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Heterogeneous Microscopic Dynamics of Intruded Water in a Superhydrophobic Nanoconfinement: Neutron Scattering and Molecular Modeling

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1 Abstract

With their strong confining porosity and versatile surface chemistry, Zeolitic Imidazolate Frameworks - including the prototypical ZIF-8 - display exceptional properties for various applications. In particular, the forced intrusion of water at high pressure (~ 25 MPa) into ZIF-8 nanopores is of interest for energy storage. Such a system reveals also ideal to study experimentally water dynamics and thermodynamics in ultra-hydrophobic confinement. Here, we report on neutron scattering experiments to probe the molecular dynam-10 ics of water within ZIF-8 nanopores under high pressure up 11 to 38 MPa. In addition to an overall confinement-induced 12 slowing down, we provide evidence for strong dynamical 13 heterogeneities with different underlying molecular dynam-14 ics. Using complementary molecular simulations, these het-15 erogeneities are found to correspond to different microscopic 16 mechanisms inherent to vicinal molecules located in strongly 17 adsorbing sites (ligands) and other molecules nanoconfined 18 in the cavity center. These findings unveil a complex micro-19 scopic dynamics, which results from the combination of sur-20 face residence times and exchanges between the cavity sur-21 face and center.

Introduction

Interfacial and confined water has become a scientific field in itself at the crossroad of physics, chemistry and biology 25 with significant implication for applied science such as in geophysics, mechanical/fluid engineering, etc.^{1,2} Such wa-27 ter, which displays marked differences with its bulk counterpart, is of importance for situations where physicochemical interactions with surfaces are at play: protein folding, biological nanochannels, nanofiltration/phase separation, catal-31 vsis, etc. ^{1,3,4} Despite the microscopic origin and nanomet-32 ric range of these physicochemical interactions, the impact 33 of interfaces on the water structure and dynamics - with a 34 pronounced dependence on confinement size, geometry and 35 chemistry – manifests itself at the macroscopic scale as illustrated by considering water flow properties. 5-8 Hydrophilic, i.e. wetting, surfaces are typically characterized by a no slip boundary condition while hydrophobic, i.e. non-wetting, 39 surfaces are known to exhibit slippage. 9,10 Such slip bound-40 ary conditions are broadly considered as a mean to overcome 41 the permeability-selectivity trade-off in membrane science 42 such as for medical applications, water desalination, blue energy harvesting ¹¹ (with the goal to compete with high performance biological selective nanochannels ³).

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Despite their significant impact at the macroscopic level, the molecular specificities of water near solid surface remain only partly elucidated. At the microscopic level, a large body of experimental and numerical studies is available on the dynamics of water confined within wetting, i.e. hydrophilic, nanoporous media. In this case, the strong molecular interactions with the hydrophilic surfaces combined with severe in-pore confinement cause a decrease in water translational diffusion coefficient - i.e. a slowing down of water dynamics which gets more pronounced with tighter confinement and/or stronger surface interactions. 12-16 In partially wetting materials, available data on moderately hydrophobic materials such as carbon nanotubes suggest a more complex scenario with strongly enhanced water transport as probed using permeability measurements. 17 While such faster transport can be rationalized by elucidating the very specific interactions between water and the host surface (weakened hydrogen-bond network, curvature-induced friction decrease, etc.), 18 it does not necessarily imply an increase in the diffusion coefficient and associated faster selfdynamics. As a matter of fact, even if the solid surface impact is less pronounced with hydrophobic surfaces, a decreased self-diffusion coefficient with respect to that of bulk water has been also reported in the literature by different groups. 13,19 Nevertheless, in all these studies dedicated to hydrophilic and so-called hydrophobic materials, the affinity of water for the solid matrices is always strong enough to promote a spontaneous water condensation within the pores at a bulk vapor pressure lower than the saturation pressure.

Beyond hydrophilic and moderately hydrophobic materials, the microscopic dynamics in superhydrophobic nanoporous solids – such as all-silica zeolites and some Metal Organic Frameworks for which a pressure much larger than the water saturation pressure is needed to force water into the porosity – remains to be elucidated. Despite a well-established non-wetting thermodynamic behavior, ²⁰ the molecular mechanisms ruling water transport in these materials involve complex phenomena which are often overlooked in such superhydrophobic environment (residence steps, in-pore relocation, etc.). In particular, the elemental dynamical mechanisms remain almost unexplored using microscopic experiments (e.g. neutron scattering, nuclear magnetic resonance). In this context, Zeolitic Imidazolate Frameworks (ZIF) are attractive candidates owing to their tunable

chemistry, variety of topology and nanoscale pore size. 21-24 ZIF-8 is made of tetrahedral zinc ions linked by imidazolate organic groups (see insert in Figure 1). These two different moieties form a cubic structure of spherical cages, 11.6 Å in diameter, interconnected by 3.4 Å narrow gates which fill with water at a pressure of the order of 25 MPa at room temperature. The ZIF-8 structure, which is superhydrophobic, presents local heterogeneities as regularly encountered in biological confinement. The large microporous volume of ZIF-8 ($\sim 0.4 \text{ cm}^3/\text{g}$)²⁵ is experimentally of interest to reach - after intrusion - a significant content of confined water with respect to bulk water. While water within ZIF-8 has been largely explored from a structural viewpoint, ^{22,26–29} its dynamics has received only very little attention. On the one hand, Han et al. 30 considered the impact of defects on the dynamics of gas water molecules in ZIF-8, on the other hand Fraux et al. 28 reported a ten-fold decrease in liquid water self diffusivity. Despite these pioneering works, we note that the details of the molecular mechanisms leading to such a dynamics slowing down remain to be identified.

Here, we report on a coupled experimental and molecular simulation investigation of the microscopic dynamics of intruded water in superhydrophobic confinement. To the best of our knowledge, we performed the first direct measurement of the diffusion coefficient of liquid water in such superhydrophobic environment by means of quasi-elastic neutron scattering (QENS). Our experiments are combined with a dual Grand Canonical Monte Carlo/molecular dynamics simulation strategy which sheds light on the heterogeneous dynamics of water confined in such hydrophobic environment. While we confirm the average slowing down of the dynamics for confined water, careful analysis of our experimental data in the light of molecular simulations unveils a complex microscopic dynamics related to two different dynamical mechanisms: reduced dynamics for water located close to strongly adsorbing sites (ligands) and bulk-like water in the cavity center.

Material and Methods

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The molecular simulation approach first involves Grand Canonical Monte Carlo simulations to determine the number of adsorbed water molecules as a function of pressure. Then starting from configurations obtained at a pressure where the ZIF-8 porosity is filled with water, we performed molecular dynamics simulations to compare bulk and confined water dynamics. A supercell consisting of a $2 \times 2 \times 2$ ZIF-8 unit cell (the cubic lattice parameter is approximately 1.7 nm) was built to conduct these calculations. Periodic boundary conditions were used along the 3 space directions to avoid finite size effects. Following previous works on water in ZIF, the TIP4P (Transferable Intermolecular Potential-4 Point) allatom model is used to describe the water molecule while both a nominal and rescaled force field proposed by Amrouche et al.31 is used to model the water/ZIF-8 molecular interactions. All details on the molecular simulation strategy can be found in the Supporting Information (S5).

QENS experiments were performed at Institut Laue-Langevin (ILL, Grenoble) on the time-of-flight spectrometer IN6-Sharp with an incident wavelength of 5.1 Å – leading to an elastic energy resolution of 70 μ eV (dynamics faster than \sim 10 ps) over the momentum transfer q-range of 0.4-2 Å⁻¹. Because of the large hydrogen incoherent cross section for neutrons, the signal arising from hydrogen motions of

water dominates the scattered intensity, therefore providing a measurement of the self-dynamics of water in the sample. Descriptions of the experimental environment together with calibration and adequate background measurements are given in the Supporting Information (S2). In brief, an optimized aluminium high pressure cell was designed 32 to prepare a sub-millimetric uniform and dense layer of ZIF-8 powder (BASF, Basolite[®] Z1200), with similar volume fractions in intra-grain pores and inter-grain bulk voids. Measurements were performed under hydrostatic pressures of 4, 20, 28 and 38 MPa which are below and above the water intrusion pressure of 25 MPa (see Figure 2a) at ambient temperature. 24 The two lower pressures correspond to a matrix empty of liquid whereas the two higher pressures correspond to a matrix filled with liquid of interest for the study of water dynamics in this superhydrophobic framework. The reproducibility of the measurements was assessed by performing several short measurements during successive intrusion cycle from low to high pressure.

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Results and Discussion

We first investigate the water dynamics, using molecular simulation, from the mean square displacements (MSD) for bulk and confined water with respect to time (Figure 1). In the long time limit t > 1 ns, the short-wave vector displacements (large distances) are dominated by translational motion where both the O and H atoms diffuse over the same distance on a given time scale. While the Fickian regime is reached on a short time scale for bulk water ($t \sim 100 \text{ ps}$), we found that much longer simulations are required to investigate this normal diffusion regime for confined water (typically, $t \sim 10$ ns). The diffusivity of water confined in ZIF-8 at 300 K is found to be smaller than its bulk counterpart; typically, $D_T \sim 3.4 \times 10^{-9} \text{ m}^2/\text{s}$ for bulk water while D_T $\sim 2.6 \times 10^{-10}$ m²/s for confined water. These two values obtained with the TIP4P water model being 30% larger than the ones obtained with the SPC/E model (extended simple point charge model)²⁸. However, importantly, the ten-fold decrease in the water self diffusivity between bulk and con-

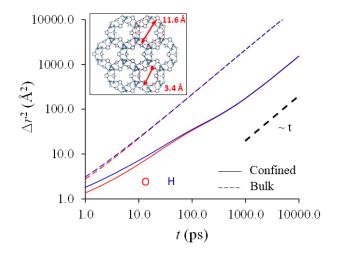


Figure 1: Mean square displacements Δr^2 as obtained using molecular dynamics for the O and H atoms of water in bulk water (dashed lines) and confined in ZIF-8 (solid lines). The black dashed line is a guide to the eye to indicate the Fickian regime where $\Delta r^2 \sim t$. The insert shows the ZIF-8 structure.

fined values is independent of the force field or its scaling (as detailed in the Supplementary Information).

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In the short time limit (t < 10-100 ps), for both bulk and confined water, the O and H mean square displacements depart from each other due to rotational effects. Interestingly, while the mean square displacements for bulk water is timescale independent in the sense that it exhibits a single regime once rotational effects are no longer noticeable, the data for confined water indicate a heterogeneous dynamics with different regimes. In more detail, between the time range dominated by rotation/translation coupling (t < 100 ps) and the normal Fickian regime ($t \sim 1$ -10 ns), a subdiffusive regime where $\Delta r^2 \sim t^\alpha$ with $\alpha < 1$ is observed.

At this stage it is useful to comfort these initial numerical results with experimental values of the diffusion coefficient D_T . The confined water dynamics is extracted from the dynamical structure factor $S(q,\omega)$. ³³ The experimental $S(\omega)$ spectra (summed over the whole q-range) at the different investigated hydrostatic pressures are shown in Figure 2b. The contribution from the ZIF-dry matrix, which is purely elastic, confirms the absence of QENS signal that could arise from imidazole ring rotation on the investigated time scale. Moreover, complementary measurements have been performed on the spectrometer IN13 (ILL), covering a shorter ($\sim 100 \text{ ps}$) timescale than IN6-SHARP. The results show that the rotational motion of the 2-methyl-imidazole rings is not influenced by the intrusion process (as it occurs on a timescale that is included in the elastic line of the present experiment). Details are presented in the Supporting Information (S1). We also observe that spectra below P_{int} =25 MPa (4 and 20 MPa) are very similar as well as those above P_{int} (28 and 38 MPa). Below 25 MPa, only inter-grain water is present outside the ZIF-8 grains as the nanopores are empty. Thus, the observed intensity increase between 20 and 28 MPa (Figure 2c) provides evidence for the presence of water within the nanopores above P_{int} . The obtained high-pressure sig-

nal is therefore a direct measurement of the water dynamics in such superhydrophobic confinement. The limited ZIF-8 compressibility, 34 which leads to minor deformations of the matrix on the considered pressure range - essentially related to a slight dilation of the order of 0.5% occurring within the narrow pressure window associated to the filling process 35 – is expected to not alter the liquid water dynamics within the pores after intrusion. The superimposition of signals measured at 28 MPa and 38 MPa confirms that once the matrix is filled, the pressure increase does not alter the water dynamics, neither directly through a potential change of water structure nor indirectly because of the flexibility of the matrix. The dynamical structure factor, $S(q, \omega)$, in the investigated energy range, can be approximated by the product of the Debye – Waller factor, accounting for the vibrational motions, with the decoupled diffusive motions, such as rotational, $S_R(q,\omega)$, and translational, $S_T(q,\omega)$, ones:³³

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$$S(q,\omega) = [S_T(q,\omega) + S_R(q,\omega)] \exp\left(-\frac{q^2\langle u^2\rangle}{3}\right)$$
 (1)

with $\langle u^2 \rangle$ the mean-square vibrational amplitude of the hydrogen atoms. In a first approach, both translational and rotational motions were described using a Lorentzian function of unconstrained half-width at half-maximum (hwhm), Γ . Γ_T and Γ_R refer to the translational and rotational motions, respectively. In the Fickian regime, the variation of Γ_T is linear with q^2 as $\Gamma_T = D_{T_{Fick}}q^2$ where $D_{T_{Fick}}$ is the translational self-diffusion coefficient. For the localized rotational motion, the width is q-independent: $\Gamma_R = 2D_R$ with D_R the rotational diffusion constant. Such a phenomenological model relying on two Lorentzian functions provides a reliable description of the water dynamics under different thermodynamical conditions. We emphasize that the appropriate background for each pressure condition is subtracted in order to isolate successively the inter-grain and intra-grain wa-

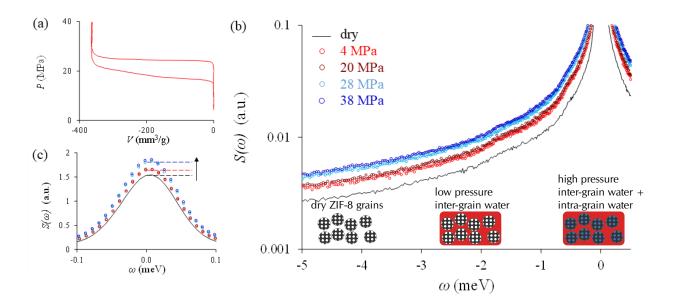


Figure 2: (a) Experimental water intrusion / extrusion cycle at room temperature in a ZIF-8 material. V(P) indicates the intruded and extruded volume V of water per g of ZIF material as a function of the exerted pressure P. (b) $S(\omega)$ summed over q [from 0.4 to 2 Å $^{-1}$] of the quasi-elastic spectra $S(q,\omega)$ obtained for dry ZIF-8 (black), low-pressure water in contact with ZIF-8 (red) and high-pressure water intruded in ZIF-8 (blue). Inset: schematics of these three situations. (c) Zoom on the elastic peak showing the overall increase of the intensity upon water addition.

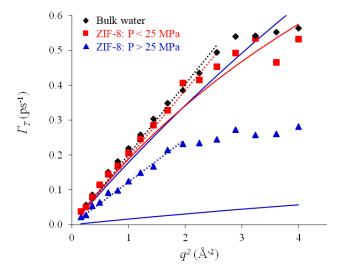


Figure 3: Half-width at half-maximum Γ_T of the translational contribution versus q^2 for bulk water (black), low-pressure water in contact with ZIF-8 (red) and high-pressure water intruded in ZIF-8 (blue). Error bars are smaller than the symbol size. The symbols refer to the results obtained with the analysis 1 (Eq. 1). The dashed segments indicate the Fickian regime where $\Gamma_T \sim q^2$. The solid lines are attributed to the results obtained with the analysis 2 (Eq. 2) using the jump diffusion model. We note the presence of two solid blue lines associated to the heterogeneous behavior of the intruded high-pressure water obtained with the analysis 2.

ter dynamics. Moreover, we assume that inter-grain water is identical at low and high pressures, as justified by the very low-pressure dependence of the bulk translational diffusion coefficient in the investigated pressure range (< 40 MPa). 36 More information on this analysis and curve fitting quality are given in the Supporting Information (S2). The obtained q^2 dependence of Γ_T is shown in Figure 3 (symbols) while the extracted parameters are given in Table 1.

The translational dynamics for bulk water and inter-grain water obey to Fick's law for $q^2 < 3 \text{ Å}^{-2}$. The corresponding self-diffusion coefficients, of $2.01 \times 10^{-9} \text{ m}^2/\text{s}$ and $1.92 \times 10^{-9} \text{ m}^2/\text{s}$ respectively, are very close to each other, and in very good agreement with values found in the literature for bulk water at 295 K³⁷ (incidentally we note that the MSD value measured with the numerical TIP4P water model is 70% larger than these experimental values). The water surrounding the grains is therefore only poorly influenced by the inter-grain boundaries on the time and space scales investigated. The deviation from Fick's law at large q is expected because of the simplified model – single Lorentzian – used for the rotational dynamics (instead of the full derivation taking into account higher Lorentzian orders and coupling between the two relaxation dynamics – see Eq. 3 and in S3 for more details).

In contrast to inter-grain water, the translational dynamics for intra-grain water follows Fick's law up to $q^2 \sim 2~\text{Å}^{-2}$ only. In the low q Fickian regime, the confinement effect leads to a decrease of both translational and rotational diffusion coefficients with respect to their bulk counterpart. The slowing down measured experimentally is only of the order of 30% with respect to the bulk values. Such a limited departure from the bulk dynamics should lead, for a single water pop-

ulation, to a linear Fickian regime that extends much beyond $q^2 \sim 2~{\rm Å}^{-2}$. As detailed in the Supporting Information (S3), the deviation from Fick's law beyond this q value, marked by the onset of a plateau for Γ_T , provides a signature of the non homogeneous behavior of intra-grain water, due to the influence of the discrete atomic structure surrounding water (also highlighted in Figure 1 with the subdiffusive regime at short times). To further analyze this departure from an effective average behavior, a more rigorous model is needed to cover and accurately describe both the Fickian and non-Fickian dynamical regimes.

Table 1: Numerical values of the extracted fitting parameters for the analysis 1. $D_{T_{Fick}}$ was calculated with the Fick's law on the appropriate q-range.

Analysis 1: eq. 1	$D_{T_{Fick}} \times 10^{-5} [\text{cm}^2/\text{s}]$	$D_R [\mathrm{ps}^{-1}]$
Bulk water	2.01 ± 0.03	2.06 ± 0.01
ZIF-8: P<25 MPa	1.92 ± 0.04	2.09 ± 0.02
ZIF-8: P>25 MPa	1.22 ± 0.02	1.75 ± 0.02

We performed therefore a deeper analysis based on a formal description of water translation through the use of the jump diffusion model ³³ coupled to isotropic rotational diffusion in a sphere of radius R (equal to the OH distance in a water molecule ~ 1 Å). Each population will then be described by the new fitting model:

$$S(q,\omega) = \exp\left(-\frac{q^2\langle u^2\rangle}{3}\right)S(q,\omega)_{T\otimes R}$$
 (2)

with 307

$$S(q,\omega)_{T\otimes R} = \left(\frac{1}{\pi} \frac{\Gamma_T}{\omega^2 + \Gamma_T^2}\right) \otimes \left(A_0(q).\delta(\omega) + \sum_{i=1}^{\infty} A_i(q).\left(\frac{1}{\pi} \frac{\Gamma_i}{\omega^2 + \Gamma_i^2}\right)\right)$$
(3)

where Γ_T is defined within the jump-diffusion model:

$$\Gamma_T = \left(\frac{D_{T_{jump}}q^2}{1 + \tau D_{T_{jump}}q^2}\right) \tag{4}$$

with τ the residence time, $A_i(q) = (2i+1)j_i^2(qR)$ where j_i is the spherical Bessel function of order i and $\Gamma_i = i(i+1)D_R$.

As expected, a single water population only is not sufficient to accurately fit our experimental data on the entire (q, w) range. On the other hand, a good fit is obtained using two water populations (ratio of 70/30 for fast and slow population, respectively). Each of these two populations are described with the model given in Eqs. 2 and 3 (fitting curves are presented in the Supporting Information (S2)). The resulting fitting parameters are shown in Table 2. The fast population is similar to bulk water while the slow population has a translational diffusion coefficient one order of magnitude lower as inferred from the MSD obtained using molecular simulation (Figure 1). As detailed in the Supporting Information (S3), at low q, the average dynamics of the two populations is similar to the one presented in Analysis 1 with an effective translational diffusion coefficient $\sim 30\%$ than that of bulk water. This moderate deviation at low *q* is related to the small proportion of the slow water population with respect to the fast one. At high q, however, the slow water popula-

Table 2: Numerical values of the extracted fitting parameters of the confined water in ZIF-8: P>25 MPa for the Analysis 2

Analysis 2: eq. 2	$D_{T_{Jump}} \times 10^{-5} [\text{cm}^2/\text{s}]$	τ [ps]	$D_R [{ m ps}^{-1}]$
ZIF-8: P<25 MPa	2.09 ± 0.05	0.55 ± 0.01	1.83 ± 0.01
ZIF-8: P>25 MPa: Population 1	1.87 ± 0.02	0.25 ± 0.05	2.18 ± 0.04
ZIF-8: P>25 MPa: Population 2	0.167 ± 0.007	2.55 ± 1.6	0.17 ± 0.01

tion, despite its limited proportion, becomes a leading term which contributes to the plateau observed for Γ_T in Figure 3 in the framework of analysis 1. Furthermore, moving now to the temporal domain using the inverse time Fourier transform of the formal model Eq. 2 (see Supporting Information S4) for the two populations, the translational term expresses as the sum of two time-decaying exponentials. At long time this translational term should be controlled by the slower exponential decay only associated to the slow population. This means that at long time a Fickian regime with a diffusion coefficient of the order of 1.7×10^{-10} m²/s is expected. This analysis of experimental data confirms, at long time, the tenfold decrease observed from numerical MSD and reveals, at short time, that the heterogeneous dynamics of confined water in ZIF-8 can be interpreted as the interplay between a slow and a main bulk-like water population.

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Attempting the same two-population decomposition using numerical data, a similar trend – even if to a less extent – is also observed when focusing on short time scales. As shown in Figure 4a, the self (incoherent) intermediate

scattering function, f(q, t) (inverse time Fourier transform of $S(q,\omega)$) extracted from molecular dynamics simulations for bulk water conforms to the classical behavior expected for Fickian diffusion. The superimposition of the curves plotted with respect to q^2t reveals that the rotational contribution is restricted to a very short time scale. This is further confirmed by the large value of $D_R \sim 1~\mathrm{ps^{-1}}$ obtained from the fit of f – inset of Figure 4a – using the inverse time Fourier transform of the formal model Eq. 2 (see Supporting Information S4). The dominant translational contribution of the form $\exp(-D_{T_{Jump}}q^2t)$ is characterized by a diffusion coefficient $D_{T_{Jump}}$ which agrees with the one inferred from MSD calculations. In contrast, as previously observed from our experimental data and molecular simulation MSD, f(q, t) for confined water display a non-Fickian behavior which can be approximated again with the contribution of a fast and a slow population (ratio 60/40) – each of them modeled with the inverse time Fourier transform of Eq. 2 (see inset of Figure 4a). Nevertheless, $D_{T_{lump}}$ for the fast population remains four times smaller than that for bulk water; this is a

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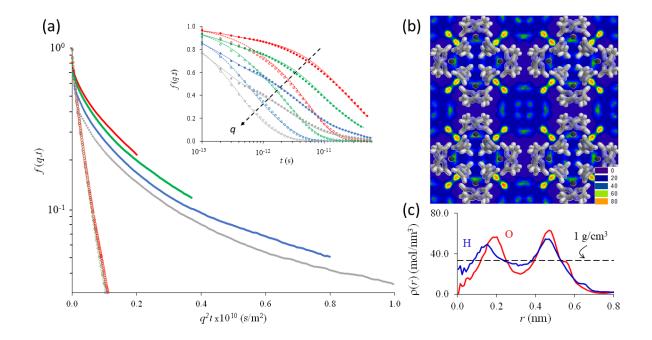


Figure 4: (a) Self (incoherent) intermediate scattering function f(q,t) as a function of q^2t for different scattering vector $\mathbf{q}=|\mathbf{q}|$ for bulk water (open circles) and for water confined in ZIF-8 (full dots). The color code is as follows: $q=0.7~\text{Å}^{-1}$ (red), $q=1.0~\text{Å}^{-1}$ (green), $q=1.5~\text{Å}^{-1}$ (blue), and $q=2.0~\text{Å}^{-1}$ (grey). Inset: f(q,t) according to t for bulk (open circle) and confined water (full dots). Dashed lines, bulk water: fit with a single water population using the inverse Fourier transform of Eq. 2 with $D_{T_{Jump}}=4.2\times10^{-9}~\text{m}^2/\text{s}$, $\tau=0.9~\text{ps}$ and $D_R=0.8~\text{ps}^{-1}$. Full lines, confined water: fit with a fast and a slow population (ratio 60/40) with $D_{T_{Jump}}=1.1\times10^{-9}/3.4\times10^{-10}~\text{m}^2/\text{s}$, $\tau=0.25/20~\text{ps}$ and $D_R=2/0.1~\text{ps}^{-1}$ (b) Contour plot showing the density of water in a ZIF-8 material as obtained using Grand Canonical Monte Carlo simulations. The density increases with color as follows: blue, green, yellow, orange and red. (c) Simulated density profiles $\rho(r)$ for the O and H atoms of water intruded in ZIF-8. r is the distance from the center of the ZIF cages shown in the contour plot in (b). The dashed line indicates the bulk density of water at room temperature.)

major difference with respect to experimental results. The rotational diffusion coefficient of the slower population being 20 times smaller than the fast population, in qualitative agreement with experimental results, the rotational contribution extends to much longer time than for bulk water and contributes to the separation of the curves observed in Figure 4a at large q^2t values. Interestingly, the functions f do not present a linear asymptotic trend at long q^2t values that would be expected in the presence of two distinct water populations only. This means that such a basic yet efficient decomposition to quantitatively analyze experimental and numerical data does not fully reflect the heterogeneity of water dynamics in such complex materials partly controlled by residence times at surfaces (see Bousige $et\ al.^{38}$ for a recent discussion on intermittent diffusion in complex nanoporous media).

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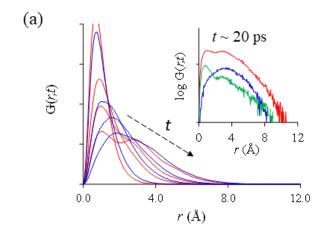
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At the microscopic level, the strong variations of water density as obtained from GCMC simulation (see Figure 4b and Figure 4c) reveal and confirm a complex heterogeneous water structure within the ZIF-8 porosity. These data, which are consistent with previous molecular simulation in the field, 22 show that water is adsorbed on specific sites, close to the ligands at a distance about 0.4 - 0.5 nm from the pore center [see marked density spots in Figure 4b], and in less adsorbing sites located closer to the pore center [see less marked density spots at a distance ~ 0.2 nm]. Although these adsorbing sites are hydrophobic enough to not trap water at ambient pressure, their relative affinity for the intruded water contribute to the marked slowing down its dynamics observed at long time.

For times much longer than the residence time, exchanges between the different adsorbing sites contribute to the complex heterogeneous dynamics and hence to the effective behavior accessible experimentally. Such a contribution from this molecular intermittence can be investigated by means of the diffusion propagator G(r,t) (inverse space and time Fourier transform of $S(q, \omega)$) which corresponds to the probability that a molecule gets located at a position r at a time t while being at r = 0 at a time t = 0. At short times (t < 10 ps), G(r,t) shown in Figure 5a for both O and H atoms of confined water is monomodal and reflects local displacements ($r \sim 0.1$ nm) which are similar to those found for bulk water on the same time scale (although of a slightly smaller amplitude in confinement). For larger times (t > 10 ps), the propagator becomes bimodal for O and H atoms, therefore indicating two characteristic translational modes coexisting on the same time scale. We show in the insert of Figure 5a the propagators for the different water populations confined in ZIF-8 (note the use of a log scale for improved visualization purpose). In more detail, we measured the propagator for molecules initially located at time t = 0 in the strongly adsorbing sites close to the ZIF-8 ligands (see density contour plot shown in Figure 4b) and that for molecules initially located in the cavity center not directly in contact with ZIF-8 ligand/surface. Such data confirm that the fast diffusion mode (i.e. subpeak in the propagator located at distance $r \sim 3.5 \text{ Å}$) corresponds to molecules located within the cavity center while the slow diffusion model corresponds to molecules initially located in the strongly adsorbing sites. Interestingly, for both contributions, some molecules undergo displacements corresponding to the other mode. This is due to the fact that, during the characteristic time t = 10 ps considered here, some molecules leave the cavity center to migrate towards the strongly adsorbing sites and reciprocally.

Additionally to the analysis of the translational motions,



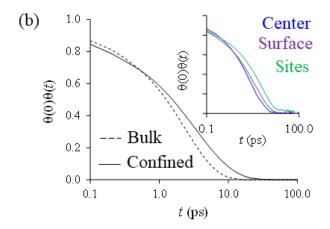


Figure 5: (a) Diffusion propagator G(r,t) as obtained using molecular dynamics at room temperature for the O (red lines) and H (blue lines) atoms of water confined in ZIF-8. The dashed line is a guide to the eye to indicate that the data are estimated at increasing times t = 1, 5, 10 and 20 ps. As discussed in the main text, the insert shows the different contributions to the G(r,t) data for the O atoms at t =20 ps: all molecules in the ZIF-8 cage (red), molecules in the cage center (blue) and molecules in strongly adsorbing sites (green). (b) Rotational autocorrelation time $\theta(t)\theta(0)$ showing the change in a water molecule time at a time t = 0 and a time t later. The dashed and solid lines are for bulk water and water confined at room temperature in ZIF-8. The insert shows the same data for confined water but for different subpopulations: molecules in the cage center (blue), molecules at the cage surface (purple), and molecules in strongly adsorbing sites (green).

we extracted the rotational relaxation time distribution for water in the different regions of the ZIF-8 cavities to clarify more specifically the impact of the surface heterogeneities³⁹ (see Figure 5b). The timescale of rotational motions enables indeed a more local analysis since molecules, over such a short timescale, do not significantly exchange between different sites. Owing to their intrinsically slow dynamics, the rotational motion of the molecules located in the strongly adsorbing sites is slower than for molecules in the pore center again with a typical rotational relaxation time a few times larger than for water in the pore center. Such a slow rotational motion only applies to molecules in the strong adsorbing sites. For molecules at the ZIF-8 cavity surface, away

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from adsorbing sites, we indeed confirm that the relaxation time distribution is close to the one found for molecules in the pore center.

450 Conclusion

By means of a coupled experimental and molecular simulation study, we provide microscopic insights into the dynamics of liquid water intruded at high pressure within the ZIF-8 453 superhydrophobic nanoporous material. For the first time, 454 the direct experimental measurement of water in superhy-455 drophobic cavities enables a comparison with a molecular 456 simulation investigation. Qualitatively, a slowing down of water is evidenced in both approaches. From an analysis of experimental data based on the contribution of two water populations, we show that at short distances, the main pop-460 ulation presents a bulk-like behavior while the slow popula-461 tion has a translational diffusion coefficient ten times smaller 462 than the bulk coefficient. This arbitrary subdivision into two 463 populations is sustained by the microscopic details obtained 464 from our molecular simulation approach. A fast population 465 is indeed identified in the cavity center while a slower population is found close to the different adsorbing sites at the cage surface (especially at the vicinity of imidazole ligands). 468 Even if water has a long residence time in these adsorbing 469 sites, exchanges with the pore center are also observed. Such 470 intermittence is found to contribute to the overall heteroge-471 neous water dynamics and could explain the discrepancy 472 between experimental and numerical values of the translational diffusion coefficient of the fast population. We believe that our findings represent an important step forward in the 475 understanding of complex dynamical behavior in superhy-476 drophobic systems which are relevant to many industrial ap-477 plications. 478

50 Supporting Information Available

The following files are available free of charge.

IN13 data; IN6 Experimental details and fitting curves; Complementary information on the analysis 1 and 2 in the experimental part; Computational details

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TOC Graphic

