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Temperature-Dependence of the Dielectric Relaxation of Water using Non-Polarizable Water Models

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Temperature-Dependence of the Dielectric

Relaxation of Water using Non-Polarizable Water

Models

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ABSTRACT

Key physical and chemical properties of aqueous fluids are determined by the structure and dynamics of the hydrogen bond network of water but we lack adequate models for the linkages between hydrogen bonding and aqueous chemistry, particularly in non-ambient conditions or in confinement. Dielectric relaxation spectroscopy (DRS) provides a sensitive approach for probing water dynamics but sound interpretation of DRS data requires molecular simulation and associated computational methods capable of accurately representing aqueous fluids and their frequency dependent, complex permittivity. Here, we test the accuracy of dielectric spectra of bulk liquid water calculated from molecular dynamics simulations using 19 non-polarizable water models at 298K. In contrast to prior studies, the simulation size, time-step and duration allow calculation of the dielectric function from 10^7 – 10^{12} Hz without assuming an analytical form. The accuracy of the prediction of the low-frequency (static) dielectric constant at room temperature is related to the water molecule dipole moment, specifically models with $\mu \geq 2.4D$ give $\epsilon(0)$ with a relative error

19 lower than 5%. However, no water model tested can fully reproduce the complex dielectric spectra
20 of water. For a subset of models, calculations of the dielectric response from -5–60C reproduces
21 the experimental trend in water dynamics with temperature but the characteristic relaxation time
22 is always under estimated. The calculated water dipole relaxation time and hydrogen-bond lifetime
23 are both exponentially decaying functions of temperature, and exhibit a linear correlation very
24 close to equality. The comparison provides new computational support for the concept that the
25 Debye relaxation of liquid water is determined by the dynamics of the hydrogen-bond network,
26 and that both are ensemble properties.

27 INTRODUCTION

28 Aqueous chemical processes are central to all hydrological and (geo)biological cycles that
29 maintain life on the planet¹⁻³ but descriptions of the molecular structure and chemical properties
30 of aqueous fluids, particularly in non-ambient conditions or in confinement, remain incomplete.
31 Key physical and chemical attributes of aqueous fluids are determined by the hydrogen bond
32 network of dipolar water molecules. For example, the solvation of ions requires hydrogen-bonded
33 water molecules to polarize and reorganize around a solute.^{4,5} However, no individual method is
34 capable of directly elucidating the structure and properties of hydrogen-bond networks. Valuable
35 insights into the hydrogen-bond network of aqueous fluids can be obtained from dielectric
36 relaxation spectroscopy (DRS), which measures the frequency-dependent, complex
37 permittivity $\varepsilon(\omega) = \varepsilon'(\omega) - i\varepsilon''(\omega)$, where $i = \sqrt{-1}$.⁶⁻⁸ The dielectric response of bulk liquid
38 water is dominated by the Debye relaxation, a large change in the permittivity that occurs in the
39 microwave range ($\nu \in 1-300$ GHz) with the inflection point in $\varepsilon'(\omega)$ at approximately 19 GHz at
40 room temperature. Hydrophilic and hydrophobic solutes, ion pair formation and other solution
41 interactions can shift relaxation frequency and introduce additional relaxations to a DRS

42 spectrum.⁶⁻⁸ In addition, water that is strongly bound to a solute or a surface typically exhibits a
43 dielectric response shifted to lower frequencies.⁹ In general, the spectrum is complex, and its
44 deconvolution into responses of individual components and distinct relaxation processes remains
45 a major challenge.^{6,9}

46 The most informative approach for interpreting DRS data would use molecular simulations that
47 accurately predict the hydrogen-bond dynamics of a system from which DRS spectra could be
48 calculated for comparison with experimental data. Methods for predicting DRS spectra from
49 atomic trajectories in molecular simulations have been developed over decades¹⁰⁻¹⁴ but remain far
50 from routine because the large frequency range of aqueous relaxation phenomena demands both a
51 small time step and a large simulation time. As a consequence, earlier studies were unable to
52 statistically sample the necessary timescales and fitted analytical response functions to the noisy
53 predicted relaxation behaviour. This process introduces ambiguity because debate remains over
54 the most appropriate approach for the decomposition of DRS data from water into one or more
55 contributions. Classical molecular dynamics (MD) simulations can increasingly access large
56 dynamical timescales enabling direct comparison with experiment.

57 In this study, we evaluate the ability of rigid, static-point-charge models, which are non-
58 polarizable, to reproduce the complex permittivity of bulk liquid water at 298K. For a subset of
59 the best performing models we also report calculations of $\epsilon(\omega)$ from -5–60C. This work
60 complements recent study by Cardona *et al.*^{14, 15} who compared different molecular models for
61 predicting the DRS of water and organic molecules as a prelude to the simulation of microwave
62 heating, and work by Segal and Schröder¹³ who compared polarizable and nonpolarizable water
63 molecules for predicting the dielectric and far-infrared spectra.

64

65

METHODS

66 Water Models

67 The interatomic potentials (force fields) that define the water models used in this study are based
68 on pair-wise non-bonded Lennard-Jones and electrostatic interactions, usually with the Lennard-
69 Jones parameters only assigned to the oxygen atom. The positive partial charges are assigned to
70 the hydrogen atoms, the negative charge is assigned to either the oxygen atom (3-point models), a
71 virtual site located on the HOH angle bisector (4-point models) or virtual sites completing the
72 tetrahedral oxygen-coordination (5-point models). The force fields have been constructed with
73 different levels of complexity and optimized to different sets of physicochemical properties¹⁶⁻¹⁸
74 that can include: structure (*e.g.*, radial distribution functions), equation of state, self-diffusion
75 coefficient, experimental gas-phase water dipole moment, liquid-phase dipole moment predicted
76 using *an initio* methods, and the static dielectric constant. However, it is currently intractable to
77 consider complex dynamical properties such as dielectric relaxation or heat capacities in force-
78 field development.

79 We chose the most popular water force-fields (*e.g.*, SPC¹⁹, SPC/E,²⁰ TIP3P,²¹ TIP4P,²¹ TIP5P²²)
80 and a few recently developed models that can reproduce the static dielectric constant of liquid
81 water at 298K (OPC,²³ OPC3,²³ TIP4Q,²⁴ TIP4P/ε,²² SPC-DC,²⁵ H2O-DC,²⁵ TIP3P-FB,²⁶ TIP4P-
82 FB²⁶). We also analysed the performance of the classical TIP n P water models that have been
83 revised for use with the particle-mesh Ewald (PME) method (*i.e.*, TIP3PF,²⁷ TIP4PEW,²⁸
84 TIP5PEW²⁹) and include one flexible water model (SPC/FW^{REF}). The parameters used in the water
85 models given in **Table 1**.

86 The simulations presented here do not consider the nuclear quantum effects that are known to
87 affect the structure and dynamics of the hydrogen-bonding structure in liquid water via tunneling,

88 proton delocalization, and intermolecular zero-point energy quantum fluctuations.³⁰⁻³⁵ The NQE
 89 effects are neglected for several reasons. First, the NQE produces the competing quantum effects
 90 on the HB network: weakening the weak HB bonds and strengthening the strong ones – overall
 91 canceling each other to a large extent in the bulk water at ambient conditions.³³⁻³⁵ Second, the ab
 92 initio path integral molecular dynamics is computationally too expensive to obtain nanosecond-
 93 range trajectories required to calculate the dielectric spectra. Finally, the NQE have relatively small
 94 effect on the average dipole moment of a water molecule in the bulk.^{30, 34}

95 **Table 1.** Force field parameters for non-polarizable water models used in our simulations.

Water model	Lennard-Jones parameters		partial charges			Geometry			Dipole moment ^c	
	σ_{O} (Å)	ϵ_{O} (kcal/mol)	q_{O} (e)	μ (D)	$q_{\text{O}/\text{X}}$ (e)	r_{OH} (Å)	$\angle\text{HOH}$ (°)	r_{OX} (Å)	μ (D)	
three-point models										
SPC	ref. ¹⁹	3.1657	0.1553	-0.8400	0.4200	1.000	109.47		2.274	
SPC/E	ref. ²⁰	3.1657	0.1553	-0.8476	0.4238	1.000	109.47		2.350	
SPC/EB	ref. ³⁶	3.1657	0.1553	-0.8476	0.4238	1.010	109.47		2.374	
SPC/FW ^a	ref. ³⁷	3.1657	0.1554	-0.8200	0.4100	1.012	107.57		2.395	
SPC-DC	ref. ²⁵	3.1577	0.1984	-0.8736	0.4368	1.000	109.47		2.423	
TIP3P	ref. ²¹	3.1506	0.1520	-0.8340	0.4170	0.957	104.52		2.348	
TIP3PF	ref. ²⁷	3.1941	0.0980	-0.8300	0.4150	0.957	104.52		2.336	
TIP3P-FB	ref. ²⁶	3.1780	0.1559	-0.8484	0.4242	0.957	104.52		2.419	
H ₂ O-DC	ref. ²⁵	3.1840	0.1417	-0.9099	0.4550	0.958	109.47		2.411	
OPC3	ref. ²³	3.1743	0.1634	-0.8952	0.4476	0.979	109.47		2.430	
four-point models										
OPC	ref. ²³	3.1666	0.2128	0.0000	0.6791	-1.3583	0.872	103.60	0.159	2.480
TIP4P	ref. ²¹	3.1537	0.1550	0.0000	0.5200	-1.0400	0.957	104.52	0.150	2.178
TIP4PEW	ref. ²⁸	3.1643	0.1628	0.0000	0.5200	-1.0484	0.957	104.52	0.125	2.322
TIP4P-FB	ref. ²⁶	3.1655	0.1791	0.0000	0.5259	-1.0517	1.012	108.15	0.105	2.429
TIP4P2005	ref. ³⁸	3.1589	0.1852	0.0000	0.5564	-1.1128	0.957	104.52	0.155	2.304
TIP4Q	ref. ²⁴	3.1666	0.1852	0.5000	0.5250	-1.5500	0.957	104.52	0.069	2.442
TIP4P/ε	ref. ²²	3.1650	0.1848	0.0000	0.5270	-1.0540	0.957	104.52	0.105	2.435
five-point models										
TIP5P ^b	ref. ²²	3.1199	0.1600	0.0000	0.2410	-0.2410	0.957	104.52	0.700	2.293
TIP5PEW ^b	ref. ²⁹	3.0970	0.1780	0.0000	0.2410	-0.2410	0.957	104.52	0.700	2.293

^aflexible water model (HOH angle flexibility modelled with a harmonic function, $k_{\angle\text{HOH}} = 37.95$ kcal/mol/rad²)³⁷

^btwo negatively charged virtual sites X with an angle $\angle\text{XOX} = 109.47^\circ$

^cexperimental value 2.4-2.95D,^{39, 40} theoretical estimation from the ab-initio molecular dynamics $\sim 3\text{D}$ ^{41, 42}

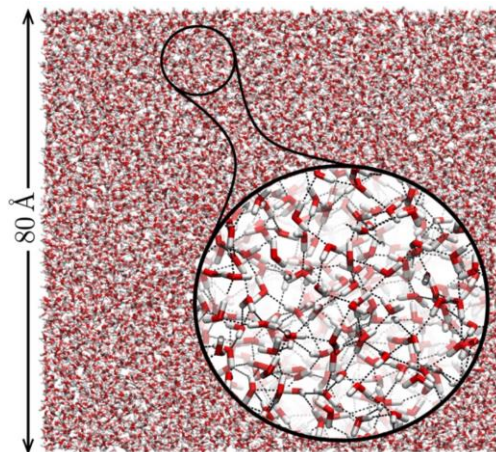
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101 Molecular Dynamic Simulation

102 Dielectric spectra and hydrogen-bond analysis were carried out by analysing classical MD
 103 trajectories obtained using *pmemd* simulation engine from the Amber18 package.⁴³ The initial

104 atom configurations were minimized using the mixture of the steepest descent and conjugate
105 gradient minimization schemes. Next, the system was brought to the desired temperature by
106 heating or cooling for 100 ps, (NVT ensemble) using Langevin thermostat (collision frequency
107 $\gamma=1.0 \text{ ps}^{-1}$). In the next step, we optimize the density and volume of the cell by running molecular
108 dynamics in the isobaric-isothermal ensemble (NPT) for additional 5 ns (Berendsen barostat with
109 the pressure relaxation time $\tau=2 \text{ ps}$, Langevin thermostat with $\gamma=1.0 \text{ ps}^{-1}$). The final simulations
110 were carried out using the optimized cell volume in the NVT ensemble for another 30 ns (Langevin
111 thermostat, $\gamma=1.0 \text{ ps}^{-1}$). The value of the collision frequency in the Langevin dynamics affects
112 the rate of convergence of static dielectric constant and controls stochastic variation between
113 simulations starting from the identical initial state as illustrated in Fig. S3. However, it does not
114 significantly alter the long-time averages, and therefore the simulations are carried out for $\gamma=1.0$
115 ps^{-1} . In all simulations we used the integrator time-step of 2 fs. The configurations were saved
116 every 0.2 ps (for calculating dielectric spectra) or 0.05 ps (for HB-analysis). The long-range
117 electrostatic interactions were calculated using the Particle Mesh Ewald summation method
118 (PME). The simulations presented here were obtained for a box of 17000 water molecules (**Fig. 1**)
119 and a short-range interaction cut-off of 16 Å – settings necessary to include the dipole-dipole
120 correlation beyond the second-solvation shell.⁴⁴ In addition, simulation of a larger number of
121 water molecules reduces the noise in the autocorrelation function and the calculated $\varepsilon(\nu)$ because
122 the fluctuations in the system properties are proportional to $1/\sqrt{N_{H_2O}}$. The final trajectory was
123 analysed using analysis codes developed in C++ and Python.



124

125 **Figure 1** Snapshot of the simulation cell with 17,000 H₂O water molecules. Inset illustrates the hydrogen-bond
 126 network in the liquid water.

127

128

129 Dielectric Properties

130 The water static dielectric constant, $\epsilon(0)$, is a collective property of an ensemble of water
 131 dipoles, which can be calculated from the equilibrium total dipole moment fluctuations, $\langle M^2 \rangle -$
 132 $\langle M \rangle^2$. The static dielectric constant is usually calculated using the following Clausius-Mosotti type
 133 equation⁴⁴

$$134 \quad \epsilon(0) = 1 + \frac{\langle M^2 \rangle - \langle M \rangle^2}{3\epsilon_0 V k_B T} \quad (1)$$

135 where ϵ_0 is the vacuum permittivity, V is the volume of an aqueous phase and M is the total
 136 dipole moment of the ensemble of molecular dipoles ($\vec{M} = \sum_i \vec{\mu}_i$).

137 The frequency-dependent dielectric constant is obtained from the Fourier-Laplace transform of
 138 the time-derivative of the normalized autocorrelation function of the total dipole moment, ϕ :⁴⁴

$$139 \quad \frac{\epsilon(\omega) - 1}{\epsilon(0) - 1} = \int_0^\infty \left(-\frac{d\phi(t)}{dt} \right) e^{-i\omega t} dt \quad (2)$$

140 where

$$141 \quad \phi(t) = \frac{\langle M(t)M(0) \rangle}{M^2} \quad (3)$$

142 By replacing the Fourier-Laplace transform by a half of the Fourier transform, and using eq. (1)
143 we obtain:

$$144 \quad \varepsilon(\omega) = 1 + \frac{\langle M^2 \rangle - \langle M \rangle^2}{6\varepsilon_0 V k_B T} \int_{-\infty}^{\infty} \left(-\frac{d\phi(t)}{dt} \right) e^{-i\omega t} dt \quad (4)$$

145 The local ordering and fluctuations of the dipole moments are usually quantified by the finite
146 (G_K) and infinite (g_K) system Kirkwood correlation factors, which are defined as:

$$147 \quad G_K = \frac{\langle M^2 \rangle - \langle M \rangle^2}{N \langle \mu^2 \rangle} \quad \text{and} \quad g_K = \frac{2\varepsilon(0) + 1}{3\varepsilon_0} G_K \quad (5)$$

148 The G_K factor measures the equilibrium fluctuations of the collective dipole moment of the
149 system and it is related to the orientational correlation function. The g_K factor measures local the
150 correlation between neighbouring dipole moments, for instance it can indicate the antiparallel
151 ($g_K < 1$), random ($g_K = 1$) or parallel neighbouring dipoles mutual orientation.

152 Analytical Debye Relaxation Models

153 The dominant water dipolar relaxation mode is accurately described by the Debye function:^{6-8,}
154 ⁴⁵

$$155 \quad \varepsilon(\omega) = \varepsilon_{\infty} + \frac{\varepsilon(0) - \varepsilon_{\infty}}{1 + i\omega\tau_D} \quad (6)$$

156 where $\varepsilon_{\infty} = \lim_{\omega \rightarrow \infty} \varepsilon'$ and τ_D is the Debye relaxation time, which is defined as the period of the
157 electromagnetic wave at the frequency of the maximum in ε'' ($\tau_D = \omega_D^{-1} = 1/2\pi\nu_D$). Quantitative
158 determination of τ_D is typically obtained by least-squares fitting but there is disagreement in the
159 literature on the number of relaxation contributions and the choice of a Debye function or a more
160 complex analytical form.^{6, 9} Here we used a model-free approach to comparing simulation and
161 experiment and read off τ_D directly from the peak position in the dielectric loss function. For visual

162 comparisons, the simulated dielectric spectra are plotted against experimental data from the
 163 literature,^{2,3} and their fit to a single Debye function.

164 **Hydrogen-Bond Lifetime**

165 Of the many definitions of the hydrogen-bond lifetime⁴⁶, here we followed Luzar and Chandler
 166 and use a reactive flux approach.^{47, 48} The interrupted hydrogen-bond lifetime is calculated from
 167 the time derivative of the hydrogen-bond correlation function, $c(t)$, given by an average over the
 168 simulation ensemble:⁴⁶⁻⁴⁸

$$169 \quad c(t) = \frac{\langle h(0)h(t) \rangle}{\langle h^2 \rangle} \quad (8)$$

170 where $h(t)$ is a binary classification function associated with each pair of water molecules: $h =$
 171 1 if a given pair is hydrogen bonded, and $h = 0$ otherwise. Thus, $c(t)$ measures the probability
 172 that two water molecules remain hydrogen-bonded at the time t if they were bonded at the time
 173 $t = 0$.⁴⁶ Two water molecules are considered to be hydrogen-bonded if oxygen-oxygen distance
 174 (r_{OO}) is not larger than 3.5 Å and hydrogen-donor-acceptor angle below 30°.⁴⁸

175 The reactive flux rate, $K(t)$, for hydrogen-bond breaking is defined as a difference between the
 176 rate of HB-breaking (k_b) and HB-(re)forming (k_f):⁴⁶⁻⁴⁸

$$177 \quad K(t) = -\frac{dc(t)}{dt} = k_b c(t) - k_f n(t) \quad (7)$$

178 where k_f, k_b are rate constants for HB (re)forming and breaking, respectively and $c(t)$ as defined
 179 above serves as an effective source term for the HB-bonded molecules. The HB (re)forming rate
 180 constant k_f is calculated from the detailed balance condition, that is:⁴⁶ $k_f = k_b \langle h \rangle / (1 - \langle h \rangle)$. $n(t)$
 181 gives the probability of breaking an existing hydrogen-bond while two water molecules remain
 182 within the HB-bond distance and represents a HB sink in eq. (7):^{46, 48}

$$183 \quad n(t) = \int_0^t -\frac{\langle \frac{dh(0)}{d\xi} \big|_0 (1-h(\xi))H(\xi) \rangle}{\langle h^2 \rangle} d\xi \quad (9)$$

184 where $H(\xi)$ is 1 if water molecules are within the HB-contact distance ($r_{00} \leq 3.5 \text{ \AA}$), or 0
185 otherwise.

186 The hydrogen-bond relaxation time is defined as an inverse of the HB-breaking rate constant
187 (i.e., $\tau_{HB} = k_b^{-1}$).⁴⁶ By knowing τ_{HB} , we can estimate the activation barrier for the breaking of the
188 hydrogen-bond. A procedure introduced by Luzar et al.⁴⁶⁻⁴⁸ assumes that the HB breaking is a
189 thermally activated process (i.e., obeys the Eyring-Polanyi equation):

$$190 \quad k_b = \frac{1}{\tau_{HB}} = \frac{k_B T}{h} \exp\left(-\frac{\Delta G_{HB}}{k_B T}\right) \quad (10)$$

191 where h is the Planck's constant, and ΔG_{HB} is the activation barrier ΔG_{HB} , which is an energetic
192 measure of the strength of the HB. It can be calculated from τ_{HB} as follows:⁴⁹

$$193 \quad \Delta G_{HB} = k_B T \ln\left(\frac{k_B T}{h} \tau_{HB}\right) \quad (11)$$

194 As noticed by Van Der Spoel et al.,⁴⁹ HB-lifetime values obtained using eq. (7) are sensitive to
195 the time-separation between subsequent configurations (Δt). Specifically, τ_{HB} decreases and k_f
196 increases rapidly with the decreasing Δt .⁴⁹ The results presented here are obtained consistently
197 for the same Δt values for each water model (i.e., $\Delta t = 50 \text{ fs}$).

198 RESULTS AND DISCUSSION

199 The calculated dielectric and hydrogen-bond network properties are given in **Table 2** and
200 compared with the experimental data² for water at 298 K. A comparison of the errors in predicted
201 static and dynamic dielectric properties is given in **Figure 2**.

202

203

204

205 **Table 2** Experimental and calculated dielectric properties of water at 298K for 19 nonpolarizable water models. The
 206 dielectric properties were obtained by analysing 30 ns molecular dynamics trajectories. The static dielectric constant
 207 ($\epsilon(0)$) is a slowly converging property (see Fig. S1, Supporting Information), the results presented here are within 2%
 208 relative error, which usually translates to $\pm 1-2 \epsilon(0)$ units.

Water model	Permittivity		Debye relaxation		HB-dynamics
	$\epsilon(0)$	$\epsilon(\infty)$	τ_D (ps)	ν_D (GHz)	τ_{HB} (ps)
SPC	66.61	7.91	4.72	33.73	6.00
SPC/E	70.80	7.49	5.66	28.10	6.89
SPC/EB	71.78	7.03	7.46	21.33	8.74
SPC/FW	83.17	7.95	6.68	23.83	7.417
SPC-DC	81.08	9.11	4.99	31.90	6.306
TIP3P	100.9	12.2	3.42	46.50	3.658
TIP3PF	91.23	9.06	4.87	32.67	4.959
TIP3P-FB	80.24	7.51	7.23	22.00	7.670
H ₂ O-DC	79.37	8.08	6.27	25.37	7.340
OPC3	80.08	7.98	5.99	26.57	7.539
OPC	79.50	8.76	5.70	27.90	7.354
TIP4P	51.56	8.34	3.33	47.83	5.353
TIP4PEW	64.61	7.65	5.43	29.33	7.160
TIP4P-FB	76.86	6.59	6.20	25.67	8.493
TIP4P2005	57.67	7.06	5.46	29.13	7.966
TIP4Q	80.87	7.91	6.77	23.50	7.630
TIP4P/ ϵ	79.18	7.71	6.76	23.53	7.986
TIP5P	92.44	7.71	7.16	22.23	6.490
TIP5PEW	98.36	8.04	6.64	23.97	6.173
Experiment ^{2,3}	78.36	5.2	8.27	19.24	

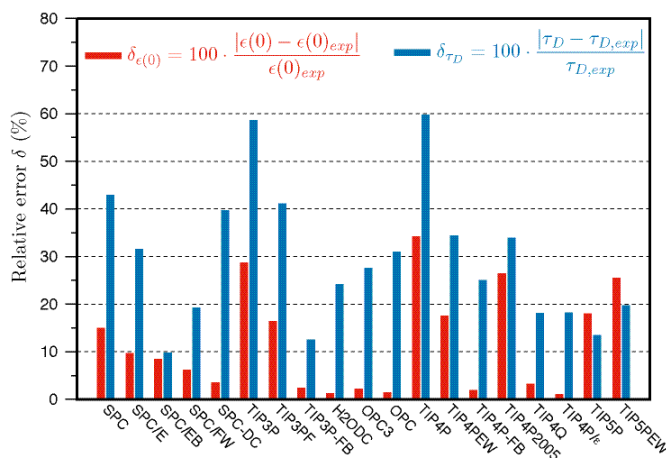
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211 Static Dielectric Constant

212 An assessment of the uncertainty in the predicted values of the static dielectric constant is
 213 required for assessing the results from the different models because $\epsilon(0)$ is a slowly converging
 214 property of molecular simulations. Fennell et al.²⁵ showed that a trajectory of at least 10 ns is
 215 needed to converge on $\epsilon(0)$. However, $\epsilon(0)$ -estimation is also sensitive to molecular dynamics
 216 details: molecular dynamics integrator, force-calculation scheme, and the temperature control. For
 217 one water model (TIP4P-FB) we examined the convergence of static dielectric properties
 218 ($\epsilon(0)$, g_K , G_K) for simulations that started from the same initial configuration and diverged in time
 219 due to the randomness introduced by the Langevin thermostat (g_K , G_K in Table S1). The simulation
 220 results obtained from 30-ns trajectories show at least 2-3% variability (**Figs. S1-S2**), which could
 221 explain why reported values of $\epsilon(0)$ for various water models vary between different simulation

222 studies (e.g.,^{24, 27, 50}). Among all 19 considered water models we can identify 9 that predict the
 223 static dielectric constant of water at room temperature that is within 5% of the experimental value.
 224 The remaining 10 water models are unable to correctly predict the static dielectric constant, usually
 225 because these models have been developed to reproduce other water properties. Cardona *et al.*¹⁴
 226 proposed scaling $\epsilon(\omega)$ to the experimentally measured value of $\epsilon(0)$, but that approach is not
 227 adopted here.

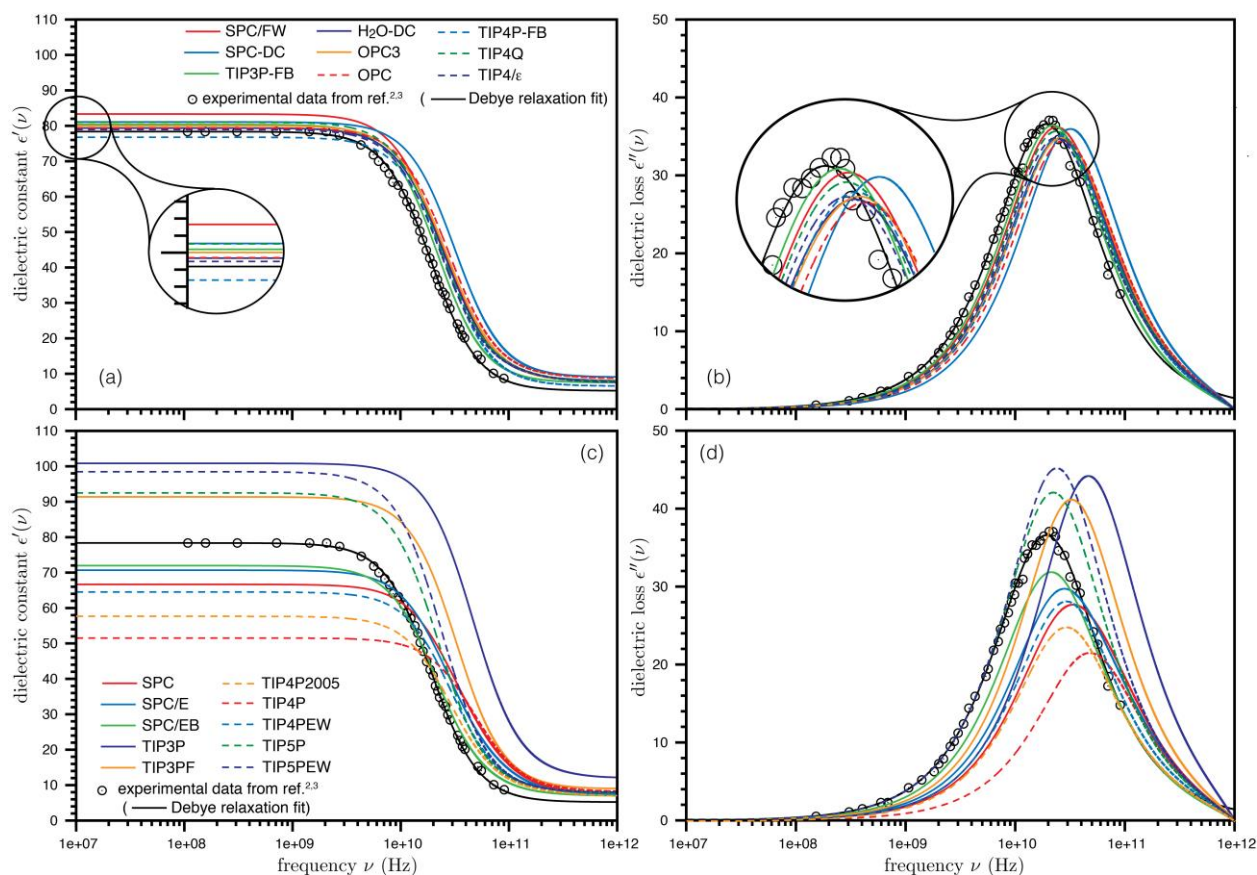


228
 229 **Figure 2.** Relative error in predicted static dielectric constant, $\epsilon(0)$, and Debye relaxation time, τ_D , at room
 230 temperature among considered water models (30 ns simulations). The experimental data, $\epsilon(0)_{exp}$, $\tau_{D,exp}$, are taken
 231 from ref.³

232 Dielectric Relaxation Spectra

233 The predicted DRS for the 9 water models that provided the best predictions of the static
 234 dielectric constant are shown in **Figure 3a**; the dielectric spectra for the remaining models are
 235 shown in **Figure 3b**. The comparison shows that accuracy in the prediction in the static dielectric
 236 constant is highly correlated with accuracy in representing the frequency-dependent dielectric
 237 relaxation. However, none of the water models exactly reproduces the dielectric spectra, as
 238 previously observed¹³. For all models, the predicted Debye relaxation time is shorter than the
 239 experimental value ($\tau_D = 8.27$ ps; $\nu_D = 19.24$ GHz). Only the 5-point models are able to reproduce

240 the low-frequency side of the dielectric loss peak at 298 K (see **Fig. 3a,b**), but they still
 241 overestimate the values of $\epsilon(0)$. Although OCP and OCP3 were intended to improve ions and
 242 proteins solvation, these models are not optimum for the frequency-dependent dielectric
 243 relaxation. TIP3P-FB seems to be the best for both static and dynamic dielectric properties, a likely
 244 consequence of the well designed and efficient optimization strategy developed in the force-
 245 balance method.²⁶



246 **Figure 3.** a) Predicted dielectric relaxation spectra of bulk liquid water using 9 water models that gave the smallest
 247 relative error in predicted static dielectric constant at 298 K. b) Simulated dielectric spectra using 10 water models
 248 with the largest relative error in predicted static dielectric constant at 298 K.
 249

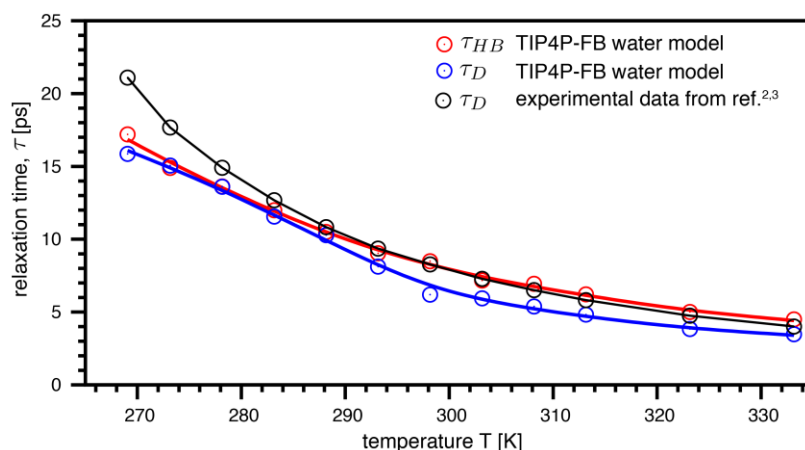
250 Water Dipole Moment

251 We find that the relative error in $\epsilon(0)$ at 298 K obtained for all water models with the dipole
 252 moment $\mu > 2.4$ is below 5%. Accordingly, we asked whether a modification to the force-field

253 parameters to obtain $\mu > 2.4$ would improve the predictions of the dielectric properties. For the
254 TIP5P water model, changing the partial charge from 0.241 to 0.2555 ($q_H = -q_x$) achieves $\mu =$
255 2.43 D. The predicted static dielectric constant at 298K is 77.2, a significant improvement over
256 the original prediction of 92.44. However, prediction of the Debye relaxation is severely worsened
257 with $\tau_D=281$ ps, almost two orders of magnitude too slow. This suggests that force-field
258 parametrization of rigid water models must consider the Debye relaxation in order to more
259 accurately capture the dynamical properties of aqueous solutions. Even models that reproduce the
260 self-diffusion coefficient, which captures an aspect of water HB dynamics, do not provide accurate
261 DRS prediction.

262 **Temperature Dependence**

263 We tested the ability of 8 of the 9 best performing models (omitting the flexible model) to
264 reproduce the frequency dependent dielectric spectra of liquid water at temperatures between -5–
265 60°C (**Figs. S4-S7**) with the results for TIP4P-FB illustrated in **Figure 4**. For these models and
266 this temperature range, the dipolar relaxation consistently occurs at higher frequency (shorter
267 relaxation time) than the experimental data but the discrepancies between the experimental data
268 and predictions for both $\varepsilon(0)$ and $\varepsilon(\nu)$ are approximately constant. Thus, although the models
269 over predict the rotational mobility of water they provide a reasonable prediction of the activation
270 barrier for water reorganization.

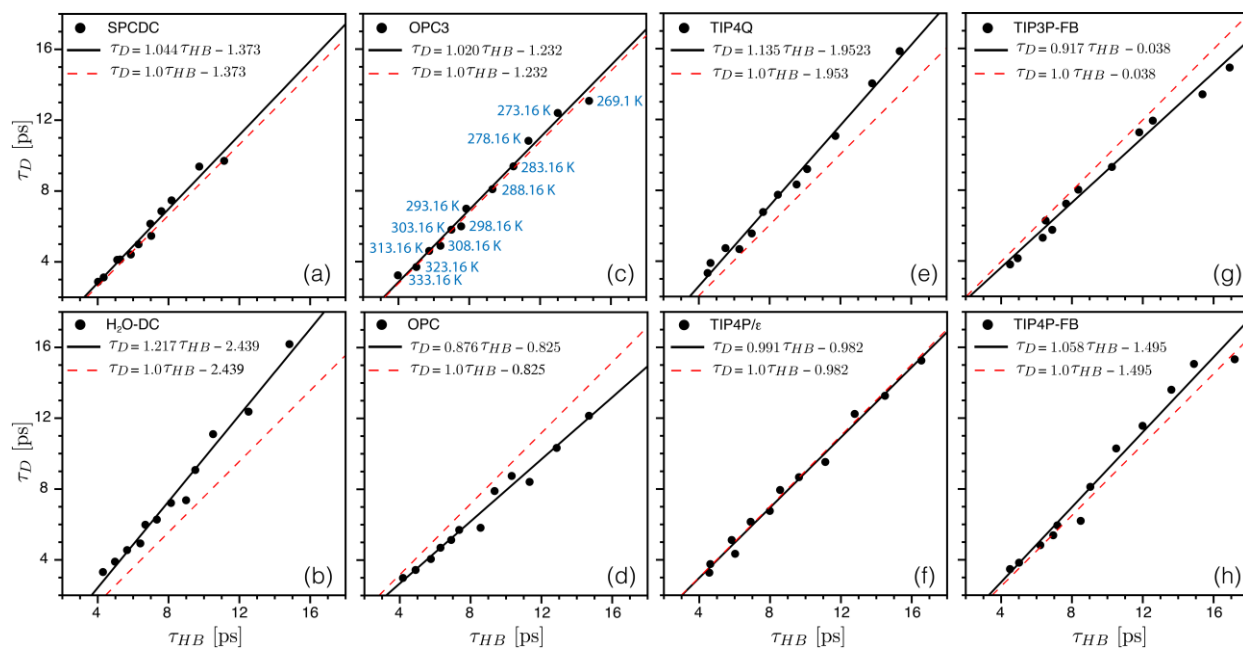


271
 272 **Figure 4** Temperature dependence of the predicted Debye relaxation time, τ_D , and hydrogen-bond lifetime, τ_{HB} , for
 273 the water TIP4P-FB model, compared with the experimental value of τ_D from refs.^{2,3}

274 The temperature series also highlight that the MD simulations do not accurately reproduce the
 275 dissipation of energy caused by water dipole reorientation at frequencies close to the Debye
 276 relaxation. The energy loss to the liquid by dielectric relaxation (called dielectric heating) is
 277 quantitatively related to the integrated area of $\epsilon''(\nu)$ (**Figs. S4-S7**). Only the TIP3P-FB closely
 278 approximates the dielectric loss function at 269 K (**Fig. S7b**).

279 Hydrogen Bond Lifetime

280 By analysing the Debye-relaxation times and hydrogen-bond lifetimes for the temperature series
 281 we find that both quantities have a similar in magnitude and have a similar temperature-
 282 dependence (illustrated for TIP4P-FB in **Fig. 4**). Plots of τ_D vs τ_{HB} show a robust linear correlation
 283 between these properties (**Fig. 5**). The constant of proportionality ranges from 0.92–1.21 and the
 284 constant offset ranges between 0.04–2.4 ps. For the best-performing model (TIP4P-FB), the
 285 relationship is the closest to equality.



286

287 **Figure 5.** Correlation plots of the Debye relaxation time (τ_D) and the hydrogen-bond lifetime (τ_{HB}) for several water
 288 models for the temperature range -5 – 60°C . Black points are predictions from individual simulations and black lines
 289 are a linear regression of the form $\tau_D(T) = a\tau_{HB}(T) + b$, where T is temperature and a and b are fitted constants
 290 displayed for each plot. Red dashed lines are regressions with $a = 1$. The temperature value corresponding to each
 291 point is shown in Panel (c) and the points follow the same order in other panels.

292

CONCLUSION

293 This work demonstrates that classical molecular dynamics simulation of the dielectric relaxation
 294 of water remains challenging and is very sensitive to the force-field parameter values, in agreement
 295 with recent studies¹³⁻¹⁵ and as previously demonstrated for the static dielectric constant.^{24, 27, 50}
 296 Even the water models that provide the most accurate predictions of $\epsilon(0)$ fail to reproduce absolute
 297 values for $\epsilon(\omega)$ including the principal relaxation frequency and the width and amplitude of the
 298 dielectric loss peak. Because the dielectric loss peak describes how water dipole reorganization is
 299 coupled to energy dissipation and heating, absolute errors in the simulation of aqueous fluids will
 300 affect predictions of important properties (heat capacity), physical phenomena (microwave
 301 heating) and any chemical phenomena involving entropy changes (*e.g.*, solvation and charge-

302 transfer processes involving solvent reorganization). Despite these limitations, the best water
303 models predicted dielectric behaviour with consistent accuracy over an important temperature
304 range for geologic, biologic and technologic systems.

305 Analysis of the temperature-dependent data revealed a close relationship between the mean
306 hydrogen-bond lifetime, calculated for simulation ensembles using the reactive flux method, and
307 the Debye relaxation frequency, also an ensemble property. This finding provides new support for
308 the concept that dipolar relaxation in liquid water is governed by the collective dynamics of the
309 hydrogen-bond network. Although current models of Debye relaxation are based on this intuitive
310 assumption,⁵¹⁻⁵⁶ it has been challenging to provide full support either from experiment or
311 simulation. Analysis of individual simulation trajectories led to the development of the *wait and*
312 *switch* model⁵¹⁻⁵⁶ in which defects in HB-network are necessary for water molecule to orient in the
313 electric field, but the detailed mechanism, such as the role for hydrogen bond defects, remains
314 debated. Time-resolved two-dimensional infrared (2D IR) spectroscopy experiments determined
315 that hydrogen bond rearrangement requires a collective neighbor response that is likely larger than
316 accessible through trajectory analysis.⁵⁷ The present work demonstrates that hydrogen-bond
317 breaking and Debye relaxation are system behaviors with a tight temporal correlation.

318

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411