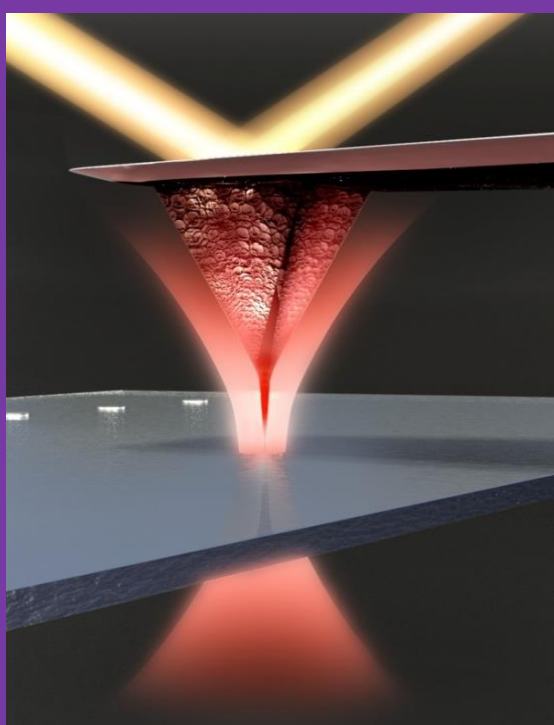




Glass: a multi-platform specimen supporting substrate for precision single molecule studies of membrane proteins



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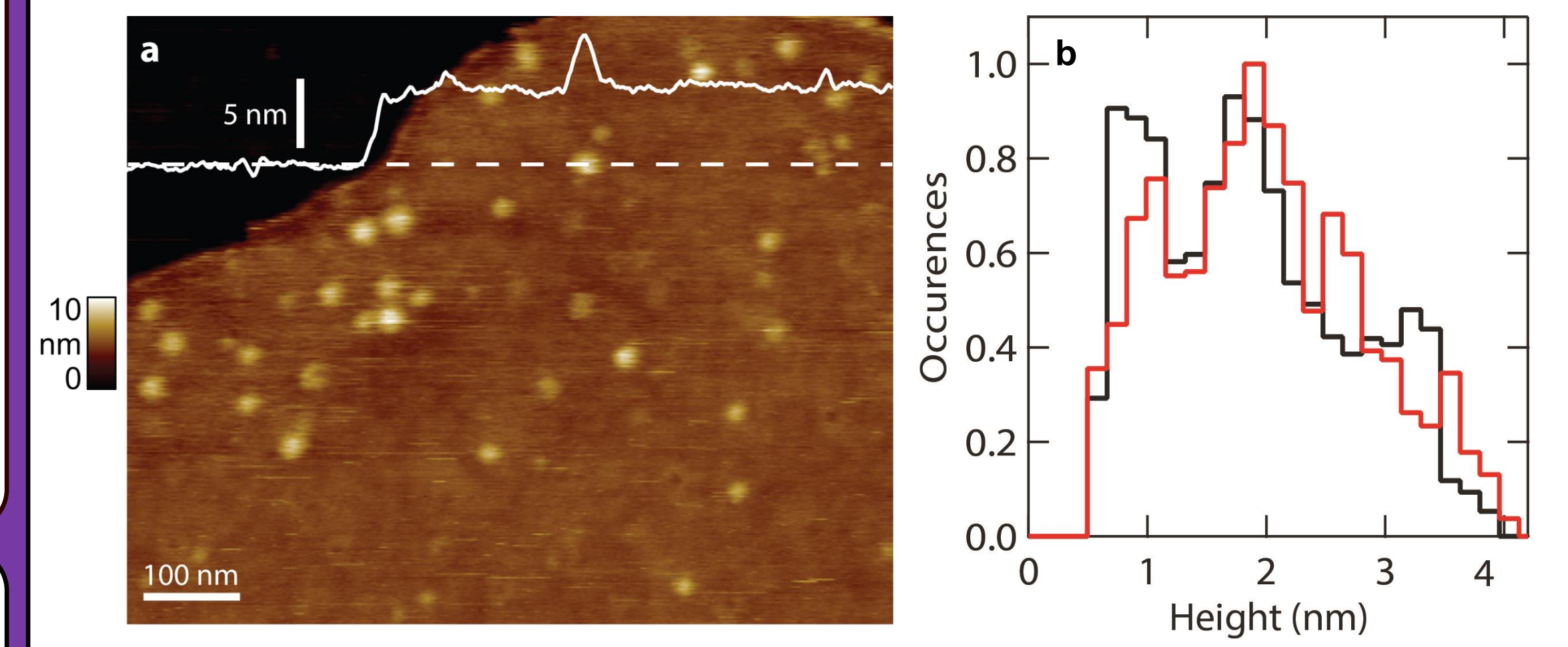
Abstract

High resolution (~1 nm lateral resolution) biological AFM imaging has been carried out almost exclusively using freshly cleaved mica as a specimen supporting surface, but mica suffers from a fundamental limitation that has hindered AFM's broader integration with many modern optical methods. Mica exhibits biaxial birefringence; indeed, this naturally occurring material is used commercially for constructing optical wave plates. In general, propagation through birefringent material alters the polarization state and bifurcates the propagation direction of light in a manner which varies with thickness. This makes it challenging to incorporate freshly cleaved mica substrates with modern optical methods, many of which employ highly focused and polarized laser beams passing through the specimen plane. Using bacteriorhodopsin from *Halobacterium salinarum* and the translocase SecYEG from *Escherichia coli*, we demonstrate that faithful images of 2D crystalline and non-crystalline membrane proteins in lipid bilayers can be obtained on common microscope cover glass following a straight-forward cleaning procedure. Direct comparison between data obtained on glass and on mica show no significant differences in AFM image fidelity. This work opens the door for combining high resolution biological AFM with powerful optical methods that require optically isotropic substrates such as ultra-stable¹ and direct 3D AFM². In turn, this capability should enable long timescale conformational dynamics measurements of membrane proteins in near-native conditions.

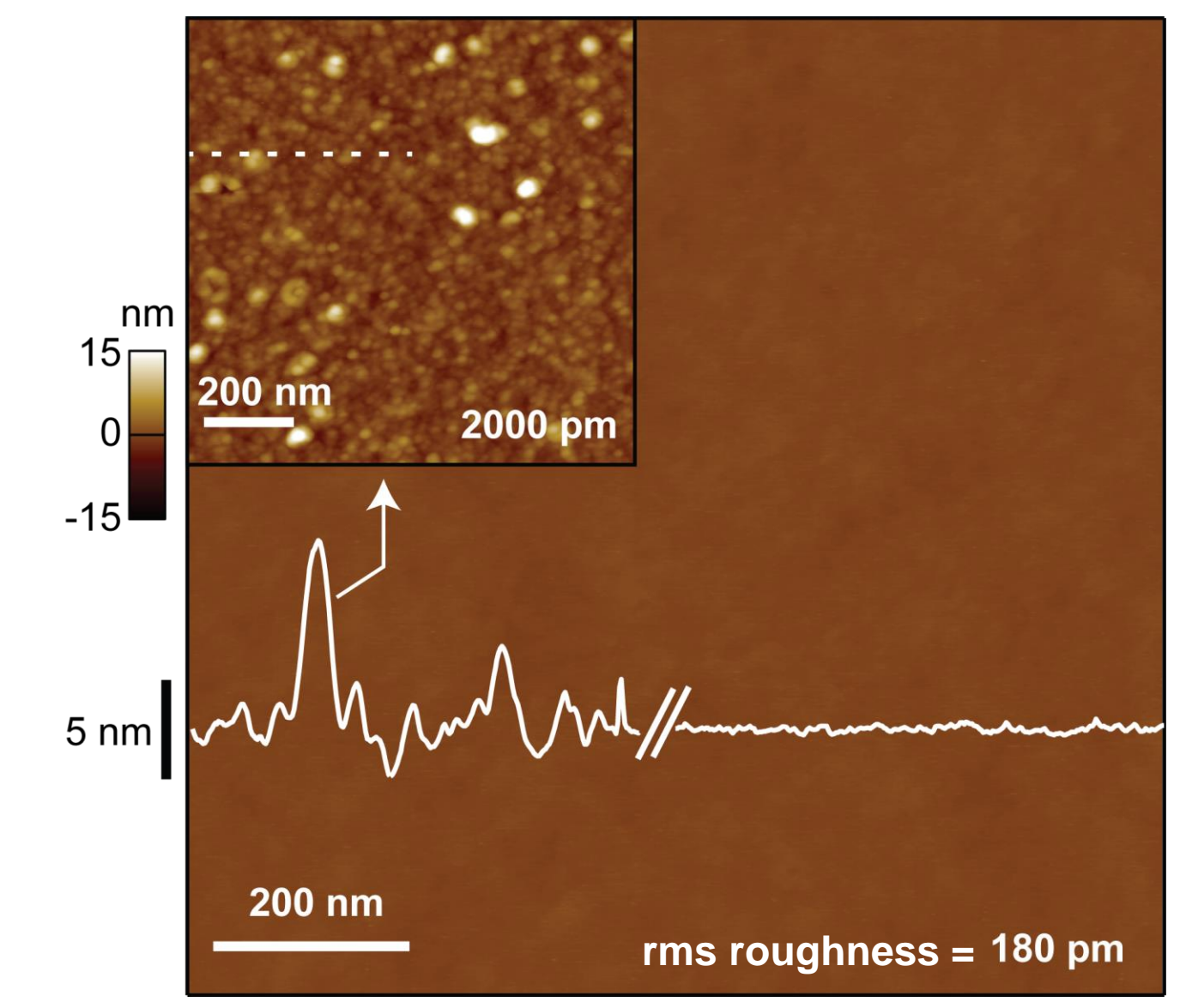
Why use glass as a substrate?

- Typical AFM substrates mica and HOPG exhibit birefringence, complicating optical paths³
- Many well established biological and biochemical assays like FRAP, FRET, Fluorescence Microscopy, FIONA and TIRF require transparent and optically homogeneous substrates³
- Recently developed techniques like US-AFM¹, 3D AFM² and polarization anisotropy methods require transparent substrates and non-birefringence due to polarization based detection
- Amorphous and optically homogeneous substrates like glass enable coupling AFM with advanced imaging techniques³

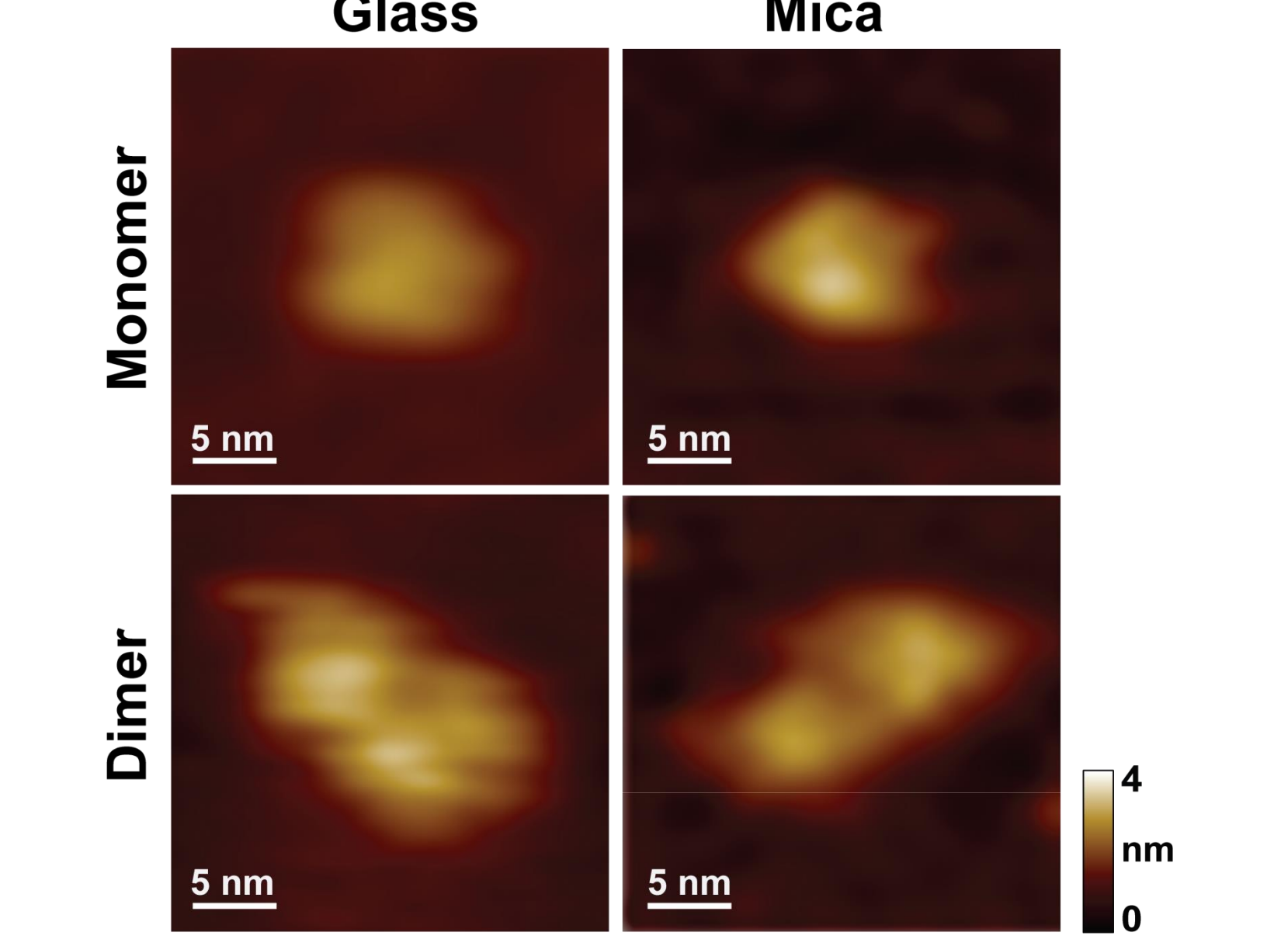
Heights of SecYEG



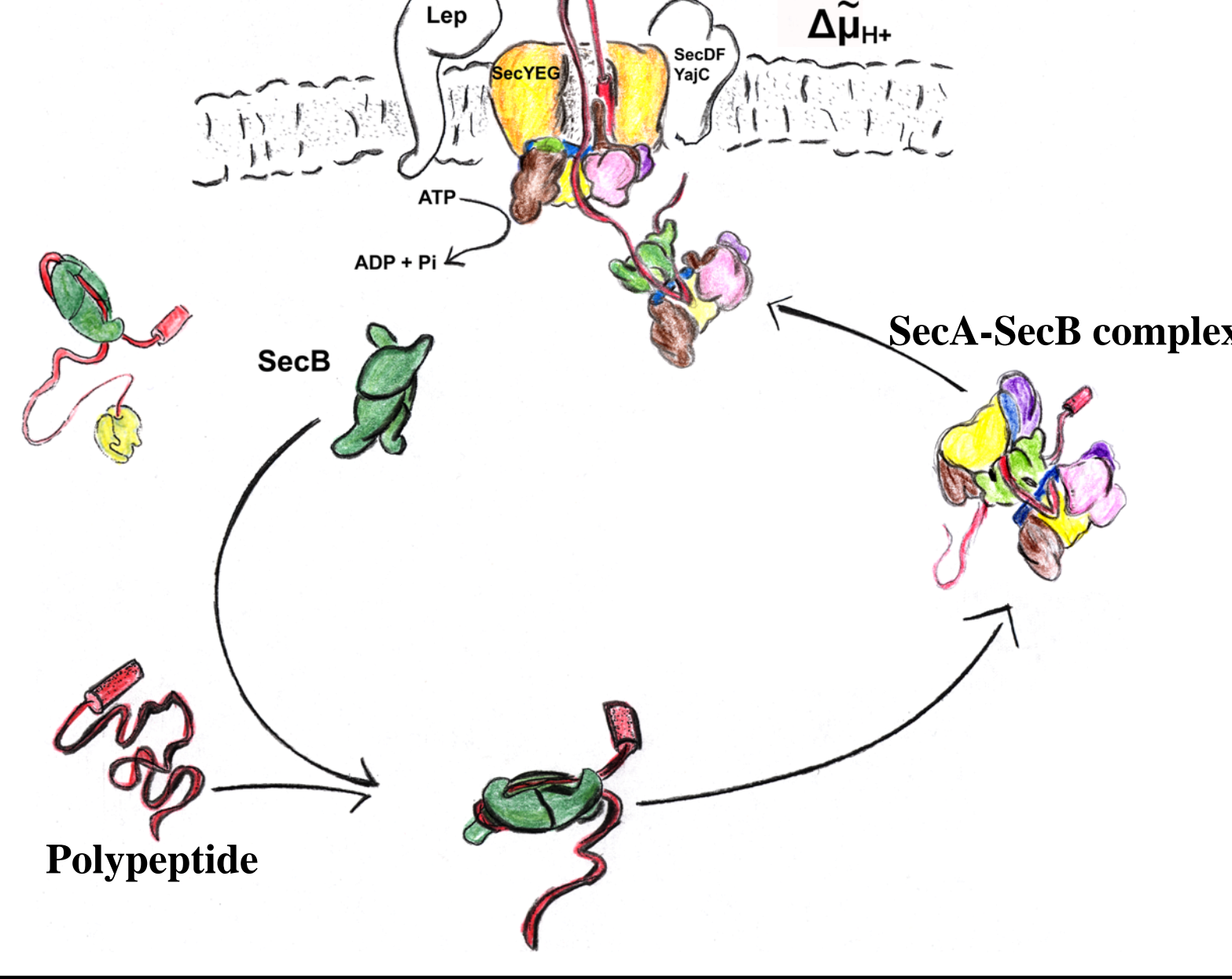
Raw glass vs. KOH treated glass



High Resolution Images of SecYEG

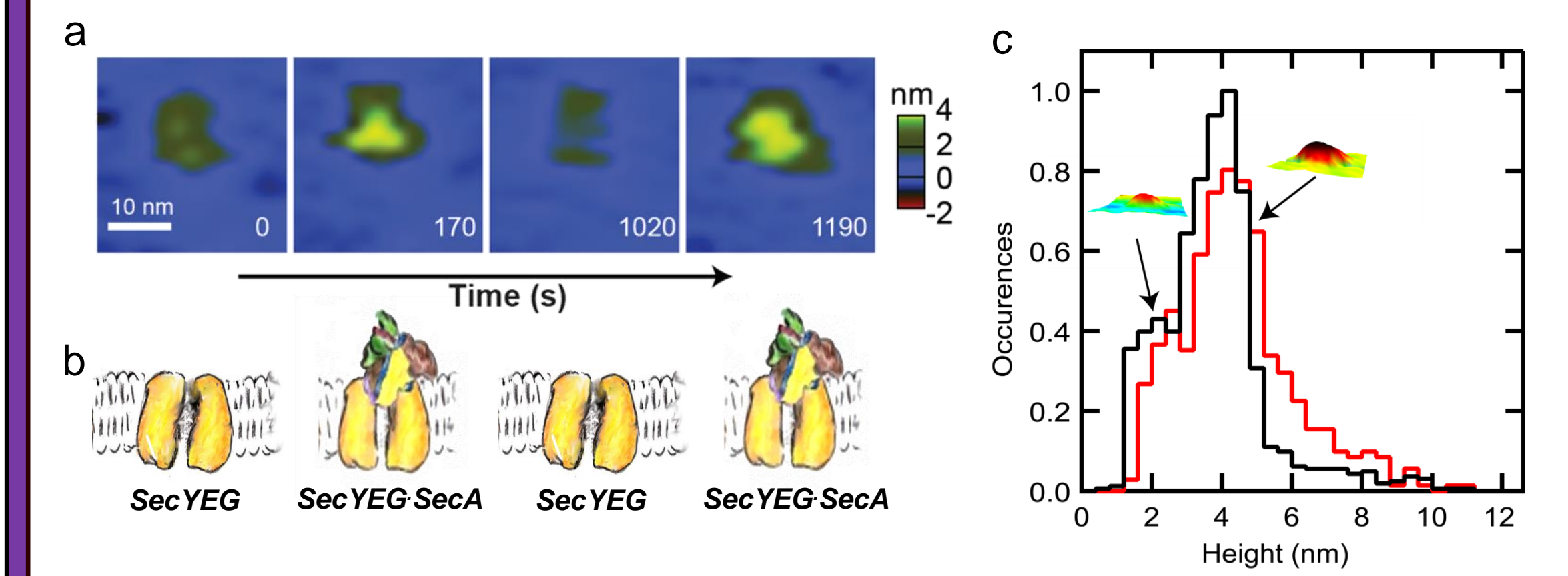


Sec-translocase

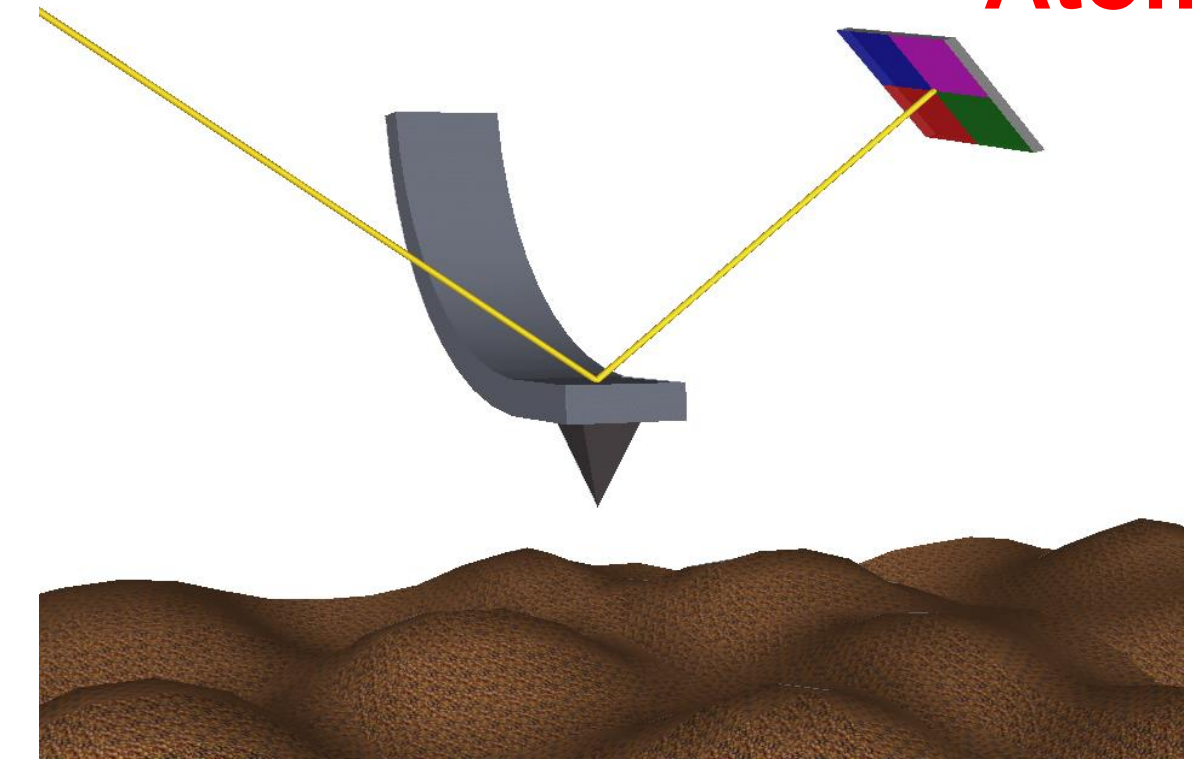


- >30% of proteins are transported from the site of synthesis into or through a membrane^{3,4}
- In *E. coli*, the Sec system orchestrates the translocation of polypeptides across membrane before they acquire stable tertiary structure and SecYEG provides the path way^{3,4}
- SecA and SecB act as chaperones to maintain newly synthesized polypeptides in a state compatible with transport⁶
- Numerous questions remain regarding the mechanistic details of translocation^{3,4,5}

SecYEG-SecA interactions

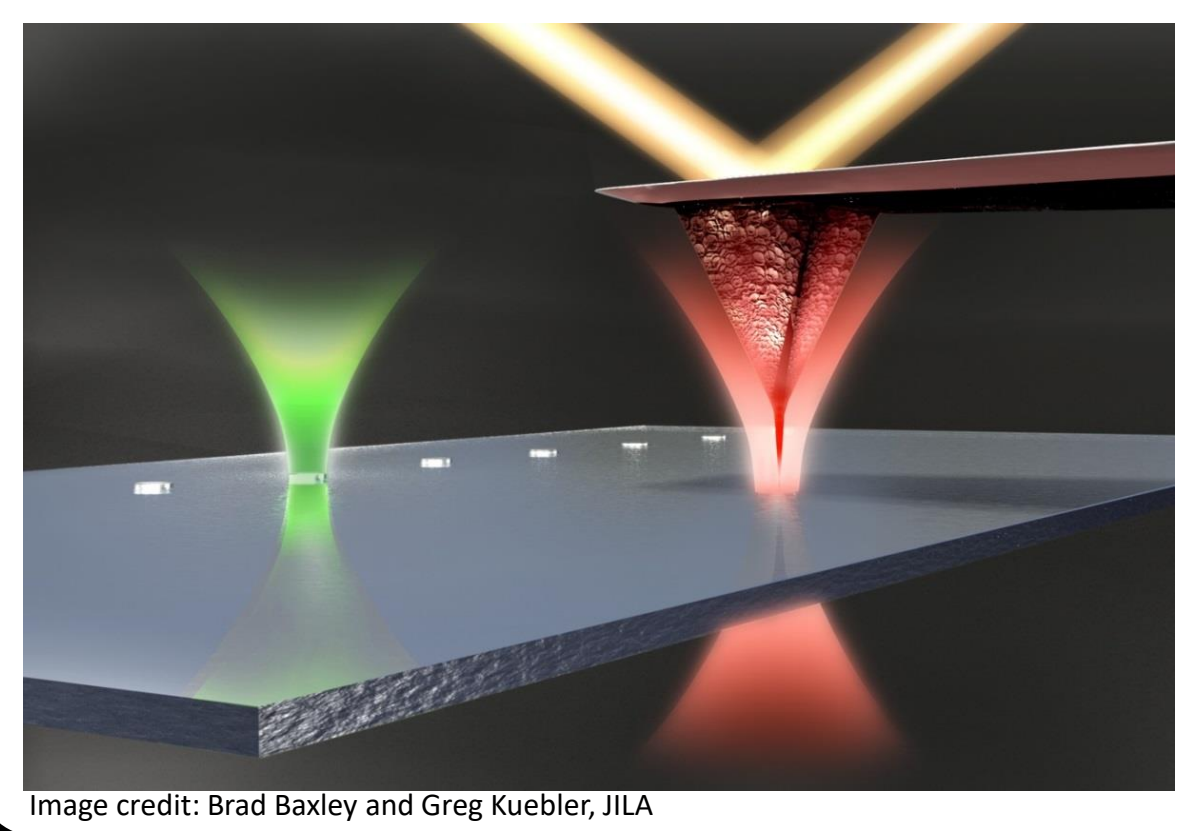


Atomic Force Microscopy (AFM)



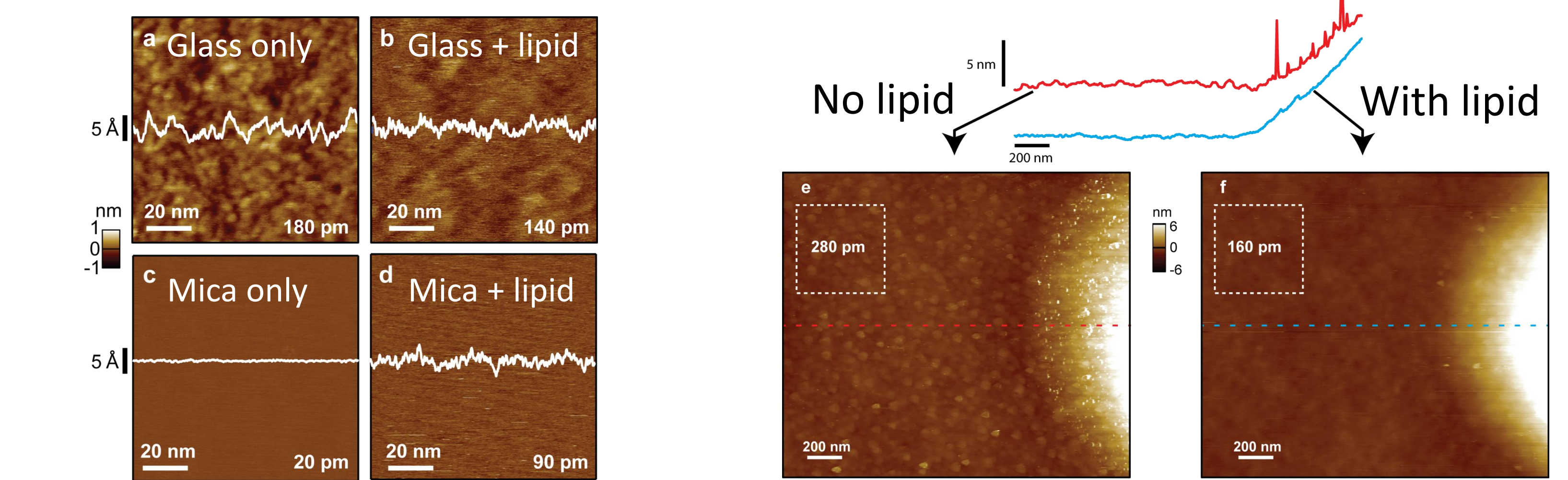
- Molecular-scale resolution³
- Imaging in near native lipid environment³
- Minimal concentration requirements³
- Single molecule dynamics³
- Drift (>1nm/min on each axis) is a common problem with commercial instruments in biological settings¹

Ultra Stable 3D AFM

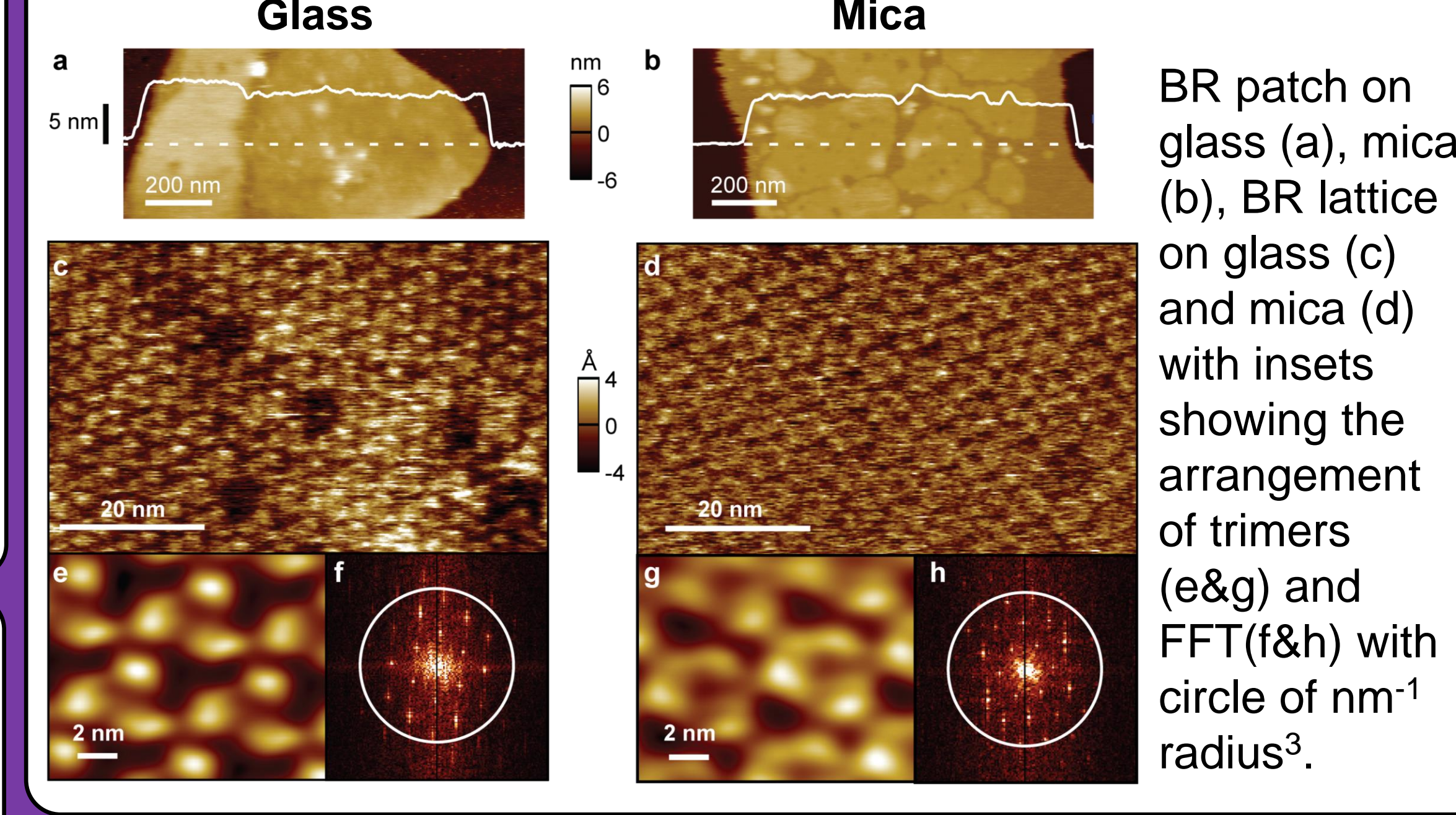


- Recent techniques^{1,2} employ two additional lasers, one focused on fiducial mark and other focused on AFM-tip for tip-sample registration and tip trajectory
- Drift can be reduced to <5pm/min at 25° C
- Direct access to **F**

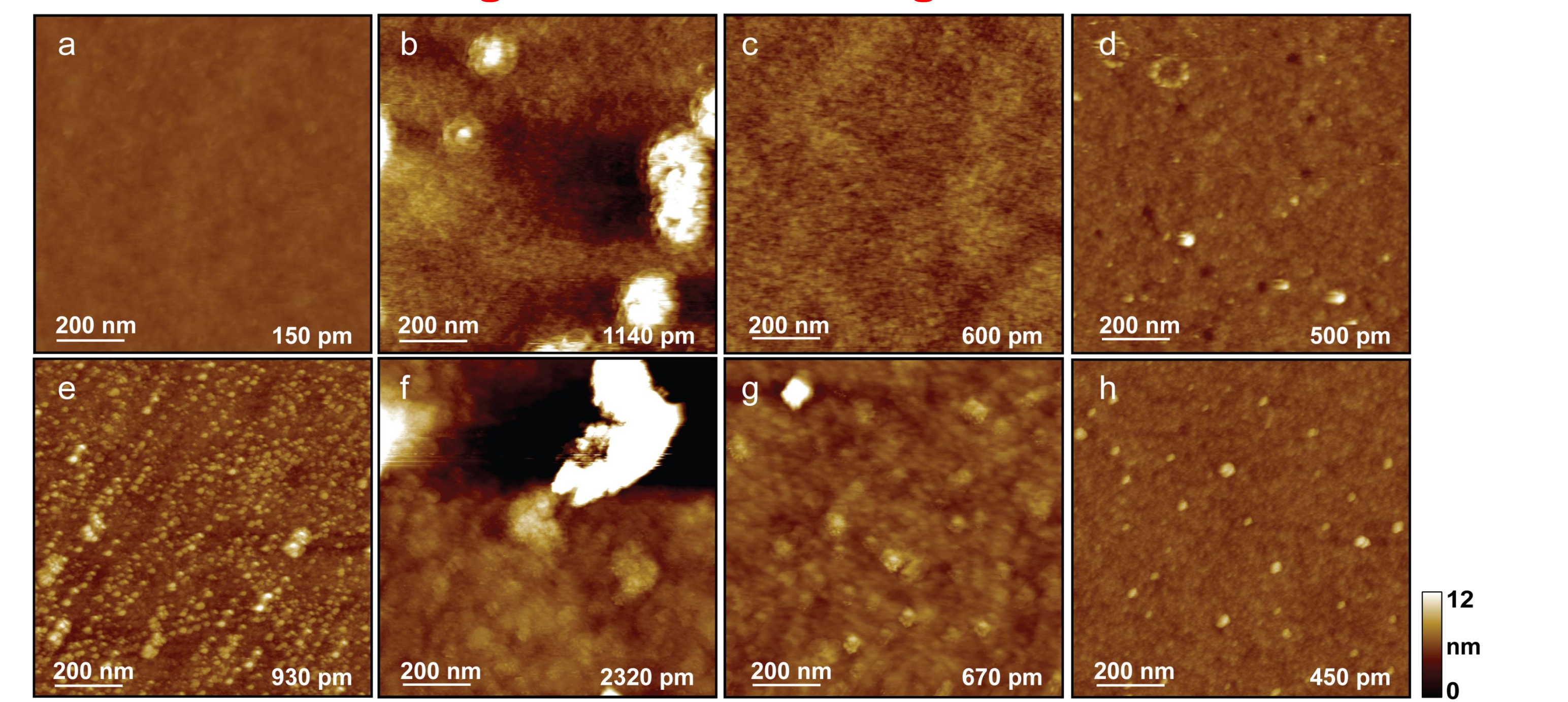
Direct visualization of reduction of surface roughness on glass



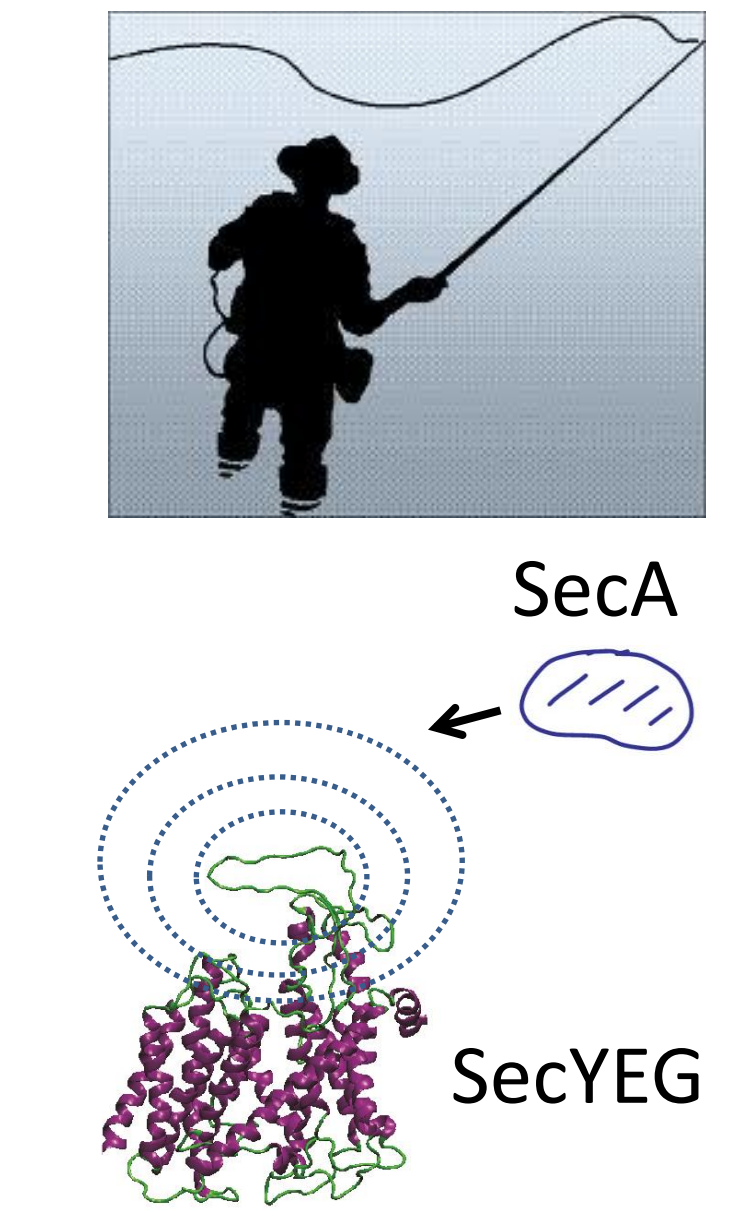
Bacteriorhodopsin on Glass and Mica



Evaluating Different Cleaning Treatments



Potential Applications



- Novel AFM modalities could measure molecular "fly fishing"⁵
- Map trajectories of protein domains in 3D
- Dynamic Loops of SecYEG, linking transmembrane helices 6-7 and 8-9, play a vital role in capturing SecA and in translocation
- Probing translocation of precursor through Sec-translocase

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Acknowledgements

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