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

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## 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 ( $\sim 25$  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 dynamics of water within ZIF-8 nanopores under high pressure up to 38 MPa. In addition to an overall confinement-induced slowing down, we provide evidence for strong dynamical heterogeneities with different underlying molecular dynamics. Using complementary molecular simulations, these heterogeneities are found to correspond to different microscopic mechanisms inherent to vicinal molecules located in strongly adsorbing sites (ligands) and other molecules nanoconfined in the cavity center. These findings unveil a complex microscopic dynamics, which results from the combination of surface residence times and exchanges between the cavity surface and center.

## Introduction

Interfacial and confined water has become a scientific field in itself at the crossroad of physics, chemistry and biology with significant implication for applied science such as in geophysics, mechanical/fluid engineering, etc.<sup>1,2</sup> Such water, 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, catalysis, etc.<sup>1,3,4</sup> Despite the microscopic origin and nanometric range of these physicochemical interactions, the impact of interfaces on the water structure and dynamics – with a pronounced dependence on confinement size, geometry and chemistry – manifests itself at the macroscopic scale as illustrated by considering water flow properties.<sup>5–8</sup> Hydrophilic, i.e. wetting, surfaces are typically characterized by a no slip boundary condition while hydrophobic, i.e. non-wetting, surfaces are known to exhibit slippage.<sup>9,10</sup> Such slip boundary conditions are broadly considered as a mean to overcome the permeability-selectivity trade-off in membrane science such as for medical applications, water desalination, blue en-

ergy harvesting<sup>11</sup> (with the goal to compete with high performance biological selective nanochannels<sup>3</sup>).

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.<sup>12–16</sup> 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.<sup>17</sup> 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.),<sup>18</sup> it does not necessarily imply an increase in the diffusion coefficient and associated faster self-dynamics. 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.<sup>13,19</sup> 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,<sup>20</sup> 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.<sup>21–24</sup> 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 ( $\sim 0.4 \text{ cm}^3/\text{g}$ )<sup>25</sup> 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,<sup>22,26–29</sup> its dynamics has received only very little attention. On the one hand, Han *et al.*<sup>30</sup> considered the impact of defects on the dynamics of gas water molecules in ZIF-8, on the other hand Fraux *et al.*<sup>28</sup> 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

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) all-atom model is used to describe the water molecule while both a nominal and rescaled force field proposed by Amrouche *et al.*<sup>31</sup> 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  $\mu\text{eV}$  (dynamics faster than  $\sim 10$  ps) over the momentum transfer  $q$ -range of 0.4–2  $\text{\AA}^{-1}$ . 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<sup>32</sup> to prepare a sub-millimetric uniform and dense layer of ZIF-8 powder (BASF, Basolite<sup>®</sup> 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.<sup>24</sup> 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.

## 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 ( $t \sim 100$  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  $D_T \sim 2.6 \times 10^{-10} \text{ m}^2/\text{s}$  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)<sup>28</sup>. However, importantly, the ten-fold decrease in the water self diffusivity between bulk and con-

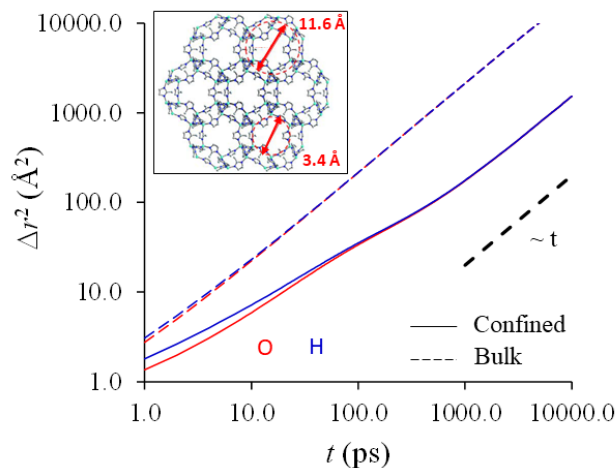


Figure 1: Mean square displacements  $\Delta r^2$  as obtained using molecular dynamics for the O and H atoms of water in bulk water (dashed lines) and confined in ZIF-8 (solid lines). The black dashed line is a guide to the eye to indicate the Fickian regime where  $\Delta r^2 \sim t$ . The insert shows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

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

192 In the short time limit ( $t < 10\text{-}100$  ps), for both bulk and  
 193 confined water, the O and H mean square displacements  
 194 depart from each other due to rotational effects. Interest-  
 195 ingly, while the mean square displacements for bulk water  
 196 is timescale independent in the sense that it exhibits a single  
 197 regime once rotational effects are no longer noticeable,  
 198 the data for confined water indicate a heterogeneous dynam-  
 199 ics with different regimes. In more detail, between the time  
 200 range dominated by rotation/translation coupling ( $t < 100$   
 201 ps) and the normal Fickian regime ( $t \sim 1\text{-}10$  ns), a subdiffu-  
 202 sive regime where  $\Delta r^2 \sim t^\alpha$  with  $\alpha < 1$  is observed.

203 At this stage it is useful to compare these initial numeri-  
 204 cal results with experimental values of the diffusion coeffi-  
 205 cient  $D_T$ . The confined water dynamics is extracted from  
 206 the dynamical structure factor  $S(q, \omega)$ .<sup>33</sup> The experimental  
 207  $S(\omega)$  spectra (summed over the whole  $q$ -range) at the dif-  
 208 ferent investigated hydrostatic pressures are shown in Fig-  
 209 ure 2b. The contribution from the ZIF-dry matrix, which  
 210 is purely elastic, confirms the absence of QENS signal that  
 211 could arise from imidazole ring rotation on the investigated  
 212 time scale. Moreover, complementary measurements have  
 213 been performed on the spectrometer IN13 (ILL), covering a  
 214 shorter ( $\sim 100$  ps) timescale than IN6-SHARP. The results  
 215 show that the rotational motion of the 2-methyl-imidazole  
 216 rings is not influenced by the intrusion process (as it occurs  
 217 on a timescale that is included in the elastic line of the present  
 218 experiment). Details are presented in the Supporting Infor-  
 219 mation (S1). We also observe that spectra below  $P_{int}=25$  MPa  
 220 (4 and 20 MPa) are very similar as well as those above  $P_{int}$  (28  
 221 and 38 MPa). Below 25 MPa, only inter-grain water is present  
 222 outside the ZIF-8 grains as the nanopores are empty. Thus,  
 223 the observed intensity increase between 20 and 28 MPa (Fig-  
 224 ure 2c) provides evidence for the presence of water within  
 225 the nanopores above  $P_{int}$ . The obtained high-pressure sig-

226 nal is therefore a direct measurement of the water dynamics  
 227 in such superhydrophobic confinement. The limited ZIF-8  
 228 compressibility,<sup>34</sup> which leads to minor deformations of the  
 229 matrix on the considered pressure range – essentially related  
 230 to a slight dilation of the order of 0.5% occurring within the  
 231 narrow pressure window associated to the filling process<sup>35</sup> –  
 232 is expected to not alter the liquid water dynamics within the  
 233 pores after intrusion. The superimposition of signals mea-  
 234 sured at 28 MPa and 38 MPa confirms that once the matrix  
 235 is filled, the pressure increase does not alter the water dyn-  
 236 amics, neither directly through a potential change of water  
 237 structure nor indirectly because of the flexibility of the mat-  
 238 rix. The dynamical structure factor,  $S(q, \omega)$ , in the inves-  
 239 tigated energy range, can be approximated by the product  
 240 of the Debye – Waller factor, accounting for the vibrational  
 241 motions, with the decoupled diffusive motions, such as rota-  
 242 tional,  $S_R(q, \omega)$ , and translational,  $S_T(q, \omega)$ , ones:<sup>33</sup>

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

243 with  $\langle u^2 \rangle$  the mean-square vibrational amplitude of the hy-  
 244 drogen atoms. In a first approach, both translational and  
 245 rotational motions were described using a Lorentzian func-  
 246 tion of unconstrained half-width at half-maximum (hwhm),  
 247  $\Gamma$ .  $\Gamma_T$  and  $\Gamma_R$  refer to the translational and rotational mo-  
 248 tions, respectively. In the Fickian regime, the variation of  $\Gamma_T$   
 249 is linear with  $q^2$  as  $\Gamma_T = D_{T_{Fick}} q^2$  where  $D_{T_{Fick}}$  is the transla-  
 250 tional self-diffusion coefficient. For the localized rotational  
 251 motion, the width is  $q$ -independent:  $\Gamma_R = 2D_R$  with  $D_R$   
 252 the rotational diffusion constant. Such a phenomenological  
 253 model relying on two Lorentzian functions provides a reli-  
 254 able description of the water dynamics under different ther-  
 255 modynamical conditions. We emphasize that the appropriate  
 256 background for each pressure condition is subtracted in or-  
 257 der to isolate successively the inter-grain and intra-grain wa-

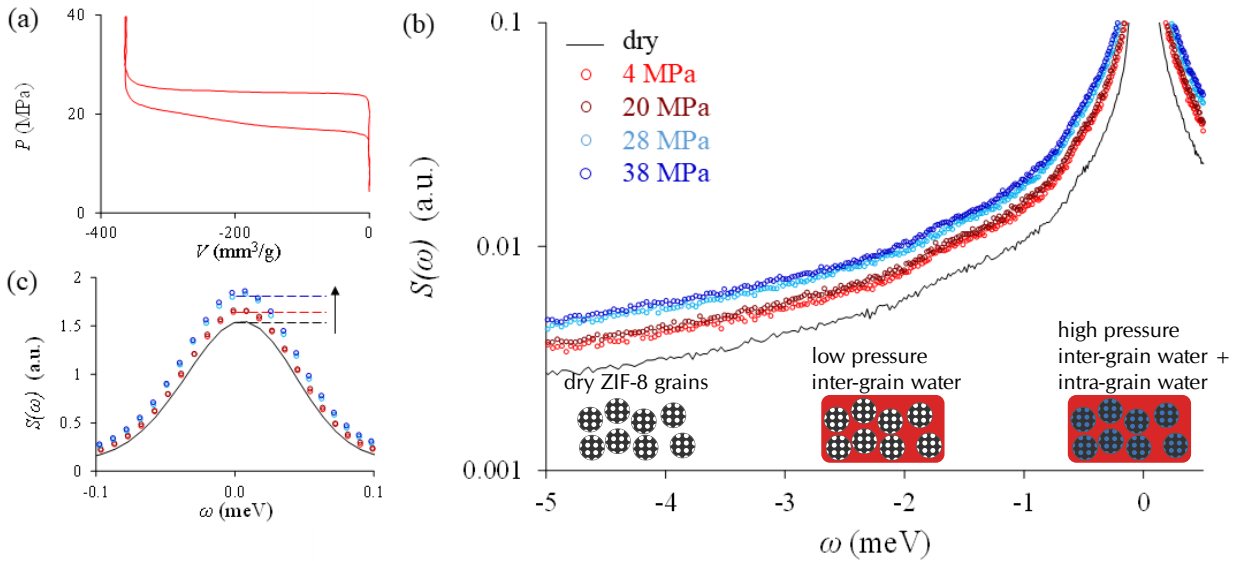


Figure 2: (a) Experimental water intrusion/extrusion cycle at room temperature in a ZIF-8 material.  $V(P)$  indicates the intruded and extruded volume  $V$  of water per g of ZIF material as a function of the exerted pressure  $P$ . (b)  $S(\omega)$  summed over  $q$  [from 0.4 to 2  $\text{\AA}^{-1}$ ] 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.

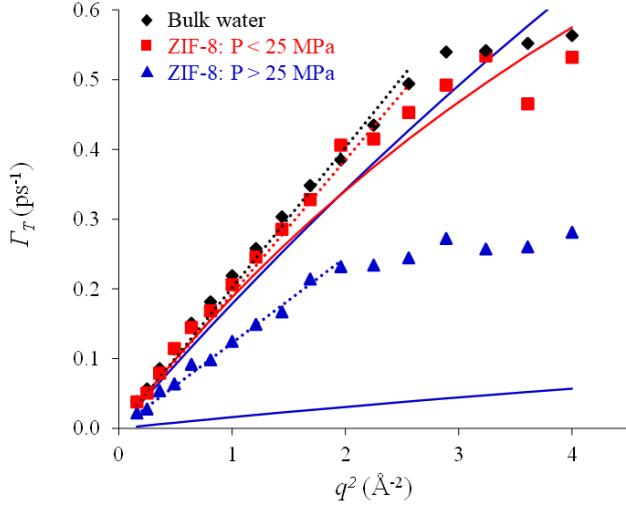


Figure 3: Half-width at half-maximum  $\Gamma_T$  of the translational contribution versus  $q^2$  for bulk water (black), low-pressure water in contact with ZIF-8 (red) and high-pressure water intruded in ZIF-8 (blue). Error bars are smaller than the symbol size. The symbols refer to the results obtained 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.

ulation, to a linear Fickian regime that extends much beyond  $q^2 \sim 2 \text{ \AA}^{-2}$ . 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  $\Gamma_T$ , 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.  $D_{T_{\text{Fick}}}$  was calculated with the Fick’s law on the appropriate  $q$ -range.**

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

We performed therefore a deeper analysis based on a formal description of water translation through the use of the jump diffusion model<sup>33</sup> coupled to isotropic rotational diffusion in a sphere of radius  $R$  (equal to the OH distance in a water molecule  $\sim 1 \text{ \AA}$ ). Each population will then be described by the new fitting model:

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

with

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

where  $\Gamma_T$  is defined within the jump-diffusion model:

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

with  $\tau$  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  $(q, \omega)$  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-

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 ( $< 40 \text{ MPa}$ ).<sup>36</sup> More information on this analysis and curve fitting quality are given in the Supporting Information (S2). The obtained  $q^2$  dependence of  $\Gamma_T$  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  $q^2 < 3 \text{ \AA}^{-2}$ . 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  $295 \text{ K}$ <sup>37</sup> (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  $q^2 \sim 2 \text{ \AA}^{-2}$  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 popu-

**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	$D_{T_{\text{jump}}} \times 10^{-5}$ [cm <sup>2</sup> /s]	$\tau$ [ps]	$D_R$ [ps <sup>-1</sup> ]
ZIF-8: P<25 MPa	$2.09 \pm 0.05$	$0.55 \pm 0.01$	$1.83 \pm 0.01$
ZIF-8: P>25 MPa: Population 1	$1.87 \pm 0.02$	$0.25 \pm 0.05$	$2.18 \pm 0.04$
ZIF-8: P>25 MPa: Population 2	$0.167 \pm 0.007$	$2.55 \pm 1.6$	$0.17 \pm 0.01$

tion, despite its limited proportion, becomes a leading term which contributes to the plateau observed for  $\Gamma_T$  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  $1.7 \times 10^{-10}$  m<sup>2</sup>/s is expected. This analysis of experimental data confirms, at long time, the ten-fold 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.

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  $q^2 t$  reveals that the rotational contribution is restricted to a very short time scale. This is further confirmed by the large value of  $D_R \sim 1$  ps<sup>-1</sup> 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  $\exp(-D_{T_{\text{jump}}} q^2 t)$  is characterized by a diffusion coefficient  $D_{T_{\text{jump}}}$  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,  $D_{T_{\text{jump}}}$  for the fast population remains four times smaller than that for bulk water; this is a

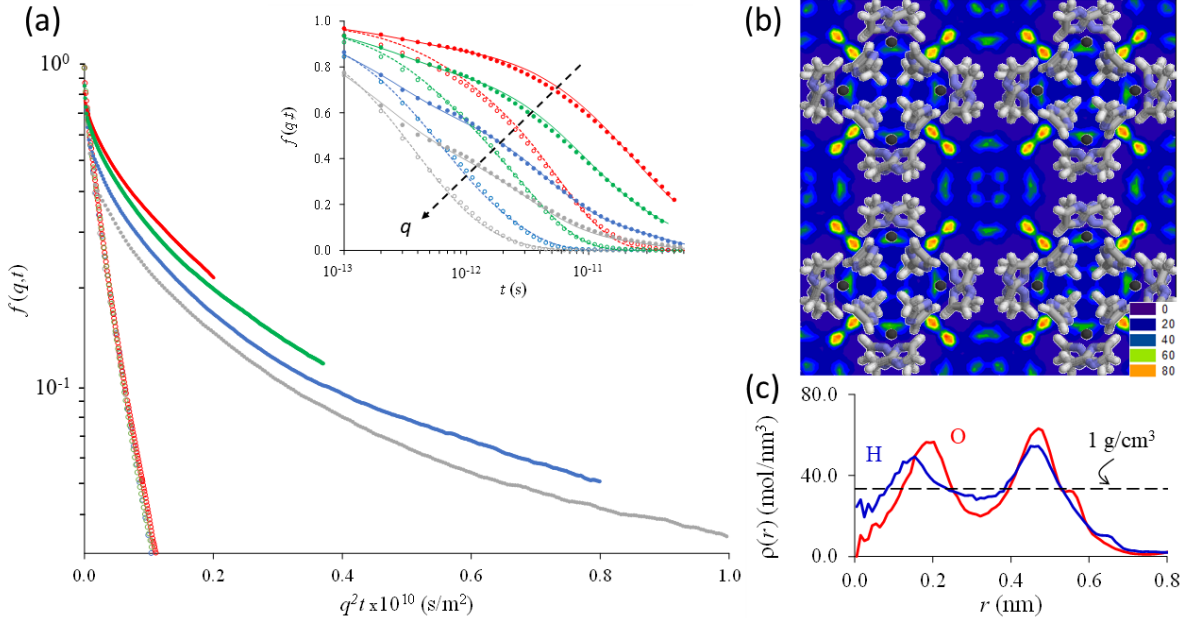


Figure 4: (a) Self (incoherent) intermediate scattering function  $f(q, t)$  as a function of  $q^2 t$  for different scattering vector  $q = |\mathbf{q}|$  for bulk water (open circles) and for water confined in ZIF-8 (full dots). The color code is as follows:  $q = 0.7 \text{ \AA}^{-1}$  (red),  $q = 1.0 \text{ \AA}^{-1}$  (green),  $q = 1.5 \text{ \AA}^{-1}$  (blue), and  $q = 2.0 \text{ \AA}^{-1}$  (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  $D_{T_{\text{jump}}} = 4.2 \times 10^{-9}$  m<sup>2</sup>/s,  $\tau = 0.9$  ps and  $D_R = 0.8$  ps<sup>-1</sup>. Full lines, confined water: fit with a fast and a slow population (ratio 60/40) with  $D_{T_{\text{jump}}} = 1.1 \times 10^{-9}/3.4 \times 10^{-10}$  m<sup>2</sup>/s,  $\tau = 0.25/20$  ps and  $D_R = 2/0.1$  ps<sup>-1</sup> (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.)

369 major difference with respect to experimental results. The  
 370 rotational diffusion coefficient of the slower population being  
 371 20 times smaller than the fast population, in qualitative  
 372 agreement with experimental results, the rotational contribu-  
 373 tion extends to much longer time than for bulk water and  
 374 contributes to the separation of the curves observed in Fig-  
 375 ure 4a at large  $q^2t$  values. Interestingly, the functions  $f$  do  
 376 not present a linear asymptotic trend at long  $q^2t$  values that  
 377 would be expected in the presence of two distinct water popu-  
 378 lations only. This means that such a basic yet efficient de-  
 379 composition to quantitatively analyze experimental and num-  
 380 erical data does not fully reflect the heterogeneity of wa-  
 381 ter dynamics in such complex materials partly controlled by  
 382 residence times at surfaces (see Bousige *et al.*<sup>38</sup> for a recent  
 383 discussion on intermittent diffusion in complex nanoporous  
 384 media).

385 At the microscopic level, the strong variations of water  
 386 density as obtained from GCMC simulation (see Figure 4b  
 387 and Figure 4c) reveal and confirm a complex heterogeneous  
 388 water structure within the ZIF-8 porosity. These data, which  
 389 are consistent with previous molecular simulation in the  
 390 field,<sup>22</sup> show that water is adsorbed on specific sites, close  
 391 to the ligands at a distance about 0.4 - 0.5 nm from the pore  
 392 center [see marked density spots in Figure 4b], and in less ad-  
 393 sorbing sites located closer to the pore center [see less marked  
 394 density spots at a distance  $\sim 0.2$  nm]. Although these adsorb-  
 395 ing sites are hydrophobic enough to not trap water at ambi-  
 396 ent pressure, their relative affinity for the intruded water con-  
 397 tribute to the marked slowing down its dynamics observed at  
 398 long time.

399 For times much longer than the residence time, exchanges  
 400 between the different adsorbing sites contribute to the com-  
 401 plex heterogeneous dynamics and hence to the effective be-  
 402 havior accessible experimentally. Such a contribution from  
 403 this molecular intermittence can be investigated by means  
 404 of the diffusion propagator  $G(r, t)$  (inverse space and time  
 405 Fourier transform of  $S(q, \omega)$ ) which corresponds to the prob-  
 406 ability that a molecule gets located at a position  $r$  at a time  $t$   
 407 while being at  $r = 0$  at a time  $t = 0$ . At short times ( $t < 10$  ps),  
 408  $G(r, t)$  shown in Figure 5a for both O and H atoms of con-  
 409 fined water is monomodal and reflects local displacements  
 410 ( $r \sim 0.1$  nm) which are similar to those found for bulk water  
 411 on the same time scale (although of a slightly smaller ampli-  
 412 tude in confinement). For larger times ( $t > 10$  ps), the  
 413 propagator becomes bimodal for O and H atoms, therefore  
 414 indicating two characteristic translational modes coexisting  
 415 on the same time scale. We show in the insert of Figure 5a  
 416 the propagators for the different water populations confined  
 417 in ZIF-8 (note the use of a log scale for improved visualiza-  
 418 tion purpose). In more detail, we measured the propagator  
 419 for molecules initially located at time  $t = 0$  in the strongly  
 420 adsorbing sites close to the ZIF-8 ligands (see density con-  
 421 tour plot shown in Figure 4b) and that for molecules ini-  
 422 tially located in the cavity center not directly in contact with  
 423 ZIF-8 ligand/surface. Such data confirm that the fast dif-  
 424 fusion mode (i.e. subpeak in the propagator located at dis-  
 425 tance  $r \sim 3.5$  Å) corresponds to molecules located within the  
 426 cavity center while the slow diffusion model corresponds to  
 427 molecules initially located in the strongly adsorbing sites. In-  
 428 terestingly, for both contributions, some molecules undergo  
 429 displacements corresponding to the other mode. This is due  
 430 to the fact that, during the characteristic time  $t = 10$  ps con-  
 431 sidered here, some molecules leave the cavity center to mi-  
 432 grate towards the strongly adsorbing sites and reciprocally.

433 Additionally to the analysis of the translational motions,

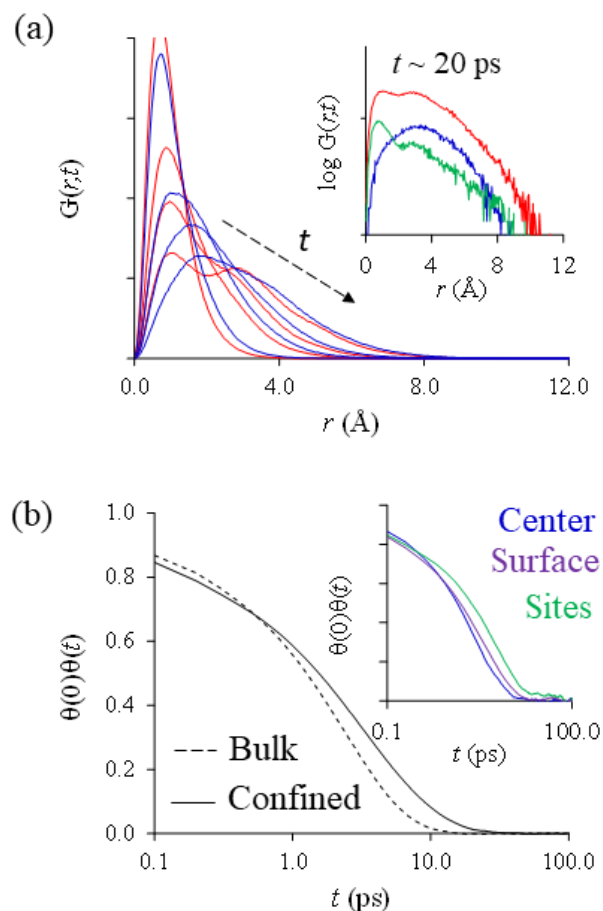


Figure 5: (a) Diffusion propagator  $G(r, t)$  as obtained using molecular dynamics at room temperature for the O (red lines) and H (blue lines) atoms of water confined in ZIF-8. The dashed line is a guide to the eye to indicate that the data are estimated at increasing times  $t = 1, 5, 10$  and  $20$  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).

434 we extracted the rotational relaxation time distribution for  
 435 water in the different regions of the ZIF-8 cavities to clarify  
 436 more specifically the impact of the surface heterogeneities<sup>39</sup>  
 437 (see Figure 5b). The timescale of rotational motions enables  
 438 indeed a more local analysis since molecules, over such a  
 439 short timescale, do not significantly exchange between dif-  
 440 ferent sites. Owing to their intrinsically slow dynamics, the  
 441 rotational motion of the molecules located in the strongly ad-  
 442 sorbing sites is slower than for molecules in the pore center  
 443 again with a typical rotational relaxation time a few times  
 444 larger than for water in the pore center. Such a slow rota-  
 445 tional motion only applies to molecules in the strong adsorb-  
 446 ing sites. For molecules at the ZIF-8 cavity surface, away

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

## 450 Conclusion

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

## 479 Supporting Information Available

480 The following files are available free of charge.

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

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