (VACF) of the metal ions obtained from AIMD (PBE-D3) simulations of the hydrated cation (single Mn+, no counterions) and of aqueous electrolyte solutions (1.9 mol.kg-1). (A) Solutions containing Li+, Na+, K+, Cs+. (B) Solutions containing Mg²⁺ and Ca²⁺.

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presence of chlorine counterions has no significant effect on the VACF profiles of the alkali metal ions and of the calcium ion. On the other hand, the inset of Fig. 3B shows a smoother decay of the VACF of Mg²⁺ in MgCl₂(aq), which is a clear indication of a looser inter-hexahedral vibration. Therefore, chlorine ions located in the second coordination shell of Mg²⁺ weaken the Mg(H₂O)₆²⁺ water 'cage' of the ion. This dynamic flexibility further highlights the effect of counterions on the dynamics of water molecules around Mg2+. A final comment concerns the sensitivity of the VACF on the density functional method used in the AIMD simulations. The VACF profiles of hydrated Ca2+ obtained from AIMD simulations using PBE, revPBE, and BLYP, with and without D3 correction, shows some differences (Fig. S5 in ESI). The PBE method generates the more marked oscillatory behaviour, to which corresponds a more rigid Ca-water coordination environment compared to the other DFT methods, explaining the absence of water exchange events around Ca2+ during the AIMD (PBE) simulation (Table 3).

Hydrogen bond statistics

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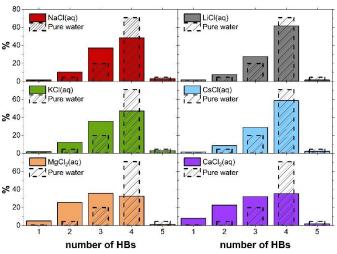


Figure 4. Percentage of water molecules that engage in n hydrogen bonds (HBs) in pure water and electrolyte solutions. Values obtained from AIMD (PBE-D3) trajectories.

Distribution of hydrogen bonds in bulk solution. The AIMD trajectories of pure liquid water and the aqueous electrolyte solutions were scanned to determine the effect of ions on the hydrogen bond (HB) structure, in terms of the percentage of molecules having n water—water HBs and the average number of HBs $(n_{\rm HB})$ per molecule (**Table S6** of ESI). We adopted the following geometrical criteria to determine the existence of a hydrogen bond between two water molecules:71,72 (i) the oxygen-oxygen distance is less than 3.5 Å; (ii) the intermolecular hydrogen-oxygen distance is less than 2.45 Å; (iii) the oxygen-oxygen-hydrogen angle is less than 30 degrees. The average number of HBs in bulk water computed from AIMD (PBE-D3) simulations is 3.75, which is in the range of experimental values (3.5-3.9) obtained using several techniques.⁷³ In comparison, the revPBE-D3 and BLYP-D3 functionals give $n_{\rm HB}$ values of 3.40 and 3.61, respectively. The average number of HBs reported in Table S6 decreases with the increase of salt concentration, a result in agreement with

previous MD studies of aqueous alkali valide salt solutions. Property of the solutions. Property of the solutions containing the doubly charged Mg^{2+} and Ca^{2+} ions, the average number of HBs is three, which is significantly lower than the 1.90 mol.kg⁻¹ solutions of LiCl (3.53), NaCl (3.40), KCl (3.37) and CsCl (3.50). Fig. 4 compares the percentage of water molecules that engage in n HBs for the solutions considered in this study. In pure liquid water, molecules are mostly engaged in four (70%) and three (20%) HBs, with a similar distribution being observed in LiCl(aq), NaCl(aq), KCl(aq) and CsCl(aq). On the other hand, in MgCl₂(aq) and CaCl₂(aq), the distribution of HBs is significantly shifted to lower n, with an almost equal proportion of molecules forming two, three and four HBs.

Hydrogen bond statistics in ionic solvation shell. Guàrdia et al. revealed significant modifications in the hydrogen bonding architecture for the water the first ionic solvation, which do not persist beyond the second shells.74 To investigate the origin of the ion-specific effects to the hydrogen bond structure, we have computed the distribution of HBs for the water molecules that, at each step of the AIMD simulation, are coordinated to the metal ion. Table 5 compares the percentage of molecules having n water-water HBs (f_n) and the average number of HBs (n_{HB}) in the first solvation shell of the monovalent (Li+, K+, Na+, and Cs+) and the divalent (Mg2+ and Ca²⁺) cations, together with the HB statistics for pure water and the water molecules coordinated to Cl-, with the latter obtained from AIMD of hydrated Cl-. For the hydrated metal ions (single M, no counterions), water molecules coordinated Mg²⁺ ($n_{\rm HB}$ = 2.20) and Ca²⁺ (2.30) display very large deviations from the HB distribution of the bulk (3.75). In comparison, the values of n_{HB} in the FSS of the more labile K⁺ and Cs⁺ are 3.06 and 3.24, respectively. The water molecules coordinated to Cl⁻ (3.61) display only minor deviations from H₂O. Fig. 5(A) highlights that the average numbers of HBs in the FSS of metal ions consistently reduces in chloride-bearing solutions compared to hydrated metal ions (single M, no counterions). The interaction between ions of opposite charge generates important differences in the distribution of HBs in the FSS. Speciation analysis of the aqueous electrolyte solutions has shown that metal ions are present as free ions only in very dilute electrolyte solutions.⁷⁵ Otherwise, ions tend to form contact and solvent shared pairs with the anions present in solution (Table 1). In electrolyte solutions, we can, therefore, consider that water molecules exist in different hydration states, depending on their relative position (coordination) to the cations and anions in solution. Classical MD simulations of MgCl₂ solutions with concentrations ranging from 0.3 mol kg⁻¹ (4 MgCl₂ units in 717 water molecules) to 2.8 mol.kg⁻¹ (33 MgCl₂ units in 633 water molecules) were also conducted to determine the fraction of water molecules that are in the bulk (or free water), coordinated to one single ion, or coordinated to both Mg2+ and Cl-. Fig. 5B shows that about 20% of water molecules are coordinated to both Mg²⁺ and Cl⁻, making cooperative effects important for most concentrations.

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Table 5. The distribution of the number of hydrogen-bonds (HBs) per water molecule in pure water and in the first solvation shell (FSS) of Li⁺, Na⁺, K⁺, Cs⁺, Mg²⁺ and Ca²⁺ obtained from AIMD (PBE-D3) simulations of hydrated ions and electrolyte solution. The values are percentages of H₂O with the given number of HBs. Concentration (b) in

	Number of HBs (%) in the FSS					
f_0	f_1	f ₂	f ₃	f_4	f 5	average
0.3	7. 3	31.1	60.9	0.5	0.0	2.54
0.3	11.1	26.8	61.0	0.8	0.0	2.51
0.2	4.7	26.1	66.6	2.4	0.0	2.66
0.6	1.7	26.8	62.5	2.4	0.0	2.58
0.7	5.9	25.9	65.8	1.6	0.0	2.62
1.3	14.1	35.0	47.3	2.2	0.0	2.35
0.2	1.5	13.5	61.3	23.4	0.1	3.06
0.5	4.0	27.3	57.6	10.5	0.0	2.74
0.5	4.2	26.0	60.7	8.5	0.1	2.73
2.0	11.9	33.3	47.0	5.5	0.2	2.43
0.0	0.6	10.4	53.3	35.0	0.6	3.24
0.9	4.2	20.5	52.6	21.7	0.2	2.91
0.5	12.2	53.7	33.6	0.0	0.5	2.20
0.3	14.3	69.4	15.9	0.0	0.0	2.01
6.0	32.4	41.1	20.6	0.0	0.0	1.76
0.2	8.4	52.2	39.1	0.0	0.0	2.30
1.4	26.7	51.2	20.2	0.1	0.0	1.91
3.8	25.0	41.3	29.9	0.0	0.0	1.96
0.0	0.3	4.6	19.8	70.7	4.6	3.75
0.0	0.5	5.5	25.4	65.8	2.8	3.61
֡֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜֜	0.3 0.3 0.2 0.6 0.7 1.3 0.2 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5	fo f1 0.3 7.3 0.3 11.1 0.2 4.7 0.6 1.7 0.7 5.9 1.3 14.1 0.5 4.0 0.5 4.2 2.0 11.9 0.0 0.6 0.9 4.2 0.3 14.3 6.0 32.4 0.2 8.4 1.4 26.7 3.8 25.0 0.0 0.3	fo f1 f2 0.3 7.3 31.1 0.3 11.1 26.8 0.2 4.7 26.1 0.6 1.7 26.8 0.7 5.9 25.9 1.3 14.1 35.0 0.5 4.0 27.3 0.5 4.2 26.0 2.0 11.9 33.3 0.0 0.6 10.4 0.9 4.2 20.5 0.5 12.2 53.7 0.3 14.3 69.4 6.0 32.4 41.1 0.2 8.4 52.2 1.4 26.7 51.2 3.8 25.0 41.3 0.0 0.3 4.6	f_0 f_1 f_2 f_3 0.3 7.3 31.1 60.9 0.3 11.1 26.8 61.0 0.2 4.7 26.1 66.6 0.6 1.7 26.8 62.5 0.7 5.9 25.9 65.8 1.3 14.1 35.0 47.3 0.2 1.5 13.5 61.3 0.5 4.0 27.3 57.6 0.5 4.2 26.0 60.7 2.0 11.9 33.3 47.0 0.0 0.6 10.4 53.3 0.9 4.2 20.5 52.6 0.5 12.2 53.7 33.6 0.3 14.3 69.4 15.9 0.0 32.4 41.1 20.6 0.2 8.4 52.2 39.1 1.4 26.7 51.2 20.2 3.8 25.0 41.3 29.9 0.0	f_0 f_1 f_2 f_3 f_4 0.3 7.3 31.1 60.9 0.5 0.3 11.1 26.8 61.0 0.8 0.2 4.7 26.1 66.6 2.4 0.6 1.7 26.8 62.5 2.4 0.7 5.9 25.9 65.8 1.6 1.3 14.1 35.0 47.3 2.2 0.2 1.5 13.5 61.3 23.4 0.5 4.0 27.3 57.6 10.5 0.5 4.2 26.0 60.7 8.5 0.0 11.9 33.3 47.0 5.5 0.0 0.6 10.4 53.3 35.0 0.9 4.2 20.5 52.6 21.7 0.5 12.2 53.7 33.6 0.0 0.3 14.3 69.4 15.9 0.0 0.0 32.4 41.1 20.6 0.0	f_0 f_1 f_2 f_3 f_4 f_5 0.3 7.3 31.1 60.9 0.5 0.0 0.3 11.1 26.8 61.0 0.8 0.0 0.2 4.7 26.1 66.6 2.4 0.0 0.6 1.7 26.8 62.5 2.4 0.0 0.7 5.9 25.9 65.8 1.6 0.0 0.2 1.5 13.5 61.3 23.4 0.1 0.5 4.0 27.3 57.6 10.5 0.0 0.5 4.2 26.0 60.7 8.5 0.1 2.0 11.9 33.3 47.0 5.5 0.2 0.0 0.6 10.4 53.3 35.0 0.6 0.9 4.2 20.5 52.6 21.7 0.2 0.5 12.2 53.7 33.6 0.0 0.5 0.0 32.4 41.1 20.6 0.0

Despite the influence to the HB network should be mainly ascribed to the specific Mg²⁺-water interaction, this confirms that the HB structure of the first hydration shell is governed by the subtle interplay between the level of hydration of the ions, and the interactions between ions, especially those of opposite charge. In more general terms, the cooperative effect of cations and anions in solution reflects a balance between solute-solvent and solute-solute interactions since relatively minor changes to the chemical characteristics result in dramatic changes to the HB distribution.

Water reorientation dynamics

The rotational dynamics of the water dipole, $\vec{\mu}$, was computed from the 1st order Legendre polynomial time correlation:⁷⁶

$$P_1(t) = \frac{\langle \vec{\mu}(0) \cdot \vec{\mu}(t) \rangle}{\vec{\mu}(0)^2} \tag{3}$$

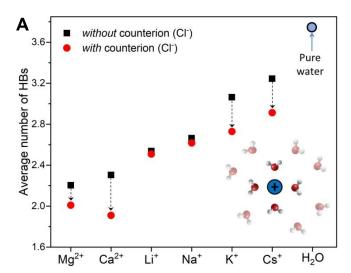
where $\vec{\mu}(0)$ and $\vec{\mu}(t)$ are the unit vectors defining the orientation of $\vec{\mu}$ at times 0 and t, respectively. The averages were computed using several time origins and overlapping intervals, each of length t = 16 ps. (Figs. S6–S9 in ESI).

Fig. 6 reports the concentration-dependent time correlation $P_1(t)$ profiles of the water molecules, obtained from the AIMD (PBE-D3) simulations of the hydrated cation (single M, no counterions) and of the aqueous electrolyte solutions. The $P_1(t)$ function starts at 1 and then decays asymptotically to zero because of the random and isotropic

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reorientation of the water molecules in solution. The early stages of fast loss of correlation are caused by his rational motion, whereas the long term decay is due to re-orientational motion and can be fit by a bi-exponential model, $\exp(-t/\tau_1) + b$ $\exp(-t/\tau_2)$. The overall time associated with this process, τ_{reor} , is given by the sum of the fitting parameters τ_1 and τ_2 . The value of τ_{reor} for pure liquid water (64 H₂O), computed at the PBE-D3 level of theory is 5.3 ps, is in good agreement with the experimental value of 4.8 ps.77

In electrolyte solutions, the presence of solvated ions accelerates the reorientation dynamics of water dipoles (faster $P_1(t)$ decay) compared to pure liquid water. However, the $P_1(t)$ profiles of NaCl(aq) and KCl(aq) do not consistently decrease with the concentration as they are also influenced by the type of ion pairs present in solution. For NaCl(ag), the fastest reorientation dynamics (fastest $P_1(t)$ decay) is observed for the 0.9 mol.kg⁻¹ solution, which only contains free Na⁺ and Cl⁻ ions. On the other hand, the 1.9 and 3.9 mol.kg-1 NaCl solutions, which contains contact and solvent-shared pairs, have a water reorientation behaviour similar to that of hydrated Na⁺ (Fig. 6). The correlation function $P_1(t)$ is, therefore, sensitive to solution speciation and can provide insights into the combined effect of cations and anions on the water reorientation dynamics. Chowdhuri and Chandra showed that in the vicinity of ions the diffusion coefficients are reduced and the orientational relaxation times are increased



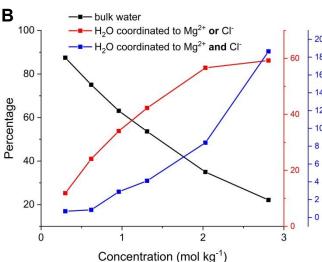


Figure 5. (A) Average number of hydrogen bonds (HBs) in the first hydration shell of metal ions obtained from AIMD (PBE-D3) simulations of hydrated metal ions (single M. no counterions) and of aqueous electrolyte solutions (1.9 mol.kg-1). (B) Distribution of the number of molecules among the water subpopulations, obtained from classical MD

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compared with pure water.⁷⁸ We have computed the reorientation dynamics of the water molecules in the first and second coordination shells of the metal ions, and of the water molecules in the bulk (beyond second coordination shells of all ions in solution). Fig. 7 compares the correlation profiles obtained from the analysis of the AIMD trajectories of the hydrated metal ions K⁺ and Mg²⁺ and of 1.9 mol.kg⁻¹ aqueous KCl(aq) and MgCl₂(aq). For hydrated K⁺, the reorientation dynamics of water dipoles in the hydration shells is similar to bulk behaviour. The reorientation dynamics of water dipoles is faster in chloride-containing solutions compared to that observed for the hydrated single metal ion: the $P_1(t)$ profiles of the water molecules in the first shell of K+ in KCl(aq) are drastically accelerated compared to the behaviour around hydrated K⁺ (single ion, no counterions). Conversely, the decay of the $P_1(t)$ profiles of the water molecules directly coordinated to Mg²⁺ are drastically slower than in the bulk, but the effect of solution composition (presence of other Mg²⁺ and Cl- in solution) is not significant. This can be explained by the much stronger ion-water interaction of Mg²⁺ compared to K⁺. The correlation function $P_1(t)$ is, therefore, sensitive to solution speciation and provide insights into the combined effect of cations and anions on the water reorientation dynamics.

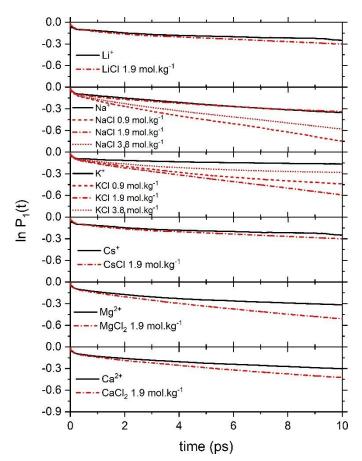


Figure 6. Profiles of the correlation function $P_1(t)$ for the water molecules obtained from AIMD (PBE-D3) simulations of hydrated metal ions (single M, no counterions) and of aqueous electrolyte solutions at different concentrations.

Conclusions

We have conducted DFT based molecular dynamics simulations of the alkali metal ions Li⁺, Na⁺, K⁺, Cs⁺ and of the alkaline earth metal ions Mg²⁺ and Ca²⁺, in pure water and electrolyte solutions containing Cl⁻ and SO_4^{2-} , with salt concentrations ranging from 0.9 to 3.8 mol.kg⁻¹, to determine the effect of solution composition on the structural and dynamic properties of the first solvation shell of metal ions. Based on the results we conclude the following:

In aqueous electrolyte solutions, the presence of the Cl⁻ and SO_4^{2-} counterions does not have a significant effect on the ion-water radial distribution functions of profiles of Li⁺, Na⁺, Mg^{2+} and Ca^{2+} .

The presence of counterions in solution influences the size of the hydration shell not only because of the formation of contact ion pairs, which reduces the coordination sites available around the metal ion, but also when cations and anions form solvent-shared ion pairs. We found, for example, that in $CaCl_2(aq)$ and $CaSO_4(aq)$, the chlorine and sulphate ions in the second coordination shell of Ca^{2+} stabilize the seven-coordinated $Ca(H_2O)_7^{2+}$ complex over the six-coordinated $Ca(H_2O)_6^{2+}$ species.

Calculation of the velocity autocorrelation function (VACF) of the metal ions has been used to probe the strength of the local ion-water interaction and the rigidity of the interatomic vibration. We found a smoother decay of the VACF of Mg^{2+} in $MgCl_2(aq)$, which is a clear indication of a looser interhexahedral vibration. Chlorine ions, located in the second coordination shell of Mg^{2+} weaken, the $Mg(H_2O)_6^{2+}$ water 'cage' of the ion. This dynamic flexibility further highlights the effect of counterions on the dynamics of water molecules around ions.

Analysis of the hydrogen bond statistics in ionic solvation shell shows how the influence to the water-water network cannot only be ascribed to the specific Mg²⁺—water interaction, but also to the subtle interplay between the level of hydration of the ions, and the interactions between ions, especially those of opposite charge.

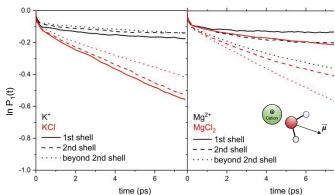


Figure 7. Profiles of the correlation function $P_1(t)$ for the water molecules in the first and second shell of K^+ and Mg^{2+} obtained from AIMD (PBE-D3) simulations of hydrated metal ions (single M, no counterions) and of aqueous electrolyte solutions (1.9 mol.kg⁻¹)

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Conflicts of interest

There are no conflicts to declare.

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