Water Nano-Diffusion through the Nafion Fuel Cell Membrane.

Baptiste Gilois
Institut de Physique de Rennes, IPR, CNRS-Université de Rennes 1, UMR CNRS 6251, 35042 Rennes, France

Florent Goujon
Université Clermont Auvergne, CNRS, SIGMA Clermont, Institut de Chimie de Clermont-Ferrand (ICCF), F-63000 Clermont-Ferrand

Alexandre Fleury, Armand Soldera
Laboratory of Physical Chemistry of Matter (LPCM), Department of Chemistry, Université de Sherbrooke, Sherbrooke, Québec, Canada, J1K 2R

Aziz Ghoufi*
Institut de Physique de Rennes, IPR, CNRS-Université de Rennes 1, UMR CNRS 6251, 35042 Rennes, France

Abstract

Nafion, an amphiphilic polymer based on fluorocarbon backbones and acid groups, is probably the most widely used fuel cell membranes. According to its hydration level it self-organizes leading to nano-cavities through which water diffuse. The diffusion of water that controls the protonic transport is then central in the conversion of chemical energy to the electrical one. Recently, a sub-diffusive and heterogeneous dynamics of water were experimentally evidenced paving the way for more efficient fuel cell membranes. Fundamentally, this dynamics which occurs at the nanoscale needs to be microscopically understood. Molecular dynamics simulations are thus performed to locally examine the water dynamics and its relation with the water and Nafion structure. The sub-diffusive regime is numerically corroborated and two sub-diffusive to diff-

*Corresponding author
Email address: aziz.ghoufi@univ-rennes1.fr (Aziz Ghoufi)
fusive transitions are found. The first is time dependent whereas the second is rather water uptake dependent. The sub-diffusive regime is ascribed to the water molecules strongly anchored close to the acid groups. We show that the sub-diffusive to diffusive time transition is the result of the elapsed time before to escape from the attractive interactions of the acid groups. The diffusive regime is recovered far from the acid groups in a homogeneous water phase that is the result of the percolation of the hydrogen bonding network. The progressive transition between sub-diffusive to diffusive regime as a function of the hydration level is due to the respective weight of diffusive dynamics that increases with respect to the sub-diffusive regime given the increase in diffusive pathways as the expense of the localized dynamics. Close to the fluorocarbon backbones the dynamics of water is also sub-diffusive but faster whereas time dynamical transition is not observed. Furthermore we highlight the existence of water corridors based on the hydrogen bonds between molecules forming single file structure in line with a sub-diffusive dynamics. These water corridors are thus possible conducting pathways of protons in a Grotthuss mechanism. Structurally we depict an interdigitated structure where the sulfonate are interleaved. Eventually, at high water uptake, we exhibit the self-organizing of Nafion leading to a phase separation between water and the Nafion membrane. We establish a specific interfacial organization of the hydrophilic sulfonate groups involving a water/Nafion interface.

Keywords: Nafion, Water diffusion, molecular simulation, anomalous dynamics

2010 MSC: 00-01, 99-00

1. Introduction

Nowadays the development of new sources of energy is one of major challenges of our society. To overcome this penury due to the intensive exploitation of our natural resource, new approaches have to be promoted in a context of sustainable development. One of the keys of this energetics problem could be
the improvement and the use of fuel cells. These are compact, light and quiet with less pollution emission. Additionally, fuel cells incorporate solid materials inducing an easy assembly and are capable to deliver a strong power with a large current density and a very good energetics performance.

Fuel cells are electrochemical devices allowing a direct conversion of the chemical energy provided from a fuel as hydrogen, methanol or hydrocarbons in electrical energy. As illustrated in Figure 1 fuel cells are built from two electrodes separated from an electrolytic ionic conductor. According to the operating temperature and then to the application (mobile or stationary applications, high or short power supplied), the electrolyte will be different. Two classes of fuel cells can be defined i) fuel cells with an operating temperature beyond 250°C such as the solid oxide fuel cell or the Molten-carbonate fuel cell and ii) fuel cells with an operating temperature below 250°C such as Proton Exchange Membrane Fuel Cell (PEMFC) or Direct-methanol fuel cell.

PEMFC are essential given their high performance, no emission in CO₂ and an use in a large range of power. Several studies have been devoted these two past decades to reduce the cost and to increase their lifetime to allow a large commercialization. Suitable functioning needs are: i) a diffusion of gases and condensation/vaporization of water into the pores of the active layers and diffusion layers, ii) an electronic transport and electrochemical conversion of the electrons, protons and reactive gases in the active layer of the electrodes and iii) transport of protons through the fuel cell membrane. The current membranes are based on perfluoro-sulfonic membranes and are built from fluorocarbon backbones and acidic groups. The oldest, most studied and used membrane is probably the one manufactured from Dupont de Nemours (US); the Nafion membrane.

Structure of hydrated Nafion is strongly investigated in order to improve its fuel cell performance but also to design new membranes. Structural studies have highlighted a complex and multi-scale structure. Additionally, a phase segregation between hydrophobic and hydrophilic moieties of Nafion was evidenced during the hydration process. One of the most impor-
tant properties of Nafion is its conductivity that is strongly connected to the hydration level and the water transport into the membrane. Indeed, as shown Figure 1, the water uptake progressively increases due to the reactivity between the protons and O₂ involving formation of water molecules. Therefore, an understanding of the mechanisms of conduction as a function of the hydration level through the membrane is crucial to design new performing membranes. Transport of protons through the Nafion membrane is then possible through the Grotthuss mechanism [13] and a vehicular one [14] implying chemical reactivity. Although this reactivity is crucial in both mechanisms, the role of water structure and the topology of the Nafion membrane, that are strongly connected to the hydration level, must be clarified because they control the conduction pathway allowing the protonic transport.

Due to progressive water filling three states of hydration can be defined, i) at low hydration (3 water molecules per sulfonate group) where water molecules solvate sulfonate groups of Nafion and protons leading to a localized water structure known as bounded (B) water, ii) at the intermediate hydration (20 water molecules per sulfonate group) such that water molecules are still close to H⁺ and SO₃⁻ but less anchored and can be considered as loosely bounded (LB) and iii) at high water loading (beyond 20 water molecules per sulfonate group) where the water molecules far from H⁺ and SO₃⁻ are then considered as free (F) [15, 16, 17]. Additionally, it has been evidenced that Nafion membrane self-assembly upon hydration leading to a peculiar topology based on channels network involving specific water conduction pathways (between channels) [6, 10, 11, 12]. An understanding of the water network and the water transport into Nafion is then essential to well apprehend the protonic conduction.

As previously evoked Nafion is a multi-scale material and the physical phenomena ruling the water transport occur from nanoscale to macro-scale [9]. Interestingly, it is now well established that mechanisms at the nanoscale strongly impact the macroscopic properties and it is then necessary to well characterize and understand the water structure and dynamics at the nanoscale. Since 50 years molecular simulations has became an useful method to capture the micro-
scopic details and to connect them with macroscopic properties using statistical physics. Several numerical and experimental works have been performed to unveil the water diffusion and the so-associated mechanism as well as the atomistic structure \[18, 19, 20, 21, 22, 23, 24, 25, 26, 12\]. Indeed, by using Nuclear Magnetic Resonance (NMR) pulsed gradient spin experiments, C. Casieri et al. have shown a sub-diffusive dynamics of confined water at the micrometer scale for temperatures \[>320\text{K}\] an at different water uptakes \[27\]. At the same time, D.T. Hallinan et al. highlighted a non-Fickian diffusion of water in Nafion from time-resolved Fourier transform infrared-attenuated \[21\]. More recently, Mabuchi and Tokumasu have also established a sub-diffusive dynamics of protons and water at low water contents below 5 ns \[28\]. According to these authors, from 5 ns the diffusion became diffusive and the self-diffusivity can be calculated. Devanathan et al. have also investigated the the relation between structure and dynamics of water and ions. By using classical molecular dynamics simulation these authors have evidenced that at low hydration levels, coordination of hydronium ions by multiple sulfonate groups prevents vehicular transport and impedes structural transport of protons through steric hindrance to hydration of the hydronium ions. \[29\] Devanathan et al. have also highlighted from quantum hopping molecular dynamics the existence of the water network percolation in Nafion and its effect on proton conductivity. \[29\].

Although these works provide insights into translational dynamics, the microscopic mechanisms and the relationship between the atomistic structure of the Nafion and dynamics of water were slightly examined. However, recently Song et al. have depicted that the surface chemistry of Nafion strongly impacts the water and ions transport. Indeed, using molecular spin probes that selectively partition into heterogeneous regions of Nafion and Overhauser dynamic nuclear polarization relaxometry, Song and co-workers have revealed that both water and proton diffusivity are significantly faster near the fluorocarbon and the acidic groups lining the water channels compared to within the water channels \[25\]. More recently, Q. Berrod et al. have also studied water motion and ion transport by combining Quasi Elastic Neutron Scattering, Pulsed Field
Gradient NMR, and atomistic simulations. They demonstrated that confinement at the nanoscale and the direct interactions with the charged interfaces produce anomalous sub-diffusion, due to a heterogeneous space-dependent dynamics within the ionic nano-channels [9].

In this work, we aim to investigate the translational diffusion of water at the nanoscale and to capture the microscopic processes ruling it. To do so molecular dynamics (MD) simulations of hydrated Nafion were carried out for different hydration states.

Figure 1: Illustration of the operating fuel cell membrane.
2. Methodology

2.1. Models and Computational Details

Nafion fuel cell membrane is a dense polymer and its molecular modeling needs specific construction to reach a relaxed configuration. To do so, Nafion membrane was built by considering 4 chains of 682 atoms of 10 monomers (see Figure 2a) by using a Monte Carlo construction based on the statistical bias developed by Theodorou & Seuter [30]. In a second time, we equilibrated the empty polymer (dehydrated Nafion polymer) using the 21 MD steps scheme proposed by Hofmann et al. [31]; Within this scheme, seven cycles of three MD simulations were performed. (1) NVT, T=600 K; (2) NVT, T=300K and (3) NpT, T=300K, where N is the number of atoms, V the volume and T the temperature. Details of each steps are provided in Table 1. The value of the pressure in the NpT simulations was gradually increased, passing from the first to the third cycle up to \( p_{\text{max}} = 50 \text{ kbar} \), and then was reduced stepwise until it reached the final value of \( p = 1 \text{ bar} \). In a last step, membrane was subsequently hydrated and 7 hydration levels were defined from the number of water molecules around a sulfonate group (\( \lambda \)), \( \lambda = 3, 5, 7, 9, 11, 14.5, 22.5 \). All hydrated membranes were relaxed from 21 MD steps scheme by using similar temperature and pressure that ones used in case of the empty polymer. Hydration was performed by randomly inserted water molecules within the relaxed empty polymer and equilibration of hydrated Nafion in statistical NpT ensemble was carried out at 1 bar and 300 K. During all processes polymer was considered as flexible.

Interestingly, isotherm of adsorption and water activities could be extracted from molecular dynamics simulations in the Grand Canonical statistical ensemble in order to consider the flexibility of the polymer. However, this task needs additional and specifics developments to carry out Grand Canonical Molecular Dynamics simulations such as the methodology of hybrid osmotic Monte Carlo simulation [32]. That is in progress and is out of the scope of this work where the water uptake is fixed.

Nafion was modeled by considering the intramolecular and intermolecular
Table 1: Details of the thermodynamic conditions for the 21 MD steps used in the process of generation of relaxed Nafion polymer.

<table>
<thead>
<tr>
<th>Step</th>
<th>ensemble</th>
<th>T (K)</th>
<th>p (kbar)</th>
<th>Time (ps)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>NVT</td>
<td>600</td>
<td></td>
<td>50</td>
</tr>
<tr>
<td>2</td>
<td>NVT</td>
<td>300</td>
<td></td>
<td>50</td>
</tr>
<tr>
<td>3</td>
<td>NpT</td>
<td>300</td>
<td>1</td>
<td>50</td>
</tr>
<tr>
<td>4</td>
<td>NVT</td>
<td>600</td>
<td></td>
<td>50</td>
</tr>
<tr>
<td>5</td>
<td>NVT</td>
<td>300</td>
<td></td>
<td>100</td>
</tr>
<tr>
<td>6</td>
<td>NpT</td>
<td>300</td>
<td>30</td>
<td>50</td>
</tr>
<tr>
<td>7</td>
<td>NVT</td>
<td>600</td>
<td></td>
<td>50</td>
</tr>
<tr>
<td>8</td>
<td>NVT</td>
<td>300</td>
<td></td>
<td>100</td>
</tr>
<tr>
<td>9</td>
<td>NpT</td>
<td>300</td>
<td>50</td>
<td>50</td>
</tr>
<tr>
<td>10</td>
<td>NVT</td>
<td>600</td>
<td></td>
<td>50</td>
</tr>
<tr>
<td>11</td>
<td>NVT</td>
<td>300</td>
<td></td>
<td>100</td>
</tr>
<tr>
<td>12</td>
<td>NpT</td>
<td>300</td>
<td>25</td>
<td>5</td>
</tr>
<tr>
<td>13</td>
<td>NVT</td>
<td>600</td>
<td></td>
<td>5</td>
</tr>
<tr>
<td>14</td>
<td>NVT</td>
<td>300</td>
<td></td>
<td>10</td>
</tr>
<tr>
<td>15</td>
<td>NpT</td>
<td>300</td>
<td>5</td>
<td>5</td>
</tr>
<tr>
<td>16</td>
<td>NVT</td>
<td>600</td>
<td></td>
<td>5</td>
</tr>
<tr>
<td>17</td>
<td>NVT</td>
<td>300</td>
<td></td>
<td>10</td>
</tr>
<tr>
<td>18</td>
<td>NpT</td>
<td>300</td>
<td>0.5</td>
<td>5</td>
</tr>
<tr>
<td>19</td>
<td>NVT</td>
<td>600</td>
<td></td>
<td>5</td>
</tr>
<tr>
<td>20</td>
<td>NVT</td>
<td>300</td>
<td></td>
<td>10</td>
</tr>
<tr>
<td>21</td>
<td>NpT</td>
<td>300</td>
<td>0.001</td>
<td>2000</td>
</tr>
</tbody>
</table>

DREIDING force field [33] that was already previously validated [34, 2]. H$_3$O$^+$ and water molecules were modeled from the model of Baaden [35, 36] and from
the TIP4P-2005 force field [37], respectively. The interactions between the polymer, water and proton were taken as a combination of the electrostatic and van der Waals (VdW) interactions. VdW contributions were modeled by considering the Lennard-Jones potentials, such that cross-interactions were calculated by means of the Lorentz-Berthelot mixing rules. The electrostatic interactions were computed by considering the Ewald sum [38, 39]. All interactions were truncated by using a cutoff of 12 Å. All of the simulations were performed using the DLPOLY software [40]. Verlet-velocity propagator combined with the Nose-Hoover thermostat and barostat were here considered [41]. Relaxation times of 0.5 ps of both thermostat and barostat were used. Acquisition phases were conducted during 20 ns with a timestep of 1 fs while the equilibration phases were lasted 50 ns. To ensure that the systems are mechanically equilibrated the local pressures of three components according to \( z \) direction were calculated. Indeed, the mechanical equilibrium is reached when the local pressures such as \( P_x(z) \), \( P_y(z) \) and \( P_z(z) \) are constant as a function of \( z \) [42]. Pressure were calculated on 20 points by using the Irving-Kirkwood method \([\text{]}\) and the averages on these 20 points with their uncertainties were calculated. We found thus the following values \( <P_x(z)> = 0.12 \pm 0.02 \) Mpa; \( <P_y(z)> = 0.16 \pm 0.02 \) Mpa and \( <P_z(z)> = 0.09 \pm 0.02 \) Mpa in good agreement with the input pressure of 0.1 Mpa. These small uncertainties show that the pressure is well converged, confirming that the simulated systems have reached a mechanical equilibrated state. These verifications were also performed for the hydrated membranes where the uncertainties were found slightly higher around 0.05 Mpa that is sufficient to conclude on the mechanical equilibrium. Let us mention that these values of uncertainties are the same order of magnitude than those found in case of liquid [43, 42].

We report in Figure 2b the density of the hydrated membrane as a function of \( \lambda \). As pointed out in Figure 2b a fair agreement with experiment was obtained that validated the so-used force fields of Nafion, proton and water and their combining.
2.2. Translational dynamics

Translational dynamics was studied from the calculation of the mean square displacement (MSD)

\[ \text{MSD}(t) = \frac{\sum_{t_0} \sum_{i=1}^{N} [\mathbf{r}_{\text{com},i}(t + t_0) - \mathbf{r}_{\text{com},i}(t_0)]^2}{NN_0t} \]  

with \( \mathbf{r}_{\text{com},i} \) the position of the centre of mass of molecule \( i \), \( t_0 \) the time origin, \( N \) the number of molecules and \( N_0 \) the number of \( t_0 \). From the MSD and by using the generalized Einstein’s relation, the diffusivity \( (D_f) \) is then extracted such that

\[ \text{MSD}(t) = D_f t^\alpha \]  

\( \alpha \) is a parameter related to the type of diffusion, \( \alpha = 1 \) corresponds to a diffusive regime, \( \alpha < 1 \) is connected to a sub-diffusive one while \( \alpha > 1 \) is linked to a super-diffusive process. The case of \( \alpha = 2 \) corresponds to a ballistic dynamics. Let us mention, that for \( \alpha \neq 1 \), the diffusion is considered as anomalous. For a diffusive (Fickian) regime the diffusivity is related to well
known diffusion coefficient (D) extracted from the Einstein’s relation in one dimension \((MSD(t) = 2Dt)\). \(\alpha\) was calculated by deriving Eq. 2 as

\[
\alpha(t) = \frac{\partial MSD(t)}{\partial t} \cdot \frac{MSD(t)}{t}
\]

(3)

3. Results and Discussion

We report in Figure 3 the instantaneous value of \(\alpha\) as a function of time \((t)\) for all water uptakes. Figure 3a shows that for \(t < 1\) ns, \(\alpha < 1\) that suggest a sub-diffusive regime in this time range. As highlighted in Figure 3b, beyond 1 ns, \(\alpha\) rises to one leading to a crossover between a sub-diffusive to a diffusive regime (or a lesser sub-diffusive regime). Interestingly, this effect seems to widen as the water uptake increases to strive for a quasi diffusive regime. At low \(\lambda\), we can assume that water molecules are probably organizing in clusters leading to the formation of nanophases where molecules locally diffusive (because they are trapped in small size environment) inducing a sub-diffusive regime. Indeed, microscopically speaking sub-diffusive regime is the result of the local diffusion and a heterogeneous structure. In second stage size of nanophases could growth as \(\lambda\) increases involving a progressive transition toward a diffusive regime. The water structure and its connection with the dynamics will be deeply discussed later.

We report in Figure 4a the average value of \(\alpha\) \(< \alpha >\) as a function of \(\lambda\) calculated between 0 and 1 ns. As shown in Figure 4a, the sub-diffusive dynamics is clearly evidenced \(< \alpha > < 1\). Additionally, Figure 4a shows an increase in \(< \alpha >\) as the water uptake increases that highlight a continuous dynamical transition. To well appreciate the loss of the sub-diffusive regime beyond 1 ns, \(< \alpha >\) was calculated between five consecutive time intervals noted \(\Delta t_i\) \((\Delta t_i =1\) between 0 and 1 ns; \(\Delta t_i =2\) between 1 and 5 ns; \(\Delta t_i =3\) between 5 and 10 ns; \(\Delta t_i =4\) between 10 and 15 ns and \(\Delta t_i =5\) beyond 15 ns\). Figure 4b displays \(\alpha\) as a function of \(\Delta t_i\) for six water contents. Figure 4b exhibits a progressive increase of \(< \alpha >\) as a function of \(\Delta t_i\) for all loadings that is in line with previous
conclusions drawn from instantaneous values of $\alpha$. This effect is found higher at low loadings because from $\Delta t_i = 3$, $<\alpha>$ is doubled. The diffusion coefficients were extracted from the linear region of the MSD related to the diffusive regime and from the total MSD trajectories (from 0 to 20 ns) and reported in Figure 3a. When the diffusion coefficient is evaluated from all MSD trajectories a fair agreement between experiment [46] and simulation was found. Indeed tendency as a function of $\lambda$ and absolute values are well reproduced. Experimentally it was not possible to de-correlate both regimes. Interestingly, when only the diffusive regime is considered the so-calculated diffusion coefficients are found weaker suggesting that molecular simulations could be a relevant tool to be combining with NMR or QENS experiments to extract the diffusion coefficient of confined liquids. To sum up; i) the sub-diffusive regime could be due to the presence of nanophases inducing localized dynamics and ii) the loss of the sub-diffusive regime could be the result of to spatial connection between nanophases inducing the diffusive pathways. We first began then to examine the existence of nanophases and their structure.

Figure 3: $\alpha$ as a function of time for all hydration levels in all range of time a) and between 1 ns and 15 ns b).
Water nanophases can be compared to the water pockets and can be assimilated to the clusters defined from their size. By using the method of Stoddard [47] the cluster size distribution was calculated by considering the hydrogen bonds (HB). HB were defined from the geometric criteria established by Luzar and Chandler [48] that consist in the distances of 2.5 Å and 3.5 Å between the hydrogen and the oxygen atoms and two oxygen atoms of two water molecules. We report in Figure 5a the cluster size distribution of water as a function of \( \lambda \). At low water uptake (\( \lambda = 3 \) and 5) all cluster sizes are equally sampled and the system does not percolate. This result sheds light on a polydisperse distribution of clusters and then the presence of small water islands. The percolation occurs from \( \lambda = 7 \) where all nanophases begin to be gathered in a single interconnected structure. This structural organization is in line with the so-observed transition between sub-diffusive to diffusive regime. Interestingly, despite the percolation of the HB network that occur at \( \lambda = 7 \), small clusters are also evidenced suggesting a mixing between the single interconnected structure and clusters, that is in good concordance with the so-observed sub-diffusive and diffusive regime. To provide an obvious picture of time evolution of clusters, the cluster size distribution was evaluated for each \( \Delta t_i \). As shown in Figure 5b, for \( \lambda = 3 \) the

Figure 4: a) \( <\alpha> \) (left axis) and translational diffusion coefficient (right axis) as a function of \( \lambda \). b) \( <\alpha> \) as a function of \( \Delta t_i \). Experimental diffusion coefficient were taken in Ref. [46].
cluster size distributions is not time-dependent that suggest an absence of cluster interconnection and an absence of clusters assembly with respect to the time. Therefore, at low $\lambda$, water molecules are trapped in the nanophases and locally diffusive in small size environment inducing a sub-diffusive regime.

![Graphs showing cluster size distributions for different hydration levels.](image)

Figure 5: a) Number of clusters as a function of the cluster size for all hydration levels. Number of clusters as a function of the cluster size for $\lambda = 3$ b) and $\lambda = 9$ c). d) Enlargement of part c) for high size clusters.

To illustrate the water nanophases, snapshots of final configuration of six hydration levels are reported in Figure 6. As shown in Figure 6 at low water uptake water clusters are observed to be located close to the sulfonate groups.
whereas at higher loadings water molecules are rather connected to form a structuring network always impacted by the hydrophilic sulfonate groups and hydrophobic fluorine backbone. It would seem that water molecules firstly adsorb on the sulfonate groups and grow to form water nanophases. As $\lambda$ increases nanophases grow to finally interconnect and form a single structure. The localized dynamics are thus always present but they statistically decreases in favor to the diffusive regions (nanophases assembly). Indeed, as shown in Figure 5: and 5 where the cluster size distributions for $\lambda = 9$ is represented, the presence of small clusters and the percolation phenomenon are observed whatever $\Delta t_i$. That suggest the permanent presence of the single interconnected water structure. This result refutes the fact that the sub-diffusive to diffusive transition is due only to the time assembly of nanophases [49].

A question arises on the origin of the sub-diffusive regime: is the result of the confined dynamics into nanophases or the local diffusion close to the sulfonate groups? Another point concerns the loss of the sub-diffusive regime from $\lambda = 5$ that can be correlated to the assembly or a growth of water nanophases leading to an increase in diffusive domains with respect to the sub-diffusive ones. This scenario does not exclude the presence of a localized dynamics whose statistical weight will decrease as the water nanophases grow.

To analyze the local dynamics close to the sulfonate groups ($\text{SO}_3^-$) the characteristic distances between water and $\text{SO}_3^-$ were examined. To do so, the radial distribution functions (RDF) between hydrogen atoms of water molecules and oxygen atoms of sulfonate groups are calculated and reported in Figure 7a. As shown in this figure two maxima can be extracted whatever the water uptake, one around 2.0 Å corresponding to a strong hydrogen bond and a second distance located at 3.4 Å connected to the second hydration shell. Difference in intensity between the three water loadings is due to a density effect [50]. RDF between oxygen atoms and fluorine atoms were also calculated to investigate the dynamics along the hydrophobic backbone. Contrary to the sulfonate groups, the first maximum is found at 4 Å for both types of fluorine (noted F1 and F2
Figure 6: Snapshots of final configurations of 6 hydration levels. Water molecules are represented such that red and white colors represent the oxygen and the hydrogen atoms, respectively. Yellow spheres are the sulfur atoms in the Nafion membrane. For clarity, other atoms of Nafion are not represented.

In Figure 2, whereas the position of the second peak changes as a function of type of fluorine. This difference is due to the proximity of the sulfonate groups. Indeed as illustrated in Figure 2, F2 is close to the sulfonate groups and then benefit from their hydrophilicity. From RDF, three regions around the sulfonate groups were drawn, < 2.0 Å corresponding to the bounded water, between 2.0 Å and 3.4 Å corresponding to the loosely bounded water and beyond 3.4 Å that is the free water. Similar decomposition was performed around the fluorine. Local dynamics have been then investigated by considering this spatial division.
Figure 7: a) Radial distribution functions between oxygen atoms of SO₃⁻ groups of the Nafion and the hydrogen atoms of water for three water uptakes. RDF between F1 atoms b) and F2 atoms c) (labels are defined in Figure 2) and oxygen atoms of water. d) Illustration of the so-considered distances, based on the RDF calculation, to define three regions around the sulfonate groups; bounded water (B), loosely bounded water (LB), and free water (F).

We report in Figure 8 the MSD of four water uptakes from \( \lambda = 3 \) to \( \lambda = 22.5 \) in three bounded (B), loosely bounded (LB) and free (F) water regions. As highlighted in Figure 8a, a sub-diffusive regime is observed in the three zones because all water molecules are anchored close to the sulfonate groups (see Figure 6 and Figure 7a). In both LB and F boundaries the sub-diffusive regime is progressively replaced by the diffusive one as the water uptake increases. That is the result of the assembly of nanophases allowing a diffusive regime through the membrane (see Figure 6). That is an evidence of the connection between the percolation process (Figures 5 and 6) previously highlighted and the water diffusion. Indeed, we have shown that the diffusion of water molecules increases.
with the water uptake i.e. with the rate of percolation and the growth of the hydrophilic part in line with the work of Benziger et al. who showed that the different percolation thresholds suggest the hydrophilic domains in Nafion grow from lamella. However as depicted in Figure 8c and 8d, the dynamics of water in the B region is still sub-diffusive during about 50-100 ps prior to becoming diffusive. The macroscopic sub-diffusive regime previously characterized by $\alpha$ could be then attributed to the water molecules strongly anchored close to the sulfonate groups. The time elapsed in the sub-diffusive regime could be probably attributed to the elapsed time before to escape from the attractive interactions of the sulfonate groups. Indeed, during the first hundred pico-seconds the MSD was found around 2-5 Å$^2$ that represent a displacement of 2.2 Å that is the same order of magnitude of size of region B.

As shown in Figure 9a, this phenomenon is independent of the water uptake that bear out an escaping process of water molecule close to the sulfonate groups. The elapsed time of water molecules close to the sulfonate groups can be quantified by calculating their residence times from the evaluation of the correlation function of the survival probability function ($P(t)$) that adopt a value of one if the water molecule labeled $j$ has been in the referred hydration shell around site $\alpha$ from time $t'$ to time $t + t'$, without getting out in between this time interval, and zero otherwise. This function is then fitted from a double exponential such that the residence time is extracted. We report in Figure 9b the correlation function as a function of time for all $\lambda$. As shown in Figure 9b, the decorrelation is obtained for $t \sim 50$ ps that is in line with the predicted time transition between both dynamical regimes. Residence times were calculated from an exponential adjustment using a double exponential. As reported in Figure 9b, the fit is in good concordance with $P(t)$ that suggest that the residence time can be extracted. We obtain the following residence times, $\tau_{\lambda=22.5}=84.4$ ps and $\tau_{\lambda=3}=63.7$ ps that is line with the time crossover observed in Figure 8.
Figure 8: Mean square displacements of water in three water regions B, LB and F around SO$_3^-$ for a) $\lambda = 3$, b) $\lambda = 7$, c) $\lambda = 11$ and d) $\lambda = 22.5$. 
Interestingly, as shown in Figure 10, the dynamics of water around fluorine atoms is found similar in three regions due to the hydrophobicity of carbon-fluorine backbone preventing the anchoring of water molecules. As for the water molecules close to the sulfonate groups a progressive transition between sub-diffusive to diffusive dynamics as the water uptake increases is also evidenced. Indeed, the minimum distance between the fluorine atoms and the sulfonate groups is found between 3.2 Å and 3.9 Å (from RDF calculations) that suggest that the F, LB and B water molecules of fluorine correspond to the LB and F water molecules of the sulfonate groups.

Figure 9: a) Mean square displacements of water in the region B around SO$_3^-$ for all uptakes. b) Correlation function of the survival probability function as a function of time allowing us the calculation of residence times.
This result, based on the position of the first minimum in RDF, suggests that the water molecules could form a conducting corridor between sulfonate groups as illustrated in Figure 11a. By calculating the number of hydrogen bonds [48] of water molecules we highlighted that the water corridor was based on the hydrogen bonds between molecules forming a single file structure (see Figure 11b). Interestingly, this structure is well known to involve a sub-diffusive dynamics [54] that is in line with the value of $\langle \alpha \rangle$ reported in Figure 4. This water corridor structure could be then a conducting pathway of protons in the Grotthuss mechanism [13]. Although classical molecular dynamics simulations can not take into account the transfer of the proton between water molecules, the existence of water corridors between sulfonate groups where each molecule are connected by the hydrogen bonds will be in favor of the transport of proton based on the Grotthuss process.
Figure 11: a) Illustration of a water corridor along the carbon-fluorine backbone b) Snapshot at $\lambda = 7$ highlighting a water corridor based on the hydrogen bonds between water molecules. Water molecules have been chosen by considering the hydrogen bond number using the geometric criteria defined by Luzar and Chandler [48].

Indeed, as shown in Figure 12, the MSD of hydronium ions are around 40-60 Å$^2$ whatever $\lambda$ i.e. a displacement of 6-7 Å involving a very strong anchoring around the sulfonate groups. At $\lambda = 7$, the MSD is 18.1 Å$^2$ while at $\lambda = 11.5$ the MSD is 13.5 Å$^2$. The difference between both water uptake corresponds to 4.6 Å$^2$ i.e. a displacement of 2 Å along the sulfonate groups. As this difference is not really significant the MSD of ions can be then considered as similar between $\lambda = 7$ and $\lambda = 11.5$. Figure 12 which report the RDF between hydronium ions and the sulfonate groups evidences several peaks between 2 and 6 Å that corroborate a strong interactions. This result is in good agreement with the work of Devanathan and coworkers that has shown a very slow proton dynamics in line with the neutron scattering experiments. [29]
Figure 12: a) Mean square displacements of $\text{H}_3\text{O}^+$ for all uptakes. b) RDF between oxygen atoms of $\text{SO}_3^-$ and hydrogen atoms of $\text{H}_3\text{O}^+$. 

The hydronium ions located beyond 6 Å in the second solvation shell of the sulfonate groups correspond probably to the ions located in the first shell of another $\text{SO}_3^-$. That is probably the result of the polymer folding. To ensure this hypothesis we report in Figure 13a the RDF between $\text{SO}_3^-$. Figure 13a shows three maxima at 2.5 Å, 3.0 Å and 5.0 Å that highlight an interdigitated structure. At high water uptake, Figure 13b exhibits the self-organizing of Nafion leading to a phase separation between water and the Nafion membrane. Interestingly, as shown in Figure 13b the hydrophilic sulfonate groups point toward the water phase involving a Nafion surface decorated with $\text{SO}_3^-$ interacting with water molecules. Therefore a water/Nafion interface is evidenced with a surface rich in $\text{SO}_3^-$ such that water molecules could jump between sulfonate sites. This result is in good agreement with the work of G.A. Voth and C.K. Knox because their very large scale simulations revealed fast intercluster 'bridge' formation and network percolation. Sulfonate groups were found inside these bridges and played a significant role in percolation. Sulfonates also strongly aggregated around and inside clusters leading to the hydrophilic-hydrophobic interface.
Translational jumps were quantified by calculating the self Van Hove function \( G_S(r,t) \) where \( r \) and \( t \) are the position and time of guest molecules, respectively, 

\[
G_S(r,t) = \frac{1}{N} \left\langle \sum_{j=1}^{N} \delta[r - (r_j(t) - r(0))] \right\rangle,
\]

where \( r_j(t) \) is the atomic position of \( j \) particle at time \( t \) and \( N \) is the number of water molecules. Figure 14a reports \( G_S(r,t) \) from 1 ps to 300 ps for a loading of \( \lambda = 3 \). The maximum of the Van Hove function remains at a distance lower than 1.5 Å from 2 ps to 300 ps, which indicates small displacements and an absence of translational jumps in this time interval. However a growth of a shoulder is observed around 3.0 Å that could correspond to the translational jumps between two sulfonate groups side by side.
4. Concluding Remarks

In this work diffusion of confined water into the Nafion membrane was examined. A sub-diffusive regime was evidenced whatever the hydration level. This regime was important at low $\lambda$ whereas it was found decreasing in favor of diffusive one as the hydration rate increased. A first progressive sub-diffusive to diffusive transition as a function of the water uptake was then evidenced. Structural analyses were then carried out to structurally understand this dynamical behavior. Cluster size study evidenced a polydisperse distribution of nanometric cluster inducing nanophases. The first hypothesis was that, at low water uptake, water molecules are trapped in these nanophases and locally diffusive in small size environment inducing a sub-diffusive regime. The percolation occurs from higher hydration level such that all nanophases begin to be gathered in a single interconnected structure. The progressive transition between sub-diffusive to diffusive regime as the increase in water uptake was then, in first time, ascribed to the structural process. More especially, we have shown
that the water molecules firstly adsorb on the sulfonate groups and grow to form these water nanophases around sulfonate groups. As well water uptake increases nanophases grows to finally interconnect and to form a single structure.

However, we have shown the presence of a permanent single interconnected water structure involving that the sub-diffusive to diffusive transition could be only the result of the time assembly of nanophases. Analyses of the local dynamics helped us to examine the molecular origin of this dynamics. From radial distribution calculations three regions around the sulfonate groups and the fluorine atoms were drawn, corresponding to the bounded (B) water, to the loosely bounded (LB) water and the free (F) water. Local dynamics have been then investigated by considering this spatial division. In the (B) boundary of sulfonate groups the sub-diffusive regime was always evidenced and was found progressively timely replaced by the diffusive one in all range of λ. The so-observed sub-diffusive regime was ascribed to the water molecules strongly anchored close to the sulfonate groups and locally diffusive in small size environment. We evidenced that the sub-diffusive to diffusive time transition was the result of the elapsed time before to escape of the attractive interactions of the sulfonate groups. This phenomenon was found independent of the water uptake. The diffusive regime is then possible from the diffusive corridors induced by the percolation of the hydrogen bonding network. On the contrary in both LB and F regions the dynamics were found rather diffusive. As a function of the location of the water molecules their dynamics can be then drastically different involving dynamical heterogeneities.

The progressive transition between sub-diffusive to diffusive regime as a function of the hydration level is ascribed to the respective weight of diffusive dynamics that increase with respect to the sub-diffusive regime due to the increase in diffusive pathways at the expense of the localized dynamics.

Close to the fluorocarbon backbones the dynamics of water molecules is also sub-diffusive but faster than the water located close to the sulfonate groups. Additionally, time dynamical transition was not observed. Dynamics of water molecules depends on their positions into the membranes (anchored in the
vicinity of sulfonate groups, close to the fluorocarbon backbones and in the nanometric channels) involving then dynamical heterogeneities. Furthermore we highlighted the existence of water corridors based on the hydrogen bonds between molecules leading to the formation of a single file structure which is in line with a sub-diffusive dynamics. This water corridor structure is assimilated to a possible conducting pathway of protons in a Grotthuss mechanism. From a Nafion standpoint, we exhibited an interdigitated structure where the sulfonate groups are interleaved. At high water uptake, we exhibit the self-organizing of Nafion leading to a phase separation between water and the Nafion membrane. Eventually, a specific interfacial organization was evidenced involving a hydrophilic surface and then a water/Nafion interface. At the microscopic scale, water molecules diffuse then close to the sulfonate groups (hydrophilic regions) and slide along the Fluorine groups (hydrophobic zones). However, at the macroscopic scale (corresponding to the NMR results) the water transport corresponds to the diffusion through the hydrophilic nanophases defined by the sulfonate groups leading to a hydrophilic surface. That suggests then an interfacial transport at the membrane/fluid interface.

Competing Financial Interest

The authors declare no competing financial interests.

Acknowledgements

We are grateful to the CNRS for its financial support through the project "Waterloo", PICS. A.F. and A.S. thank Calcul Québec and Compute Canada to provide computational resources, through the financial support of the Canadian Foundation Innovation (CFI). Their work was supported by the Université de Sherbrooke, the Fonds Québécois de la Recherche sur la Nature et les Technologies (FRQNT), and the Natural Sciences and Engineering Research Council of Canada (NSERC).
References


• Water sub-diffusion at the nanoscale in the Nafion fuel cell membrane
• Time and water-uptake sub-diffusive to diffusive cross-over
• Molecular origin of the anomalous dynamics of water nano-confined into the Nafion
• Correlation between Nafion structure and the dynamics of water
• Self-assembly, interdigitated organization and water/Nafion interface
A.G, B.G., F.G. and A.F. performed all Molecular Dynamics simulations. All authors were involved in the interpretation of water transport. The manuscript was written by A.G and A.S.
Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.