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# The interfacial layer breaker: a violation of Stokes' law in high-speed Atomic Force Microscope flows

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## Abstract

Structured water near surfaces is important in non-classical crystallization, biomineralization, and restructuring of cellular membranes. In addition to equilibrium structures, studied by atomic force microscopy (AFM), high-speed AFM (H-S AFM) can now detect piconewton forces in microseconds. With increasing speeds and decreasing tip diameters, there is a danger that continuum water models will not hold, and molecular dynamics (MD) simulations would be needed for accurate predictions. MD simulations, however, can only evolve over tens of nanoseconds due to memory and computational efficiency/speed limitations, so new methods are needed to bridge the gap. Here we report a hybrid, multi-scale simulation method, which can bridge the size and timescale gaps to existing experiments. Structured water is studied between a moving silica AFM colloidal tip and a cleaved mica surface. The computational domain includes 1472766 atoms. To mimic the effect of long-range hydrodynamic forces occurring in water, when moving the AFM tip at speeds from  $5 \times 10^{-7}$  to 30 m/s, a

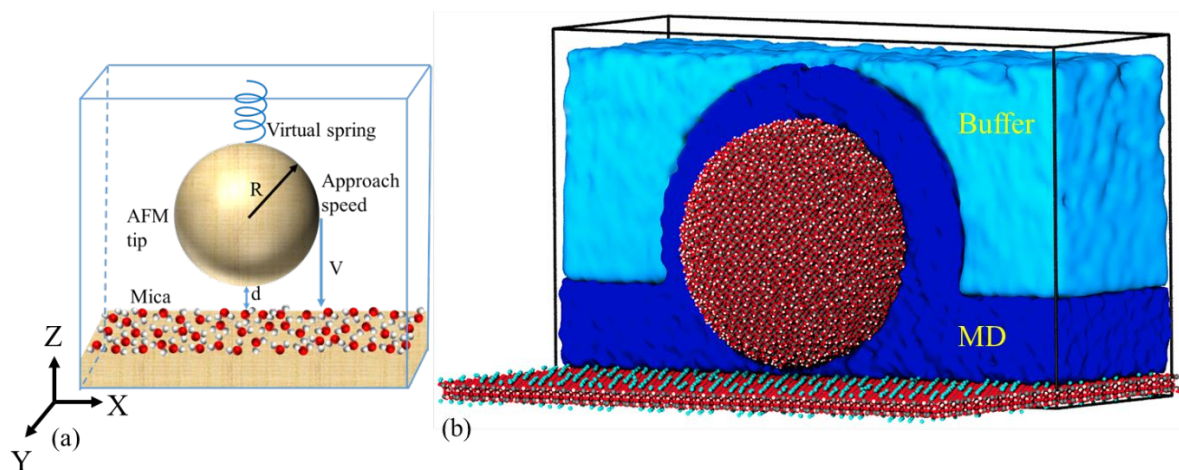
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hybrid multiscale method with local atomistic resolution is used, which serves as an effective open-domain boundary condition. The multiscale simulation is thus equivalent to using a macroscopically large computational domain with equilibrium boundary conditions. Quantification of the drag force shows breaking of continuum behaviour. Non-monotonic dependence on both tip speed and distance from the surface imply breaking of the hydration layer around the moving tip at timescales smaller than water cluster formation and strong water compressibility effects at the highest speeds.

## **Introduction**

The intricacy of water structure at nanoscale distances from a material surface has been observed by both experiments and equilibrium molecular dynamic (MD) simulations [1-6]. The study of this structure is important in non-classical crystallization [7-9], biomineralization [10], and tribochemical wear [11]. Interfacial water properties are also pertinent for the restructuring processes of cellular membranes [12-14], organismal signalling [15]. The advent of advanced High-Speed Atomic Force Microscopy (H-S AFM) opens new opportunities for measuring the dynamic piconewton forces, which occur within microseconds and nanometre structural rearrangements with subsecond resolution [16, 17]. AFM tools are used to detect the equilibrium structure of water layers, but the dynamic behaviour is still beyond the experimental reach in the sub-microsecond and nanosecond region typical of structural rearrangements and bio-catalytic molecular systems [18-20]. While the timestep of molecular dynamics (MD) simulations is orders of magnitude too small and so is their system sizes to model such processes, which involve collective dynamics of many water molecules, the hybrid multi-scale methods, which combine MD with continuum fluid dynamics simulations [21-24] provide an attractive opportunity for modelling of realistic-size systems while resolving details of the molecular ordering. Here we report the detection of water structuring between a silica AFM colloidal tip and a cleaved mica surface. The quantification of the drag force on the tip as a function of tip-wall distance and tip velocity shows non-monotonic force, which becomes negative for high-speed regimes, implying variations not only viscosity but also density. At the highest speeds

these are consistent with breaking of the hydration layer around the moving tip and formation of a layered water structure in the sub-nanometer gap between the tip and the substrate surface.



**Figure 1.A)** Schematic representation of one-half of the symmetric computational domain (26 nm x 19 nm x 15.6 nm). The symmetry plane corresponds to the top boundary in the z direction. The silica tip is represented by a sphere ( $R = 5 \text{ nm}$ ),  $d$  is a distance between the tangential plane of sphere and the potassium ions (K) layer of the mica surface. The sphere is suspended by a massless spring from the top of the box, and the resulting force  $F$  on tip is measured in accordance with Hooke's law,  $F = -k \cdot \Delta z$ , where  $k$  and  $\Delta z$  are the virtual spring constant and the displacement. The approach speed  $V$  of the tip is varied. **B)** A cut through the simulation domain showing the decomposition of the hybrid multiscale simulation domain into several regions. The silica tip and mica surface are modelled at the full-atom resolution. In addition, a part of the water volume in a rectangular slab, which covers the mica surface with a 3.7 nm thickness and the water in a spherical shell, which covers the silica tip with a 2.79 nm thickness are also modelled at all-atom resolution (water in the pure atomistic MD region is shown in dark blue). Outside of the pure atomistic region, a multi-scale interface region is placed where the MD particles are subjected to a

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**hydrodynamic force including the Brownian motion. The thickness of the multi-scale region is selected to include at least a few cut-off radii of van der Waals forces for best performance (see Supporting Information for further details). The multi-scale interface and the outer region, where water molecules are fully dominated by hydrodynamics forces (Buffer) are shown in light blue. The hydrodynamic forces are consistent with the macroscopic Stokes solution in a large water volume surrounding the MD domain so that no reflection of pressure waves occurs at the boundaries of the multiscale domain.**

## **Method**

Figure 1 (a) shows the schematic representation of the simulated AFM device, where a silica tip functionalized by hydroxy group on surface with a radius of 5 nm that corresponds to a mid-range tip size compared to the values reported in the AFM experiments [25, 26]. The tip is attached by a massless virtual spring and suspended above mica substrate at a distance  $d$ . The simplification of AFM device as a massless spring attached to a spherical tip is a well-established practice in molecular modelling for it simplifies comparison of the numerical simulation with the analytical Stokes' solution in the continuum hydrodynamics theory [16, 27]. In both cases, the drag force acting on a sphere moving in viscous liquid is considered. Details of the force comparison between the results of the multi-scale calculations and predictions based on the Stokes' law are presented in the following paragraph. Following [28, 29], a detailed atomistic model of the mica substrate ( $\text{K}[\text{Si}_3\text{AlO}_8]\text{Al}_2\text{O}_2(\text{OH})_2$ ) has been implemented. In the model, the Si $\rightarrow$ Al charge defects are distributed in agreement with the solid state Nuclear Magnetic Resonance data [30]. The silica tip and mica substrate are fully immersed in water. For water-mica, water-tip and silica-mica modelling, the recently developed INTERFACE force field is applied which was shown to provide good results for interfacial systems [31]. Figure 1 (b) illustrates the partition of simulation domain in the multiscale model, where the external boundary conditions in the continuum part of the simulation are prescribed in accordance with the macroscopic Stokes model by introducing the Fluctuating Hydrodynamics (FH) force in the equations of MD particles.

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The centre of the domain including the water layers on material surfaces is simulated at the all-atom resolution, hence all the interactions at all-atom level are accurately captured as in the conventional MD method. The continuum flow effects are enforced by the moving tip via the effective boundary conditions using the multiscale method (see the method details in Supporting Information). Therefore, the input parameters of the model are the tip-substrate distance,  $d$  and the AFM speed,  $V$  while the output is the force on the AFM tip (Fig. 1).

It should be noted that the bulk hydrodynamic flow effect induced by the moving AFM boundary decays slowly with the distance. Furthermore, the simulation time needed for averaging out the thermal velocity fluctuations of water atoms ( $\sim 100$  m/s), which amplitude is much larger than the hydrodynamic signal, is macroscopically large too. Altogether this means that using the standard all-atom MD approach with equilibrium boundary conditions would require a very large computational domain in space and time, which is prohibitively expensive. Therefore, the implemented multiscale method essentially truncates the computational domain, while preserving the atomistic resolution where it is critical to account for short-range interatomic interactions.

To calculate the total force exerted on the AFM tip, individual forces in the all-atom region from the surrounding water atoms and the mica substrate are summed up and time averaged. The averaging volume comprising the water and mica atoms around the AFM tip, which defines the force exerted on the AFM tip, is governed by short-range interatomic interactions between those atoms and the microscopic AFM tip. This microscopic volume is the effective volume, which corresponds to the statistical ensemble averaging used in the current simulations. The collective force associated with the pure hydrodynamic effect is computed as a difference between the total AFM force with and without the AFM tip motion.

To validate the FH/MD model, the tip velocity is set to zero, and the resulting force-distance curve solution is compared with the solution obtained from the all-atom molecular dynamics simulation performed for the same system with periodic boundary

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conditions. The force is calculated in accordance with Hooke's law,  $F = -k \cdot \Delta z$ , where  $k$  and  $\Delta z$  are the virtual spring constant and the displacement in the  $z$ -direction. The distance  $d$  in the force-distance curve is defined as the distance between the tangential plane of sphere and the potassium ions (K) layer of the mica surface. Notably, as shown in Fig. S1 (a) in Supporting Information, the difference between the FH/MD and the reference all-atom MD solution is within 0.05 nN. Furthermore, the sensitivity of the MD solution to the averaging window, 2 ns versus 3 ns is also approximately within the same 0.05 nN error bar. (See Fig. S1 (b)). It can be noted that this difference could be further reduced by running the simulations for a longer time to better converge the models statistically.

The above error of the multiscale model solution is much smaller than the uncertainty of the computed force associated with the thermal fluctuations of water molecules in the volume corresponding to the tip and the typical averaging time of MD solutions. On the one hand, the uncertainty in the AFM force measurement due to the Brownian motion can be estimated as the difference between the forces applied on the AFM sphere and its mirror image in the upper part of the symmetrical computational domain (which fluctuates randomly for different AFM distances and speeds in accordance with random nature of the Brownian force, as shown in Fig. S2 in Supporting Information). On the other hand, for the case when the AFM tip is approaching the mica surface slowly and is considered at a distance greater than the hydration layer from the mica surface, the uncertainty in the force associated with the Brownian motion can be estimated as a perturbation of Stokes' law,  $\delta F \sim 6\pi R\mu \cdot \text{Var}(V')$ . Here  $\mu$  is the dynamic viscosity of bulk water and  $V'$  is the thermal velocity fluctuation. The variance corresponds to averaging over the Grand-Canonical ensemble in accordance with the Fluctuating Hydrodynamics theory [21],  $\text{Var}(V') = \sqrt{\frac{k_B T}{\rho_0 V_0}}$ , where  $V_0 = 4\pi R^3/3$  is the characteristic control volume,  $k_B$  is the Boltzmann constant,  $T$  is the temperature, and  $\rho_0$  is the bulk water density at equilibrium conditions. For the large AFM distance of 0.77 nm and the low speed of  $5 \times 10^{-7} \text{ m/s}$ ,

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where Stokes' law should be approximately valid, the theoretical estimate gives  $\delta F = 0.234 \text{ nN}$ . It can also be noted that the latter uncertainty associated with the Brownian motion in the small molecular system considered is in the order of the positive drag force predicted by the continuum Stokes' theory at medium-high AFM speeds ( $F^{Stokes} = 0.283 \text{ nN}$  at 0.3 m/s). This explains why the classical positive hydrodynamic drag effect is not resolved in the AFM force-distance curve (Fig. S2).

Furthermore, the estimated uncertainty due the Brownian motion is in good agreement with the numerical simulation result, i.e., one half of the magnitude of the force difference between the AFM sphere and its mirror image in the same conditions,  $0.3 \text{ nN}$  (Fig. S2 in Supporting Information). Hence, in the remaining part of the article, the numerically computed one half of the absolute difference between the computed forces on the AFM tip and its mirror image will be used as an estimate of the standard deviation of the statistical error for each tip speed and mica-tip distance.

Results of additional calculations with the FH/MD method for a test molecular liquid system are included in Supporting Information (Fig.S5). The test system shares a few pertinent features with the water flow over the AFM tip, such as flows through the hybrid FH/MD interface region, whilst allowing a direct comparison with a reference all-atom MD solution in this case. The comparison shows that the moving multiscale interface does not lead to noticeable numerical artefacts in computing the molecular diffusion coefficient.

## Results

Properties of water close to the mica surface simulated with the FH/MD solution are analysed separately in Fig. 2. To reduce the effect of the AFM tip, the stationary regime is considered where the tip is fixed at a distance from the mica of  $d = 0.77 \text{ nm}$ , which is much larger than the radius of van der Waals forces. From Fig. 2 (a), it can be noted that the water atoms are absorbed next to the mica cations, one per vacancy of the mica crystal, in accordance with the experimental and previous numerical studies [32, 33]. Notably, the periodic structure formed by absorption of

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water atoms in the mica structure is crucial for the hydration layer formation. For example, as demonstrated in the atomic force microscope experiments using frequency modulation [2] and equilibrium all-atom MD simulations [4, 34], the hydration layer is responsible for the amplitude of the repulsive peak of the AFM force-distance curve at  $d = 0.5 - 0.6$  nm when the tip velocity is zero (Fig. 4 (b) and Fig.S1 (a)). Further discussion of the force-distance curve can be found in the Supporting Information.

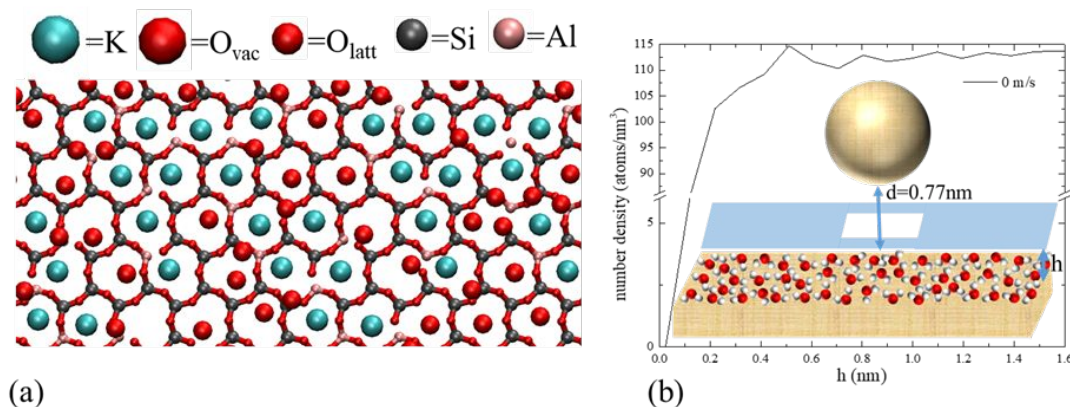
Fig. 2 (b) shows the distribution of the number density of water in the (wall-normal) z-direction from the mica surface. The density profile is calculated by averaging of the mass of water molecules contained in each vertical bin between the mica surface and the maximum height (0.05 nm) of the water column considered in the mica-tip gap over 2 ns. It should be reminded that, as in Fig.2a, the point of interest here is the water density profile close to the mica and away from the AFM tip. Therefore, to exclude the effects of non-uniform water confinement between the mica surface and the AFM tip, the water volume corresponding to the square area of 10 nm x 10 nm under the AFM tip was excluded from the averaging when computing the water density profile on mica. Notably, the resulting density profile has a prominent peak in the vicinity of the mica surface followed by several smaller peaks and deeps. Again, this is in good qualitative agreement with typical interfacial water structure observed experimentally.

Having validated the force-distance curve of the multiscale model for the stationary AFM regime, and shown the known water layering near a mica surface, we investigated the fate of water layering in a dynamic context. We performed a series of simulations for a range of approach speeds of the AFM tips from  $5 \times 10^{-7} m/s$  (typical speed of the AFM cantilever in experiments) to  $30 m/s$  (hypothetical high-speed AFM regime).

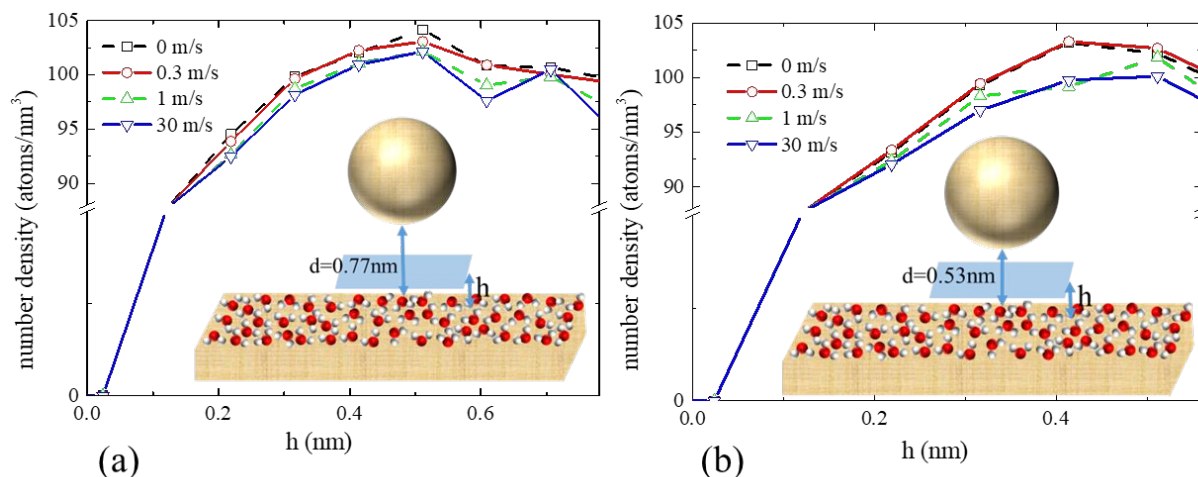
We find that though layering under the tip close to the mica is disturbed due to the presence of the moving curved geometry. For the relatively large mica-tip distance,  $d=0.77$  nm, the higher the tip speed, the more layering could be observed, which is especially notable at 30 m/s, which corresponds to an oscillation in the density profile at around 0.6 nm in Fig. 3(a). Fig. 3 (b) shows the variation of water density with height



for a different AFM tip-mica separation distance  $d=0.53$  nm and a range of approach speeds, 0-30 m/s. For such small mica-tip distance, regardless of the tip speed, the hydration layer cannot fit in the gap due to the proximity of the AFM tip.



**Figure 2.A) The first mica layer in contact with the water. The interface is composed of potassium ions from the mica (blue) and oxygen atoms adsorbed in vacant potassium cavities in the mica (bigger red  $O_{vac}$ ). Oxygen (smaller red  $O_{latt}$ ), silicon (grey) and aluminium (pink) make the rest of the mica lattice. B) Plot of average number density of water atoms in planar 0.05 nm thick slices immediately above the mica (and excluding the 10x10 nm square area below the sphere, (white)) as a function of the slice' height,  $h$  in the  $z$ -coordinate direction. The potassium ions' plane of mica surface is taken as  $h = 0$ , and densities are plotted up to a height of 1.6 nm. The mica-tip distance from the same reference is,  $d = 0.77$  nm.**



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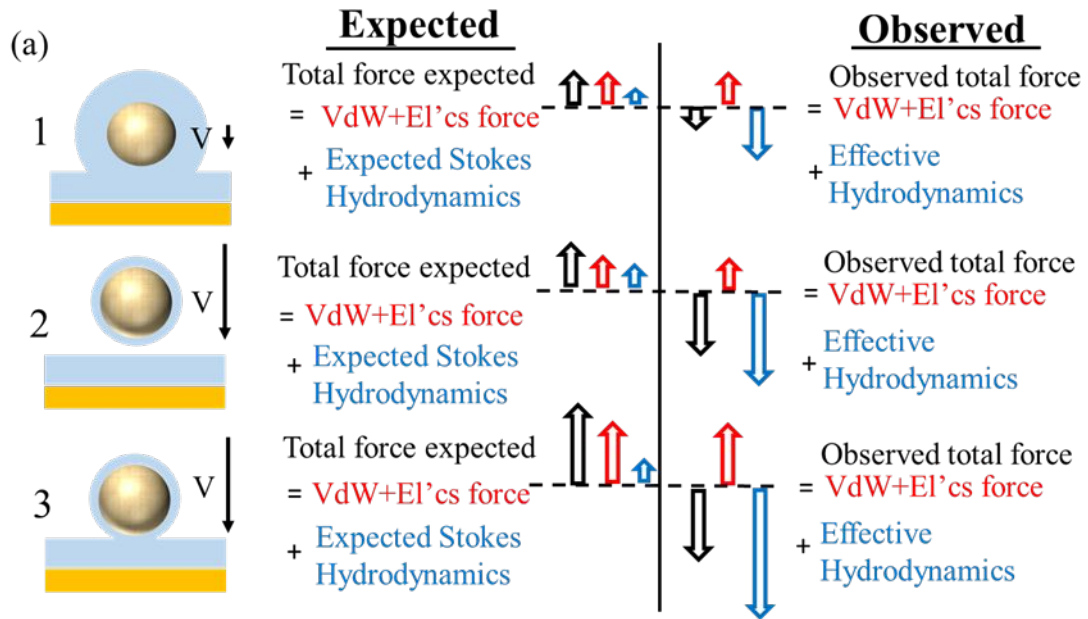
**Figure 3. Averaged number density of water atoms in a planar slice below the tip sphere (blue, 10 nm x 10 nm) as a function of the slice' height,  $h$  in the  $z$ -coordinate direction; the blue area in this figure corresponds to the white area in Figure 2(b). Bin thickness used in the averaging is 0.05 nm. (a) and (b) correspond to 0.77 nm and 0.53 nm mica-tip distance,  $d$ , respectively. Note the distinctly layered structure of the water density distribution at 30 m/s tip speed in (a).**

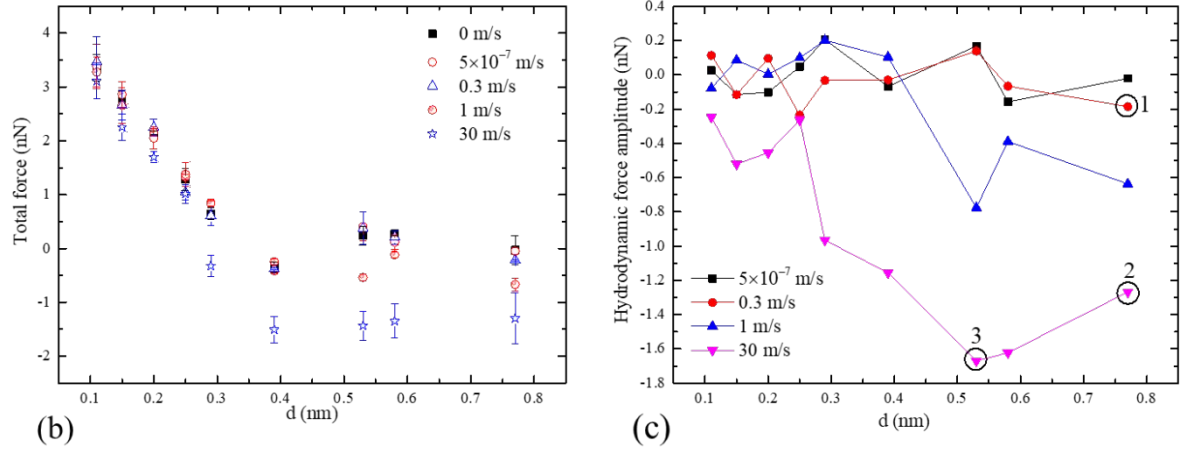
In turn, the dynamic layering of water near the tip leads to significant, non-intuitive departures from the expected forces on the AFM tip in accordance with the continuum hydrodynamics.

Simulation results for various tip/mica distances,  $h$  and AFM speeds,  $V$  are summarised in Fig.4 (b) and (c). Fig.4 (a) provides a roadmap to these results in the form of sketches. On the left, three typical tip/mica configuration regimes are presented: (1) the equilibrium situation when the tip is moving at a low speed with hydration layers formed on both the surfaces and (2) and (3) when the tip moving at high speed further or closer to the mica, so that the hydration layer on the tip does not have time to form. The second and the third columns show schematics of the force balances, where the total force includes both the conventional interatomic, van der Waals and electrostatic forces, which mostly depend on the tip-mica distance, and the collective effect of moving water molecules (the hydrodynamic force). The second column corresponds to the classical Stokesian hydrodynamics, which ignore the non-continuum effects such as the hydration layer, thereby always predicts a positive total repulsion force on the tip. In accordance with this model, the hydrodynamic drag of a small sphere moving in a viscous liquid only depends on the liquid viscosity, sphere diameter and speed, but critically is in the direction opposite the body velocity [35]. The third column shows the actual balance of the forces in accordance with the simulation FH/MD results. In accordance with the multiscale simulations, the hydrodynamic force is not only of a different magnitude but also in the opposite direction compared to the Stokes theory. The potential role of slip was excluded as a cause, as according to the continuum theory,

the drag force reduces by 1.5 times when the hydrophilic nonslip condition on the spherical tip surface is replaced by the hydrophobic full-slip [36]. And regardless of the wall boundary condition on a moving sphere down, the Stokes' drag should always correspond to a repulsive force. However, the actual simulation results show an attractive hydrodynamic force. It can be hypothesised that the collective attractive force may be associated with destroying the hydration layers around the fast-moving microscopic body, which redistributes pressure around the tip with creating an upstream rarefaction zone in the water volume by analogy with the ice breaker.

The total forces experienced on the nanometer tip when moving at different speeds up to 30 m/s are shown with their error bars in Figure 4 (b). This dataset is used to compute the hydrodynamic interactions by subtracting the well-defined interatomic (van der Waals and electrostatic) interactions from the total force. We plot the resulting hydrodynamic forces in Fig. 4 (c), showing they are non-monotonic, large in magnitude, especially in the highest velocity case, 30 m/s, and opposite to their usual direction in accordance with the Stokes model. Typical flow regimes (1), (2), and (3) discussed in Fig. 4(a) are clearly marked on the force distribution curves.





**Figure 4. A) Schematic of the sphere approaching the structured water layer above the mica plane; the thickness of the interfacial water layers depends on the mica-tip gap and the tip speed and comparison of the total tip force predicted by classical hydrodynamic theory and FH/MD. For  $d = 0.77 \text{ nm}$ , at small speeds (denoted by Num. 1), a hydration layer occurs on the tip surface. At high speed (denoted by Num. 2), the hydration layer on the AFM tip breaks down because of water molecules depleting from it to bulk thereby resulting in occurrence of layered water structure, which corresponds to the single hydration layer on the mica surface, while the hydration layer on the mica stays the same. As the tip approaches the mica at high speed even closer (denoted by Num. 3) both the hydration layer on the tip and the mica are broken down because of the confinement. Centre and right: comparison of the total force based on the predicted hydrodynamic force from Stokes' law and the numerical simulation result.**

**B) Total force on the AFM tip as a function of distance from the mica surface for velocity 0,  $5 \times 10^{-7}$ , 1, and 30 m/s. The error bar is the absolute difference of total force between the AFM tip and its symmetrical image on the top. The total force includes van der Waals (VdW) and electrostatic (El'cs) force from water and mica substrate for the zero tip velocity. For other velocities, the total force also includes hydrodynamic force because of the incorporated flow. C) The effective Hydrodynamic drag force on the sphere, i.e. the difference between the computed**

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**AFM forces in the stationary regime and the force in the flow regime measured for the same mica-tip distance.**

The answer to the puzzling behaviour of the hydrodynamic force lies in the violation of the continuum theory due to breaking of the hydration layer on the moving tip surface. When the tip-mica distance  $d$  is in the range of a small integer number of hydration layers, the forces can no longer be described by the continuum theory. Further, the formation of such hydration layer structures is associated with the collective behaviour of water clusters, which require a small but finite time to form, in the order of  $10^{-11}$  s according to [37]. At the same time, the tip approaching the mica surface at a speed of 30 m/s passes through the characteristic length scale of water structure, such as the equilibrium intermolecular distance between the oxygen atoms, 0.31 nm in about  $10^{-11}$  s. This shows that, for the very high approach speeds, there is little time for the hydration layers to form on the AFM tip. This observation is consistent with the decreased number density around the tip, as shown in Fig. S3 in the Supporting Information. Furthermore, there is another important physical time scale for this problem, which is related to water compressibility. As the water squeezed between the mica surface and the approaching tip is pushed and displaced to the tip sides, the propagating pressure waves need to cover the distance equal at least to the tip diameter (10 nm). Assuming the pressure waves propagate in water at normal speed of sound conditions (1500 m/s) the time for a single wave to traverse the tip is also similar  $0.67 \times 10^{-11}$  s. This means the water compressed under the approaching AFM tip at a speed of 30 m/s does not have time to settle to the equilibrium pressure state, which would require many passing-through cycles of the pressure wave. Altogether, this suggests that, depending on the tip speed and size, the water in the sub-nano-meter gap between the approaching tip and the mica can be highly compressible, thereby potentially leading to the formation of strong non-divergent flows and rarefaction under the tip. In turn, this generates a negative pressure zone below the spherical tip, which cannot be explained by the classical Stokes model that assumes water incompressibility. The compressibility of water is another potential cause for the oscillation in the water

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density at the 30 m/s tip speed, as shown earlier in Fig. 3 (a). In accordance with the numerical solution, the characteristic size of this oscillation, which occurs close to the AFM tip surface, is about 0.2 nm. By combining this space length with the tip speed (30m/s), one obtains the corresponding time scale of the oscillation,  $0.67 \cdot 10^{-11}$  s, which is remarkably close to the water compressibility time scale discussed above.

## Conclusions

In conclusion, we demonstrate a hybrid continuum - molecular dynamics (FH/MD) method capable of simulating 1472766 atoms, in a  $7700 \text{ nm}^3$  box size system for 2 to 3 ns with MD-like precision, while the outer continuum part of the model, which drives the continuum bulk flow effects is many orders of magnitude larger. Hence, the key feature of the multiscale solution used here is that it allows bridging the gap in time scales between the fast atoms/ thermal fluctuations and the slow continuum/ AFM motion.

Notably, the cantilever structure above the AFM tip in the multiscale simulation was approximated by a massless spring model consistently with the molecular dynamics literature [16, 27]. This is a reasonable approximation because it is equivalent to focusing on the localized flow effects around the microscopic tip, which exhibit the non-Stokesian behaviour due to the breach of the continuity assumption at high-speed AFM regimes. These effects are treated as a perturbation to the continuum bulk flow imposed away from the moving AFM tip. At the same time, the size of the full AFM system does not influence the simulation time significantly, as any increases beyond the immediate vicinity of the mica and the tip are treated only by continuum mechanics. This is because: (1) majority of the atomistic forces are short ranged (a few nanometers) and (2) the important length scales of the problem such as the AFM tip radius and the distance between the AFM tip and the substrate wall are 5 nanometer and a sub-nanometer, respectively. Therefore, in static conditions, the entire AFM system could be modelled at all atom resolution within a modest MD box at equilibrium boundary conditions. What cannot be modelled by pure MD methods is the slow motion of the AFM tip, which generates substantial bulk-flow hydrodynamic effect on the boundary

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conditions of the water atoms surrounding the AFM tip. Because this hydrodynamic effect is much slower than thermal fluctuations, one would require a prohibitively long time averaging/ MD simulation times to separate this collective effect from thermal noise. At the same time, the simulation of the bulk-flow effects using a continuum fluid dynamics method is many orders of magnitude faster than the equivalent all-atom MD model, and which speed-up is utilised in the suggested multiscale model. For a 5 nm sphere, 0.5 - 0.77 nm from a mica surface, moving towards it at 1 - 30 m/s, we show that the continuum Stokes' law expression for drag force fails spectacularly. Rather than a drag force opposite the direction of motion, the effective hydrodynamic force is in the direction of the motion, and can be significantly larger than the respective electrostatic and van der Waals contributions. The collective force of water molecules deviates from the continuum, where the time to rearrange molecules during the high-speed AFM tip motion falls below the time of formation of water clusters and the time for the water density around the tip to equilibrate. With decrease in tip sizes and higher frequencies current high-speed AFM (HS-AFM) experiments are starting to approach velocities considered here. FH/MD simulations would be critical for interpreting the force components once the operation velocity in future advanced H-S AFM experiments reaches 1 m/s.

The current simulation results can be viewed as a high-speed extension of recent experimental observations from [38], which revealed that the atomic-scale features observed in force-distance curves in AFM experiments reflect the inhomogeneous perturbation introduced by the tip and the substrate in the liquid.

### **Supporting Information**

FH/MD method and its validation for the force-distance AFM curve; Validation of the FH/MD method for a test molecular system with moving the atomistic/continuum interface.

### **Acknowledgements**

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The work of F.L. was supported by the China Scholarship Council (CSC). I.A.K. and S.A.K. gratefully acknowledge the funding under the European Commission Marie Skłodowska-Curie Individual Fellowship Grant No. H2020-MSCA-IF-2015-700276 (HIPPOGRIFFE). S.S. gratefully acknowledge EPSRC fellowship EP/R028915/1. The work was also supported by European Commission in the framework of the RISE program, Grant No. H2020-MSCA-RISE-2018-824022-ATM2BT.

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## References

- (1) Umeda, K.; Zivanovic, L.; Kobayashi, K.; Ritala, J.; Kominami, H.; Spijker, P.; Foster, A. S.; Yamada, H. Atomic-resolution three-dimensional hydration structures on a heterogeneously charged surface. *Nat. Commun* **2017**, *8* (1), 2111. DOI: 10.1038/s41467-017-01896-4.
- (2) Fukuma, T.; Ueda, Y.; Yoshioka, S.; Asakawa, H. Atomic-Scale Distribution of Water Molecules at the Mica-Water Interface Visualized by Three-Dimensional Scanning Force Microscopy. *Phys. Rev. Lett.* **2010**, *104* (1), 016101. DOI: 10.1103/PhysRevLett.104.016101.
- (3) Argyris, D.; Ashby, P. D.; Striolo, A. Structure and Orientation of Interfacial Water Determine Atomic Force Microscopy Results: Insights from Molecular Dynamics Simulations. *ACS Nano* **2011**, *5* (3), 2215-2223. DOI: 10.1021/nn103454m.
- (4) Tsukada, M.; Watanabe, N.; Harada, M.; Tagami, K. Theoretical simulation of noncontact atomic force microscopy in liquids. *J. Vac. Sci. Technol. B* **2010**, *28* (3), C4C1-C4C4. DOI: 10.1116/1.3430541.
- (5) Watkins, M.; Reischl, B. A simple approximation for forces exerted on an AFM tip in liquid. *J. Chem. Phys.* **2013**, *138* (15), 154703. DOI: 10.1063/1.4800770.
- (6) Fukuma, T.; Reischl, B.; Kobayashi, N.; Spijker, P.; Canova, F. F.; Miyazawa, K.; Foster, A. S. Mechanism of atomic force microscopy imaging of three-dimensional hydration structures at a solid-liquid interface. *Phys. Rev. B* **2015**, *92* (15), 155412. DOI: 10.1103/PhysRevB.92.155412.
- (7) Gerrard, N.; Gattinoni, C.; McBride, F.; Michaelides, A.; Hodgson, A. Strain Relief during Ice Growth on a Hexagonal Template. *J. Am. Chem. Soc.* **2019**, *141* (21), 8599-8607. DOI: 10.1021/jacs.9b03311.
- (8) Michaelides, A.; Morgenstern, K. Ice nanoclusters at hydrophobic metal surfaces. *Nat. Mater.* **2007**, *6* (8), 597-601. DOI: 10.1038/nmat1940.
- (9) Wu, S.; He, Z.; Zang, J.; Jin, S.; Wang, Z.; Wang, J.; Yao, Y.; Wang, J. Heterogeneous ice nucleation correlates with bulk-like interfacial water. *Sci. Adv.* **2019**, *5* (4), eaat9825. DOI: 10.1126/sciadv.aat9825.
- (10) Lu, H.; Huang, Y.-C.; Hunger, J.; Gebauer, D.; Cölfen, H.; Bonn, M. Role of Water in CaCO<sub>3</sub> Biomineralization. *J. Am. Chem. Soc.* **2021**, *143* (4), 1758-1762. DOI: 10.1021/jacs.0c11976.



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- (11) Wang, X.; Kim, S. H.; Chen, C.; Chen, L.; He, H.; Qian, L. Humidity Dependence of Tribochemical Wear of Monocrystalline Silicon. *ACS Appl. Mater. Interfaces* **2015**, *7* (27), 14785-14792. DOI: 10.1021/acsami.5b03043.
- (12) Mondal, J. A.; Nihonyanagi, S.; Yamaguchi, S.; Tahara, T. Structure and Orientation of Water at Charged Lipid Monolayer/Water Interfaces Probed by Heterodyne-Detected Vibrational Sum Frequency Generation Spectroscopy. *J. Am. Chem. Soc.* **2010**, *132* (31), 10656-10657. DOI: 10.1021/ja104327t.
- (13) Moilanen, D. E.; Spry, D. B.; Fayer, M. D. Water Dynamics and Proton Transfer in Nafion Fuel Cell Membranes. *Langmuir* **2008**, *24* (8), 3690-3698. DOI: 10.1021/la703358a.
- (14) Zhao, Q.; Majsztrik, P.; Benziger, J. Diffusion and Interfacial Transport of Water in Nafion. *J. Phys. Chem. B* **2011**, *115* (12), 2717-2727. DOI: 10.1021/jp1112125.
- (15) Khesbak, H.; Savchuk, O.; Tsushima, S.; Fahmy, K. The Role of Water H-Bond Imbalances in B-DNA Substate Transitions and Peptide Recognition Revealed by Time-Resolved FTIR Spectroscopy. *J. Am. Chem. Soc.* **2011**, *133* (15), 5834-5842. DOI: 10.1021/ja108863v.
- (16) Rico, F.; Russek, A.; González, L.; Grubmüller, H.; Scheuring, S. Heterogeneous and rate-dependent streptavidin–biotin unbinding revealed by high-speed force spectroscopy and atomistic simulations. *Proc. Natl. Acad. Sci.* **2019**, *116* (14), 6594-6601. DOI: 10.1073/pnas.1816909116.
- (17) Valotteau, C.; Sumbul, F.; Rico, F. High-speed force spectroscopy: microsecond force measurements using ultrashort cantilevers. *Biophys. Rev.* **2019**, *11* (5), 689-699. DOI: 10.1007/s12551-019-00585-4.
- (18) Banerjee, D.; Pal, S. K. Direct observation of essential DNA dynamics: melting and reformation of the DNA minor groove. *J. Phys. Chem. B* **2007**, *111* (36), 10833-10838.
- (19) Andreatta, D.; Pérez Lustres, J. L.; Kovalenko, S. A.; Ernsting, N. P.; Murphy, C. J.; Coleman, R. S.; Berg, M. A. Power-law solvation dynamics in DNA over six decades in time. *J. Am. Chem. Soc.* **2005**, *127* (20), 7270–7271.
- (20) Brauns, E. B.; Madaras, M. L.; Coleman, R. S.; Murphy, C. J.; Berg, M. A. Complex local dynamics in DNA on the picosecond and nanosecond time scales. *Phys. Rev. Lett.* **2002**, *88* (15), 158101.
- (21) Korotkin, I.; Karabasov, S.; Nerukh, D.; Markesteijn, A.; Scukins, A.; Farafonov, V.; Pavlov, E. A hybrid molecular dynamics/fluctuating hydrodynamics method for modelling liquids at multiple scales in space and time. *J. Chem. Phys.* **2015**, *143* (1), 014110.
- (22) Markesteijn, A.; Karabasov, S.; Scukins, A.; Nerukh, D.; Glotov, V.; Goloviznin, V. Concurrent multiscale modelling of atomistic and hydrodynamic processes in liquids. *Philos. Trans. Royal Soc.* **2014**, *372* (2021). DOI: 10.1098/rsta.2013.0379.
- (23) Korotkin, I. A.; Karabasov, S. A. A generalised Landau-Lifshitz fluctuating hydrodynamics model for concurrent simulations of liquids at atomistic and continuum resolution. *J. Chem. Phys.* **2018**, *149* (24), 244101. DOI: 10.1063/1.5058804.

- 
- (24) Li, F.; Korotkin, I.; Farafonov, V.; Karabasov, S. A. Lateral migration of peptides in transversely sheared flows in water: An atomistic-scale-resolving simulation. *J. Mol. Liq.* **2021**, *337*, 116111. DOI: <https://doi.org/10.1016/j.molliq.2021.116111>.
- (25) Heath, G. R.; Scheuring, S. High-speed AFM height spectroscopy reveals  $\mu$ s-dynamics of unlabeled biomolecules. *Nat. Commun* **2018**, *9* (1), 4983. DOI: 10.1038/s41467-018-07512-3.
- (26) Uchihashi, T.; Kodera, N.; Ando, T. Guide to video recording of structure dynamics and dynamic processes of proteins by high-speed atomic force microscopy. *Nat. Protoc.* **2012**, *7* (6), 1193-1206. DOI: 10.1038/nprot.2012.047.
- (27) Hu, X.; Nanney, W.; Umeda, K.; Ye, T.; Martini, A. Combined Experimental and Simulation Study of Amplitude Modulation Atomic Force Microscopy Measurements of Self-Assembled Monolayers in Water. *Langmuir* **2018**, *34* (33), 9627-9633. DOI: 10.1021/acs.langmuir.8b01609.
- (28) Heinz, H.; Koerner, H.; Anderson, K. L.; Vaia, R. A.; Farmer, B. L. Force Field for Mica-Type Silicates and Dynamics of Octadecylammonium Chains Grafted to Montmorillonite. *Chem. Mater.* **2005**, *17* (23), 5658-5669. DOI: 10.1021/cm0509328.
- (29) Heinz, H.; Vaia, R. A.; Farmer, B. L. Interaction energy and surface reconstruction between sheets of layered silicates. *J. Chem. Phys.* **2006**, *124* (22), 224713. DOI: 10.1063/1.2202330.
- (30) Heinz, H.; Suter, U. W. Surface Structure of Organoclays. *Angew. Chem. Int. Ed.* **2004**, *43* (17), 2239-2243. DOI: 10.1002/anie.200352747.
- (31) Heinz, H.; Lin, T.-J.; Kishore Mishra, R.; Emami, F. S. Thermodynamically Consistent Force Fields for the Assembly of Inorganic, Organic, and Biological Nanostructures: The INTERFACE Force Field. *Langmuir* **2013**, *29* (6), 1754-1765. DOI: 10.1021/la3038846.
- (32) Martin-Jimenez, D.; Garcia, R. Identification of Single Adsorbed Cations on Mica-Liquid Interfaces by 3D Force Microscopy. *J. Phys. Chem. Lett.* **2017**, *8* (23), 5707-5711. DOI: 10.1021/acs.jpcclett.7b02671.
- (33) Park, S.-H.; Sposito, G. Structure of Water Adsorbed on a Mica Surface. *Phys. Rev. Lett.* **2002**, *89* (8), 085501. DOI: 10.1103/PhysRevLett.89.085501.
- (34) Kobayashi, K.; Liang, Y.; Amano, K.-i.; Murata, S.; Matsuoka, T.; Takahashi, S.; Nishi, N.; Sakka, T. Molecular Dynamics Simulation of Atomic Force Microscopy at the Water-Muscovite Interface: Hydration Layer Structure and Force Analysis. *Langmuir* **2016**, *32* (15), 3608-3616. DOI: 10.1021/acs.langmuir.5b04277.
- (35) Brenner, H. The slow motion of a sphere through a viscous fluid towards a plane surface. *Chem. Eng. Sci.* **1961**, *16* (3), 242-251. DOI: [https://doi.org/10.1016/0009-2509\(61\)80035-3](https://doi.org/10.1016/0009-2509(61)80035-3).
- (36) Luo, H.; Pozrikidis, C. Effect of surface slip on Stokes flow past a spherical particle in infinite fluid and near a plane wall. *J. Eng. Math.* **2008**, *62* (1), 1-21. DOI: 10.1007/s10665-007-9170-6.
- (37) O. L. Lange, L. K., E.-D. Schulze. *Water and Plant Life Problems and Modern Approaches*; Springer, 1976.

(38) Hernández-Muñoz, J.; Uhlig, M. R.; Benaglia, S.; Chacón, E.; Tarazona, P.; Garcia, R. Subnanometer Interfacial Forces in Three-Dimensional Atomic Force Microscopy: Water and Octane near a Mica Surface. *J. Phys. Chem. C* 2020, 124 (48), 26296-26303. DOI: 10.1021/acs.jpcc.0c08092.

### TOC Graphic

