

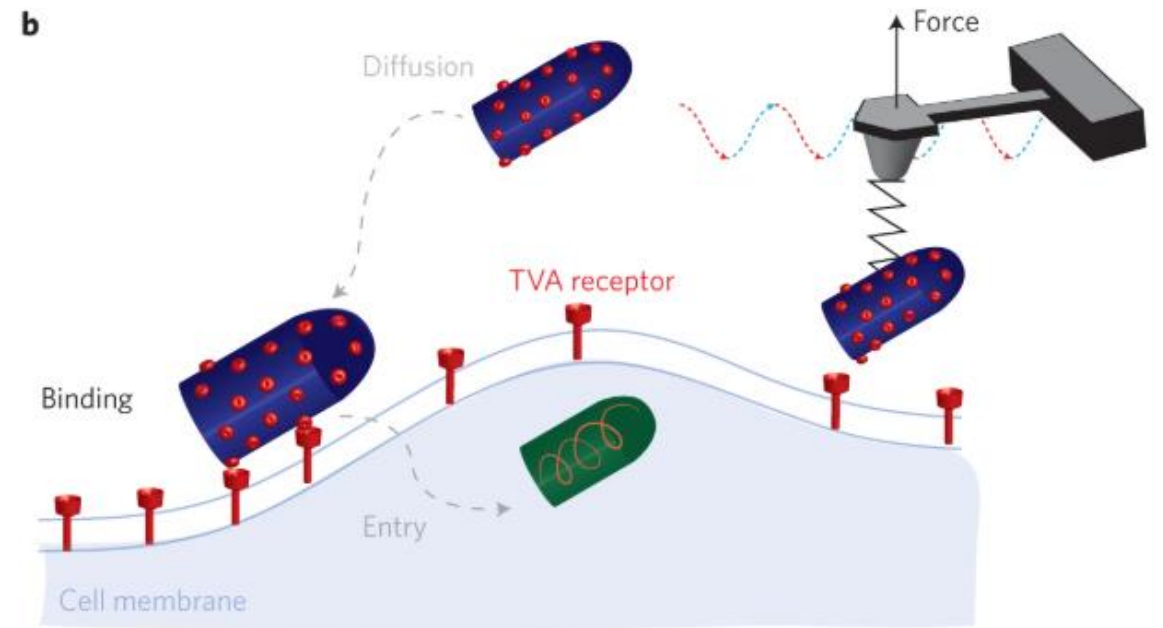
High-speed AFM and other special applications

Luca Elizabet Kosik, Miklós Cervenak, Dominik Sziklai, Andrea Balogh-Molnár, Bence Fehér, Soma Yamamoto, Hiroki Konno, Noriyuki Kodera, Holger Flechsig, Romain Amyot, Hedvig Tordai, Levente Herényi, Ana Cuervo, Miklós Kellermayer, Bálint Kiss

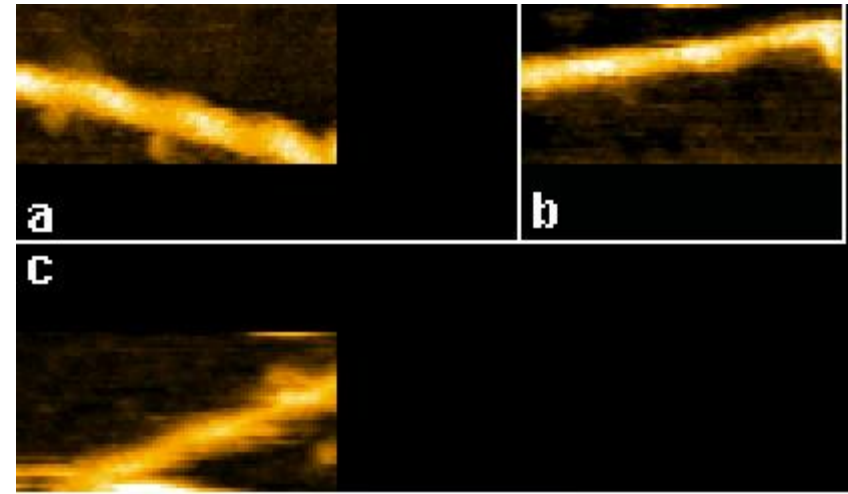


Special approaches in AFM

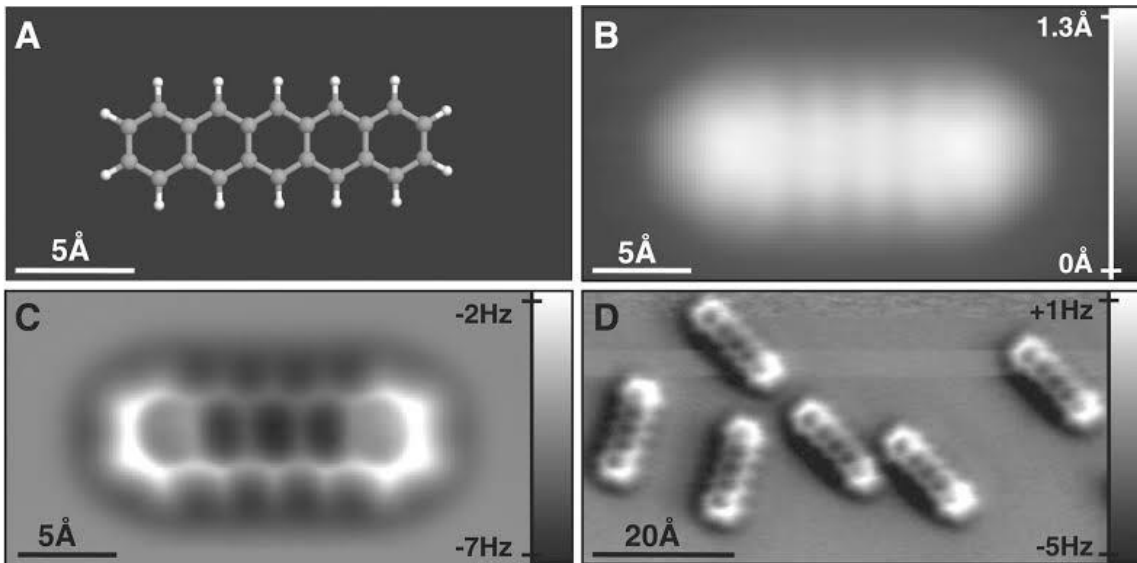
- Tip functionalization
- Surface modification
- Atomic resolution
- High-speed scanning
- Nanoendoscopy



Alsteens, D., Newton, R., Schubert, R. *et al.* Nanomechanical mapping of first binding steps of a virus to animal cells. *Nature Nanotech* **12**, 177–183 (2017).
<https://doi.org/10.1038/nnano.2016.228>



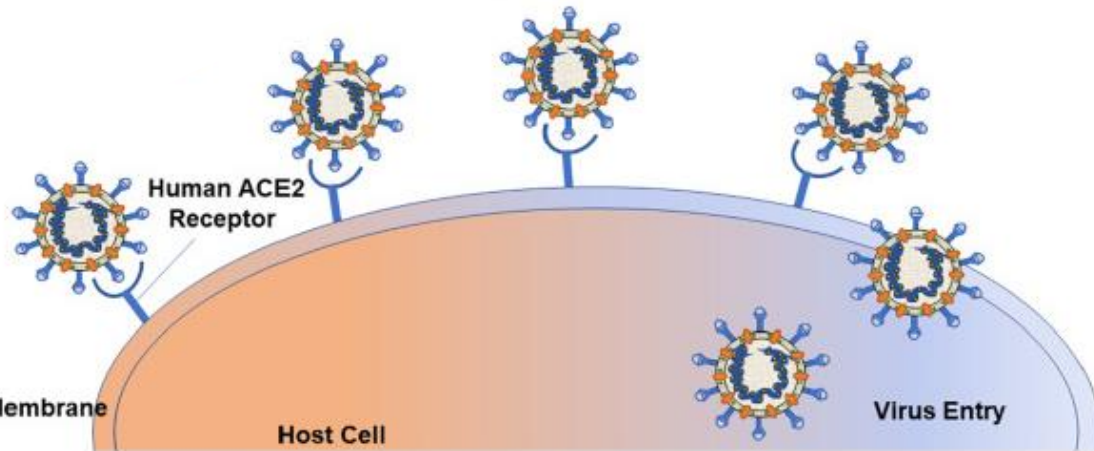
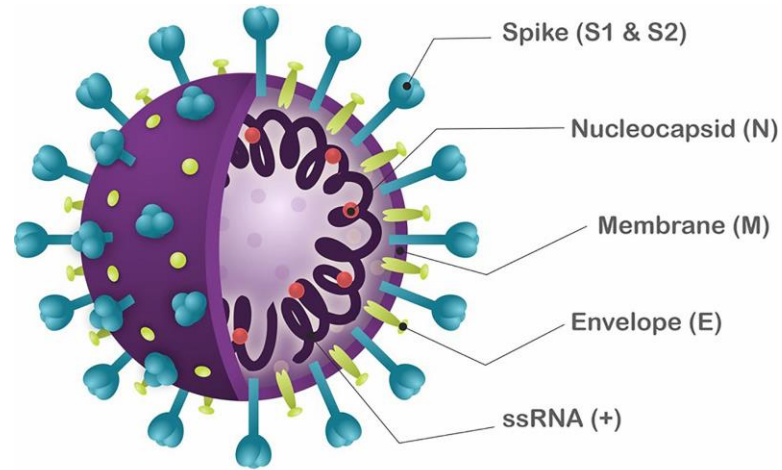
Kodera, N., Yamamoto, D., Ishikawa, R. *et al.* Video imaging of walking myosin V by high-speed atomic force microscopy. *Nature* **468**, 72–76 (2010). <https://doi.org/10.1038/nature09450>



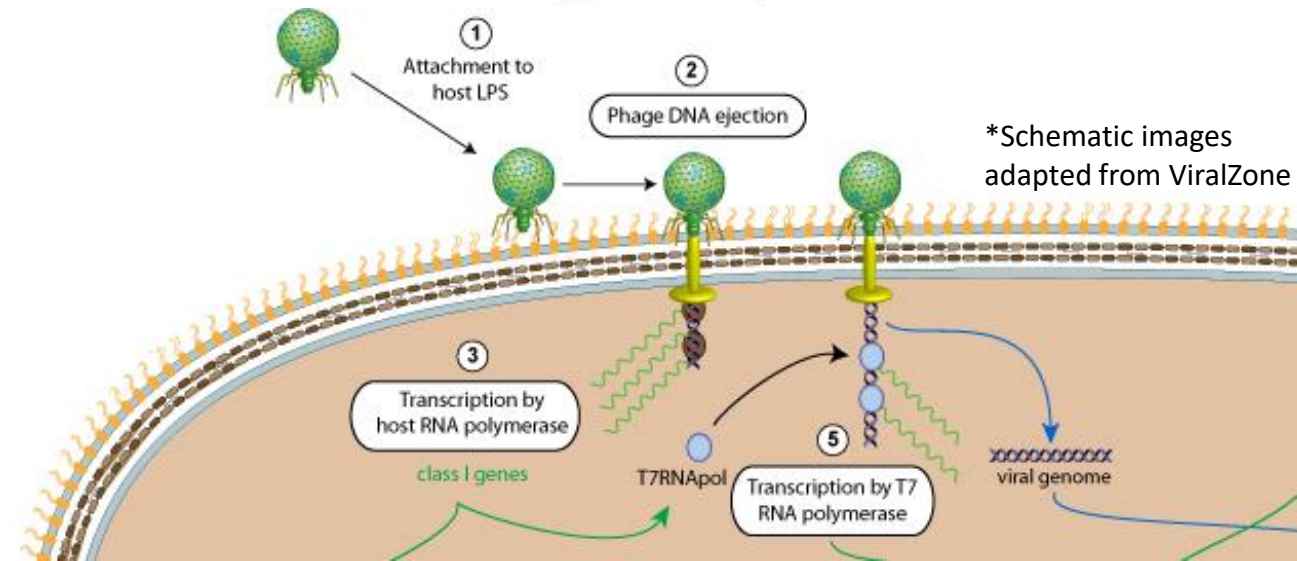
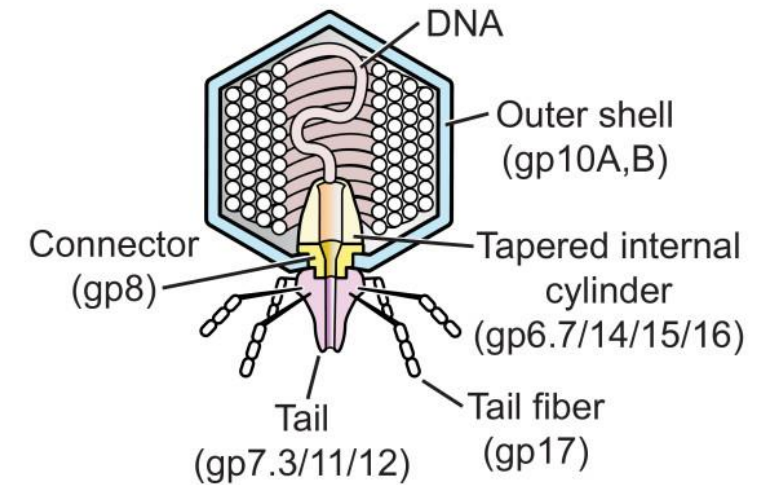
Leo Gross *et al.*, The Chemical Structure of a Molecule Resolved by Atomic Force Microscopy. *Science* **325**, 1110–1114 (2009). DOI:10.1126/science.1176210

Two viruses with two different infection strategies

SARS-CoV-2

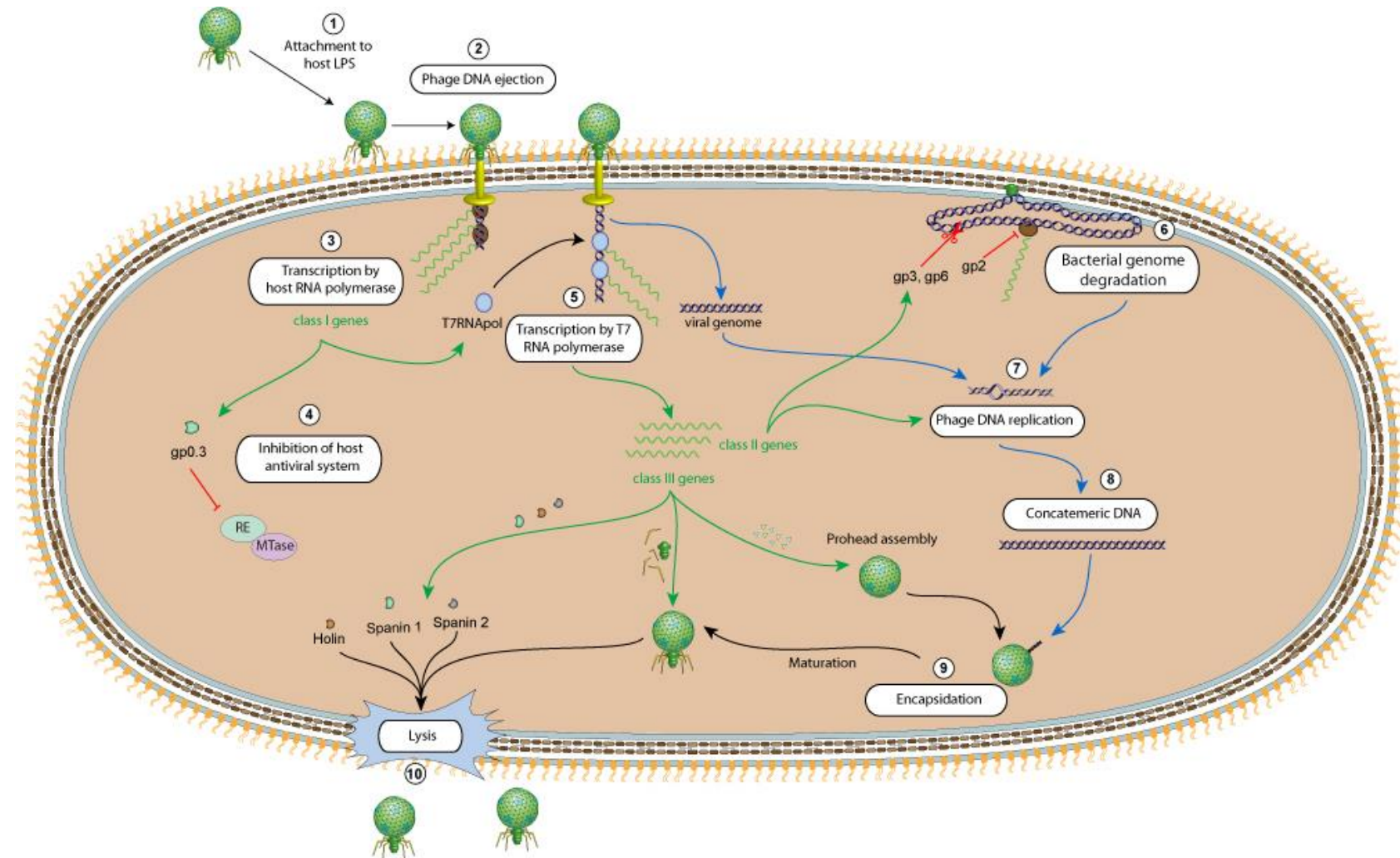
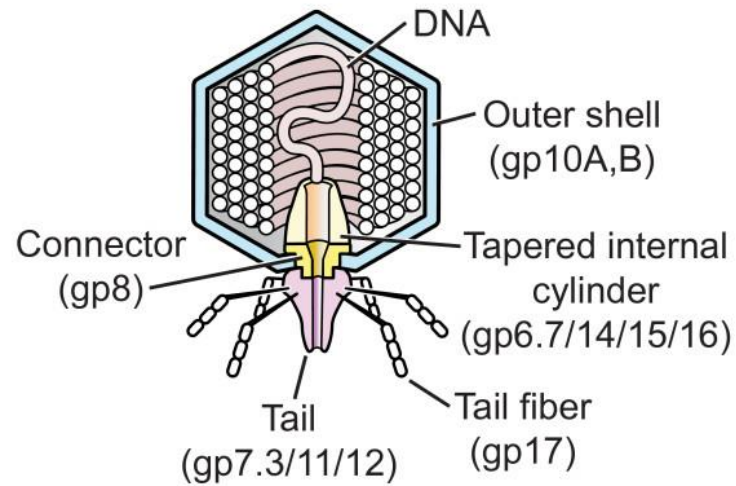


T7 bacteriophage



T7 bacteriophage

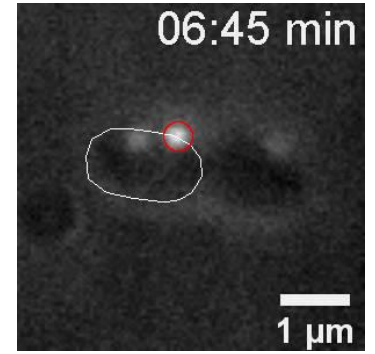
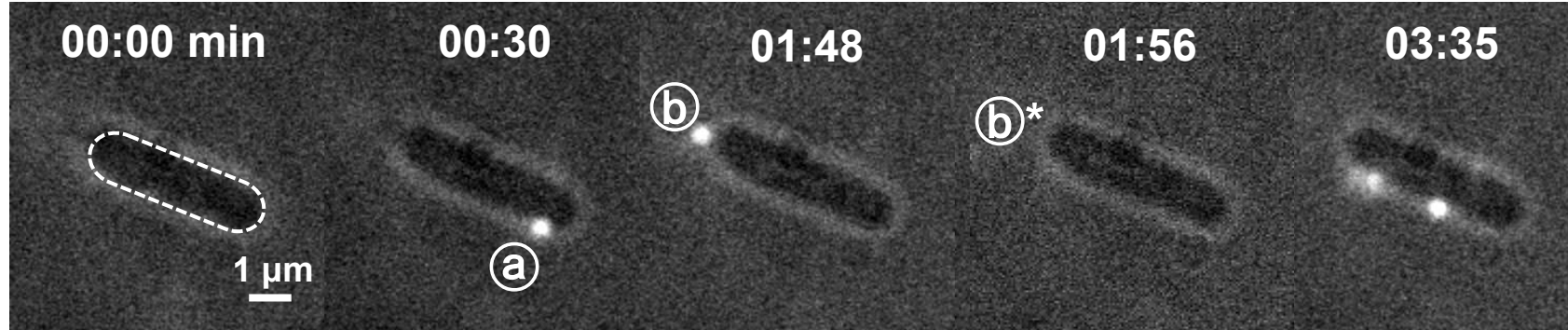
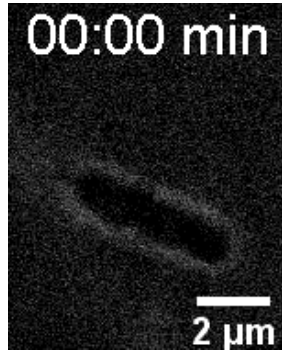
T7 bacteriophage



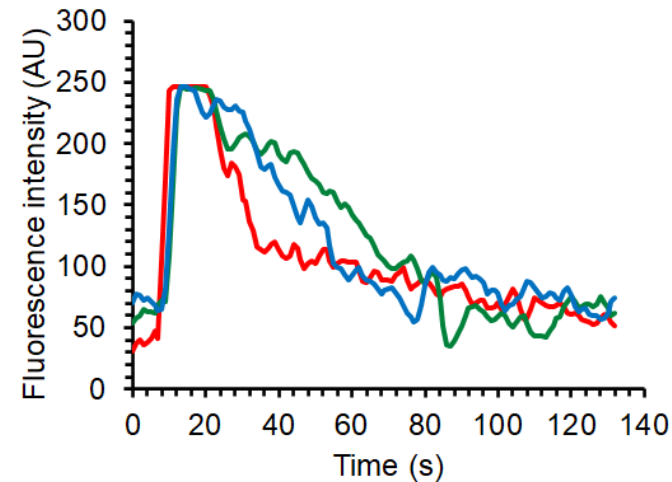
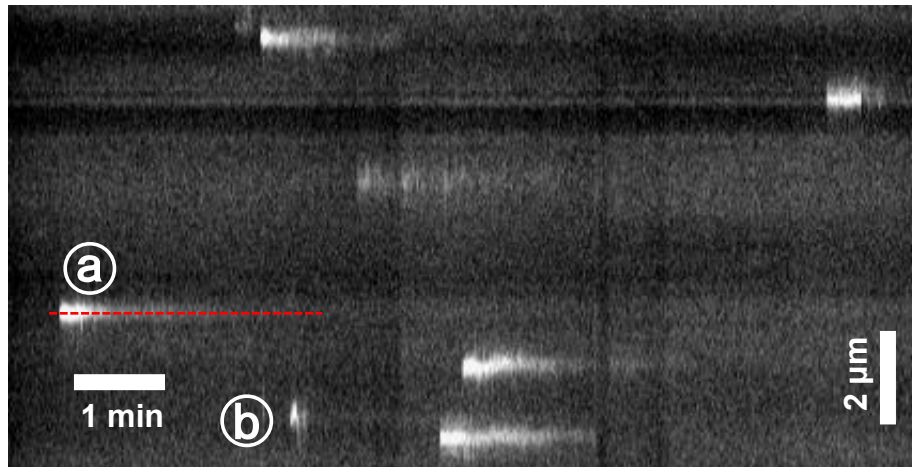
*Schematic images adapted from ViralZone

T7 phage docking is a dynamic, two-dimensional searching process

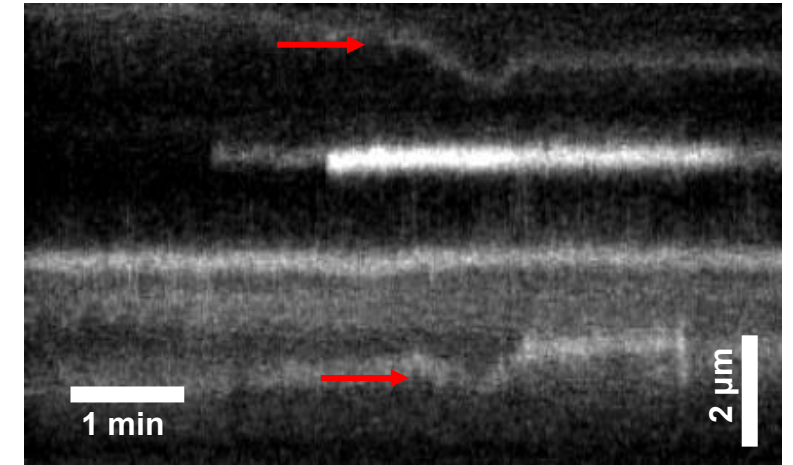
Surface binding



Docking and DNA injection

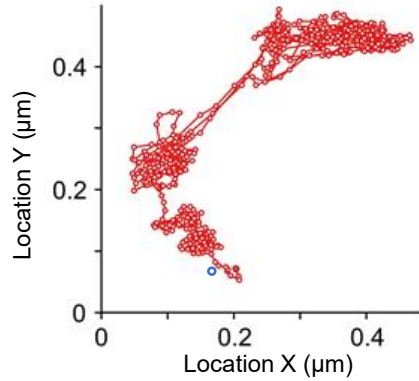
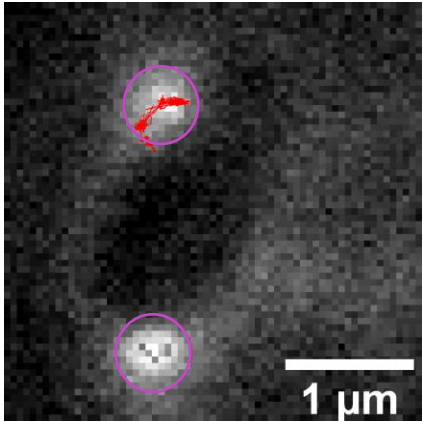


Surface exploration

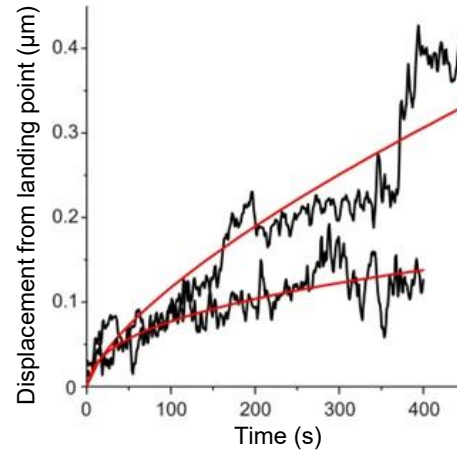


Phage docking is a dynamic, two-dimensional searching process

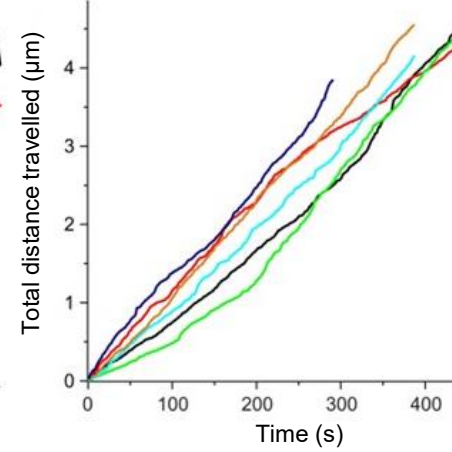
Exploration trajectory



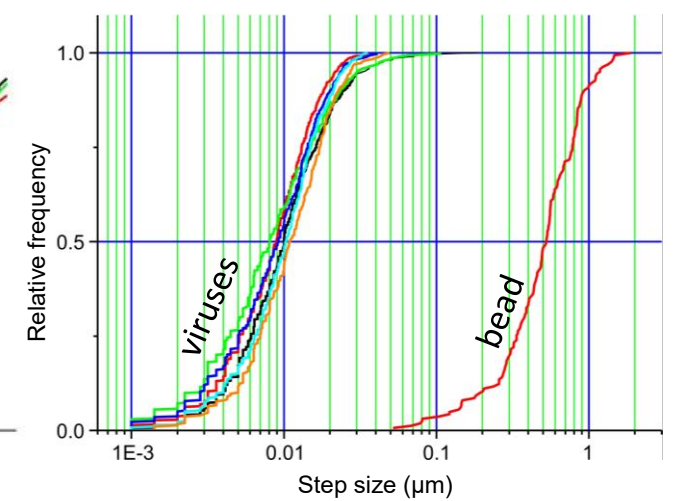
Displacement from landing



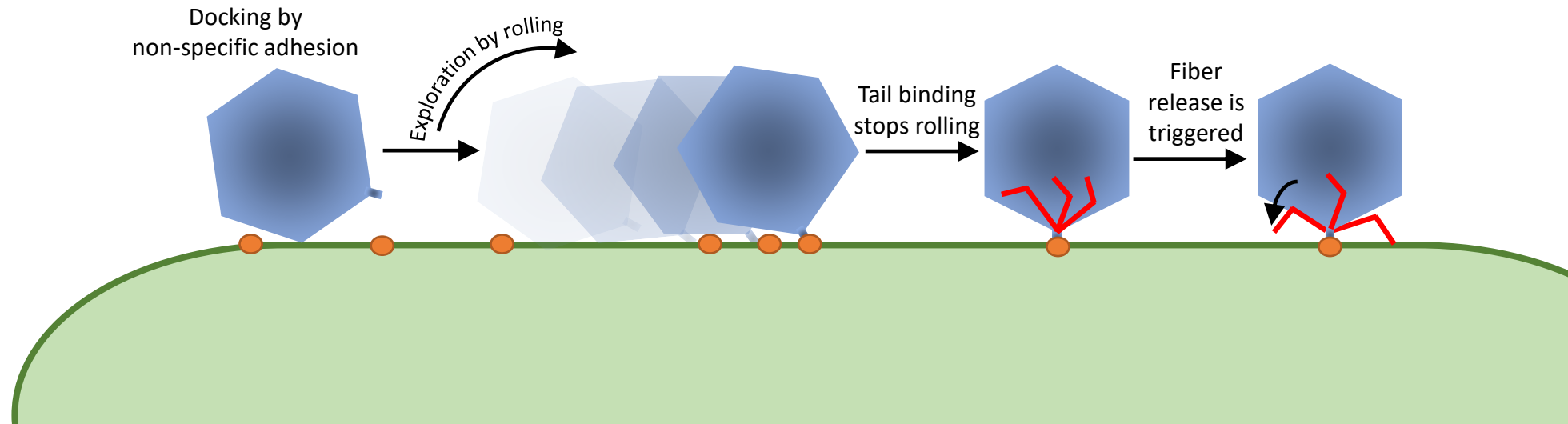
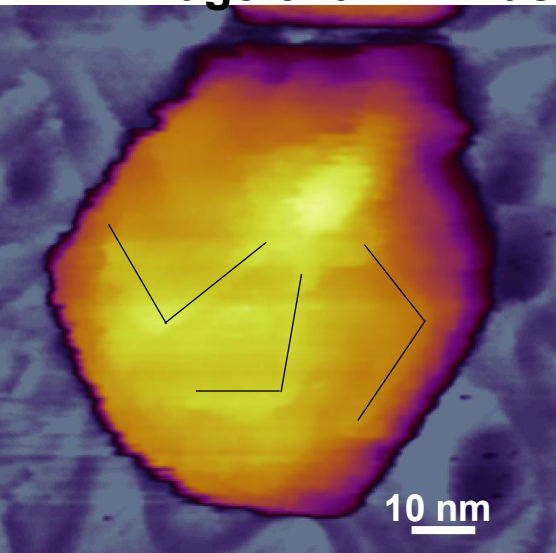
Distance travelled over time



Nanoparticle step sizes

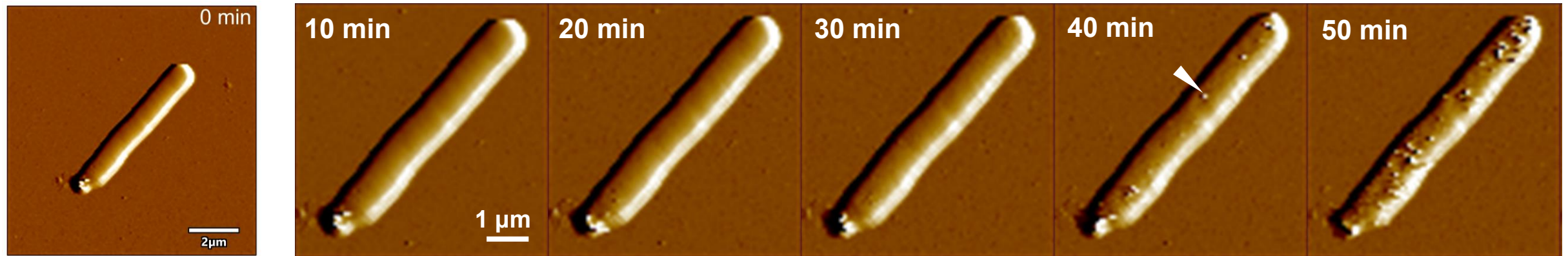


AFM image of a T7 virus

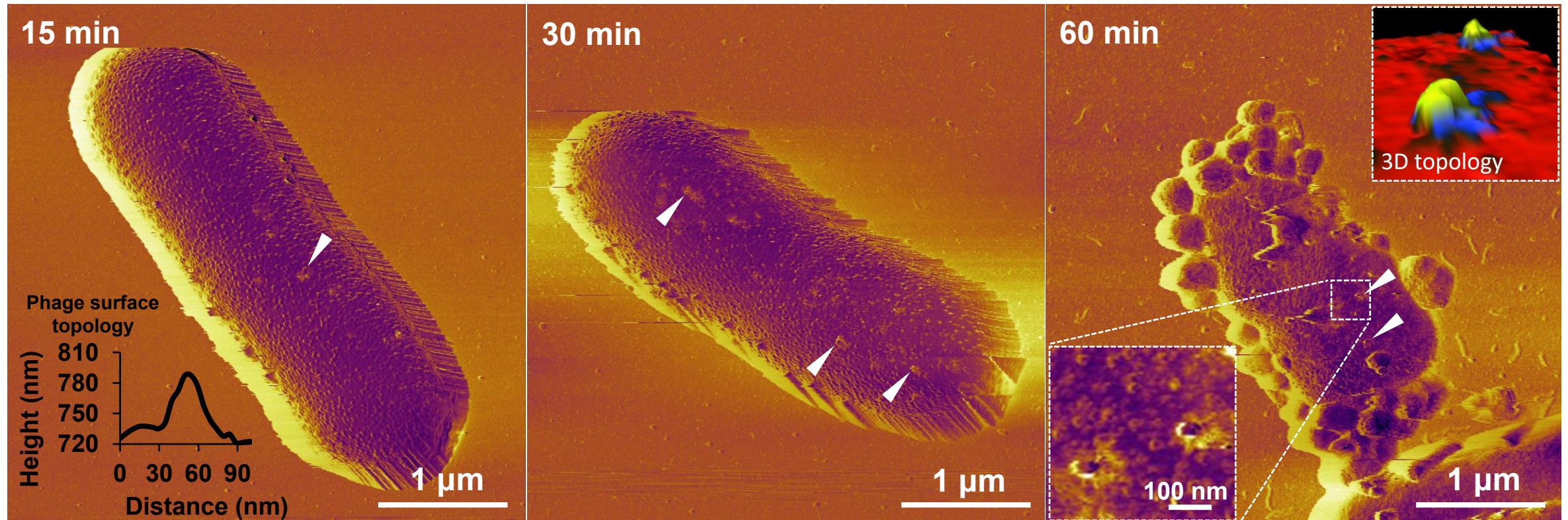


Phages saturate the bacterial surface by binding via their tail fibers

Native

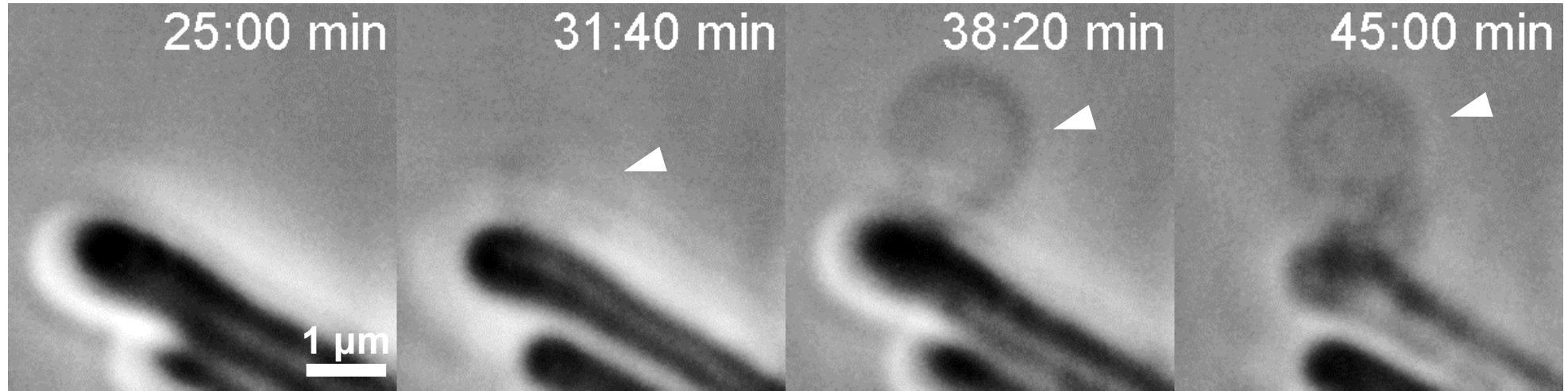
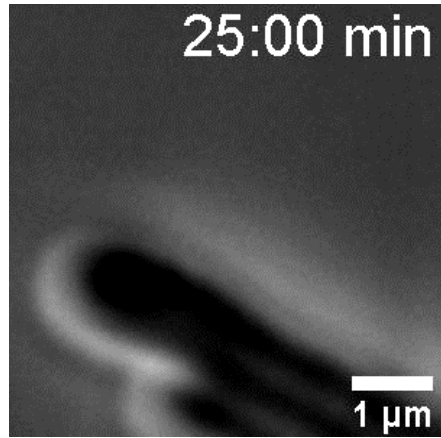


Chemically fixed

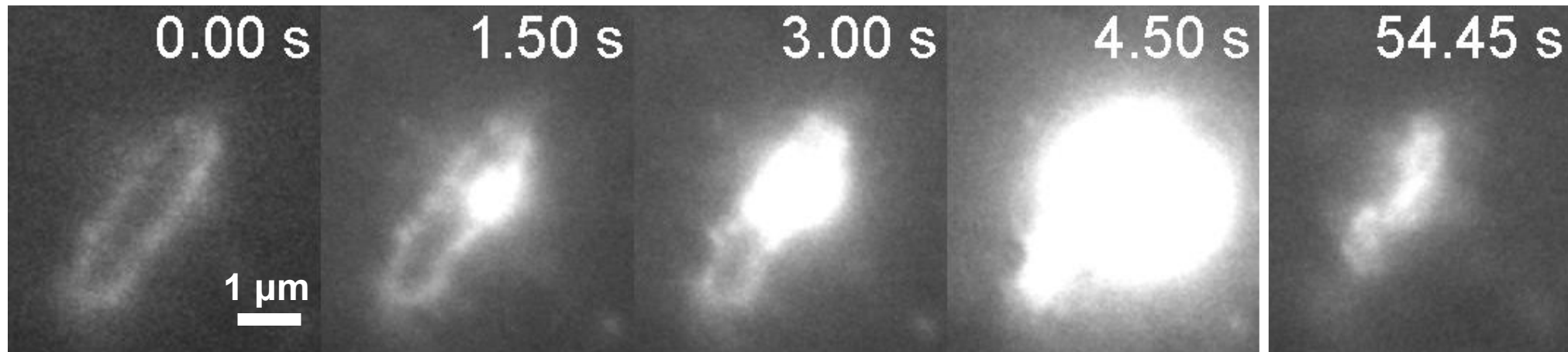
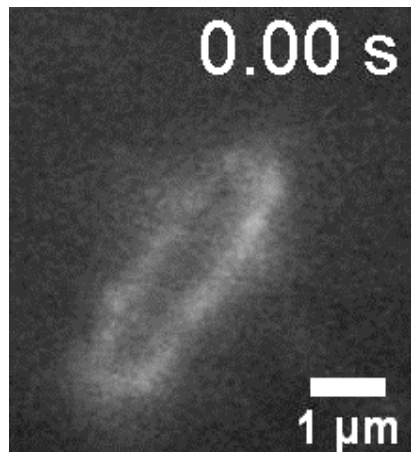


Viral infection leads to bleb formation and a violent host cell lysis

Bleb formation

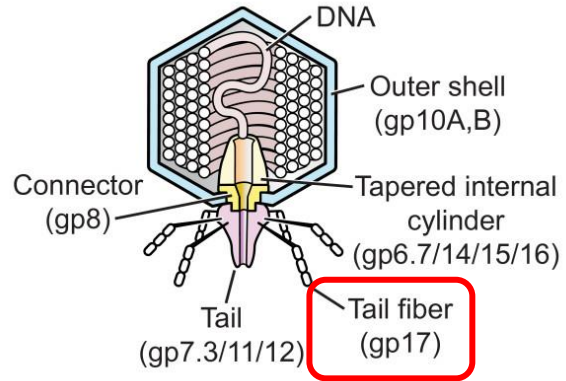


Bacterial lysis

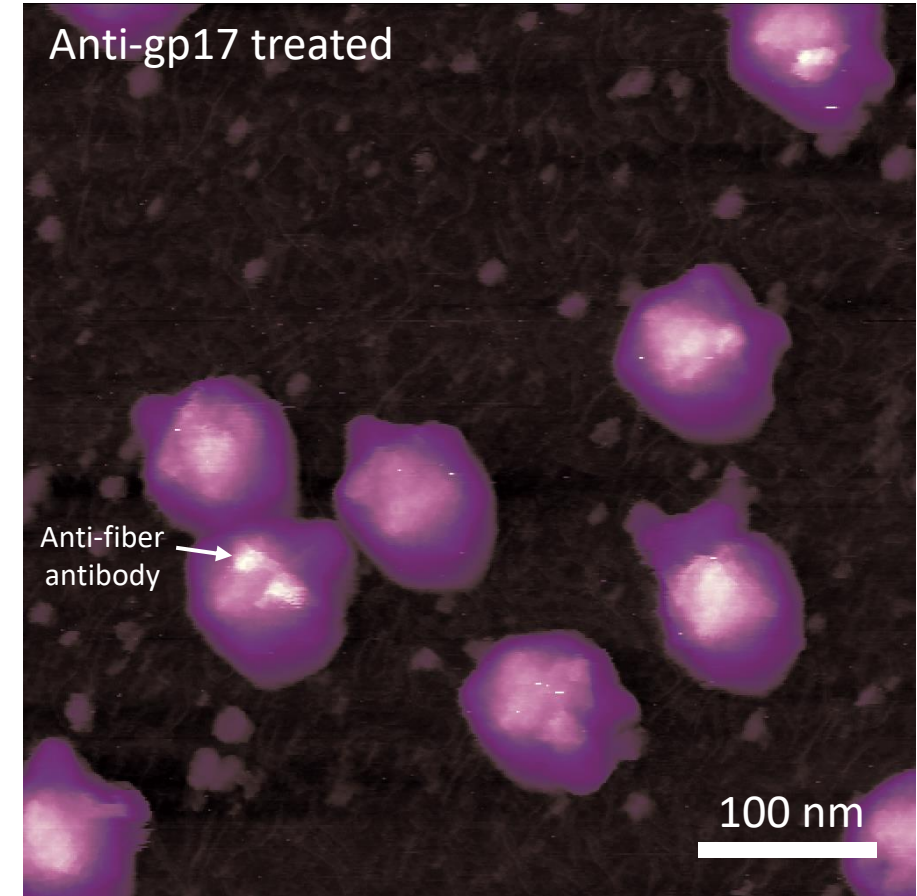
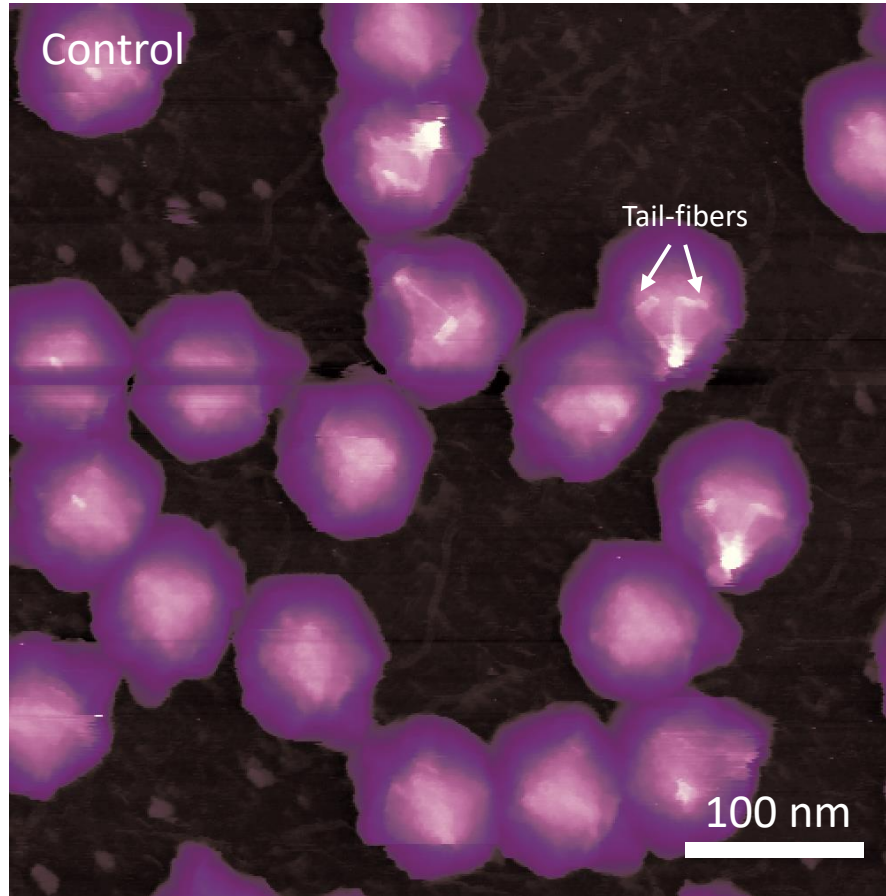


T7 tail fiber nanomechanics

T7 bacteriophage

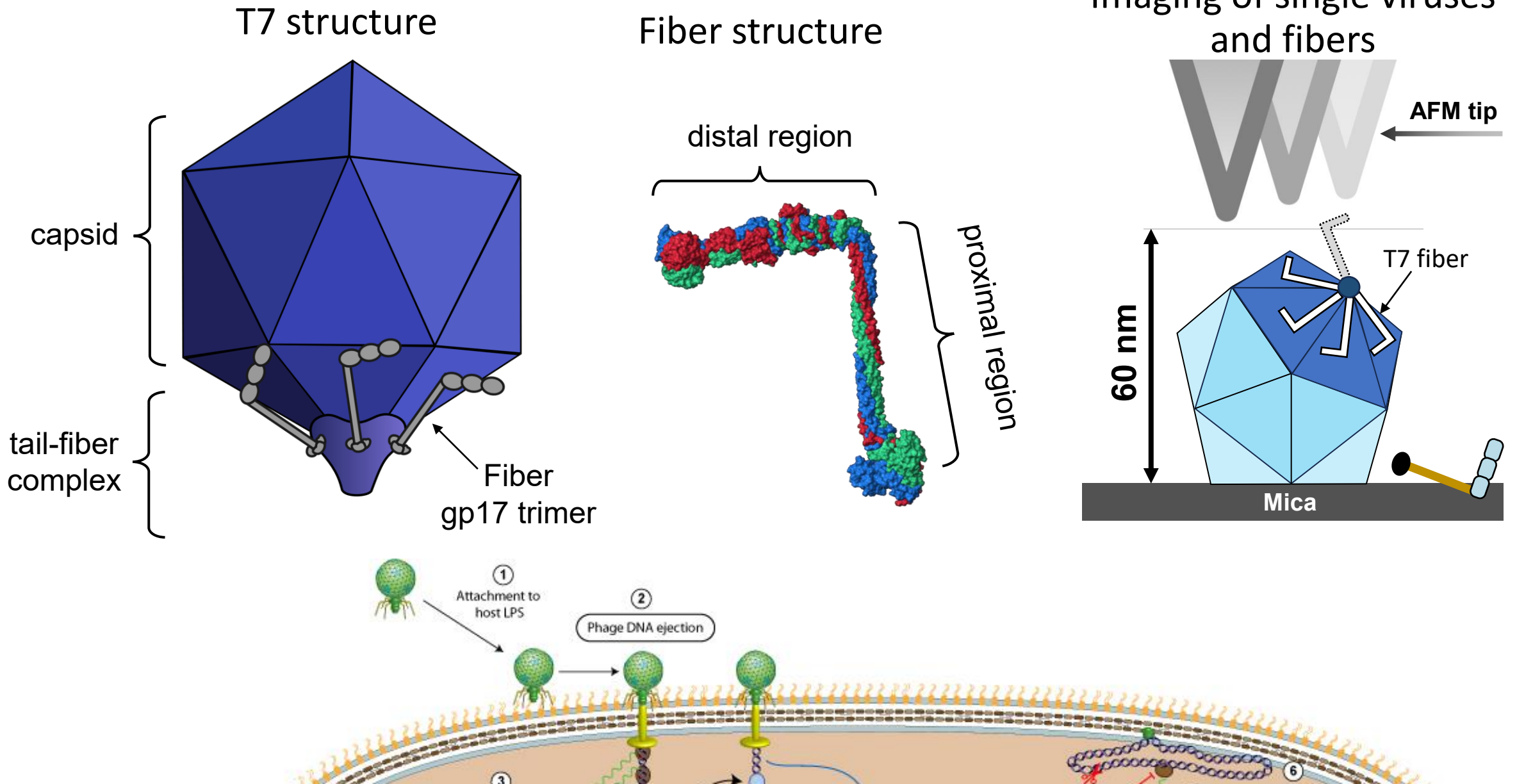


Tail fiber specific antibody, but what is its exact target?



Antibody binds to the fibers' proximal region

T7 phage and fiber structure

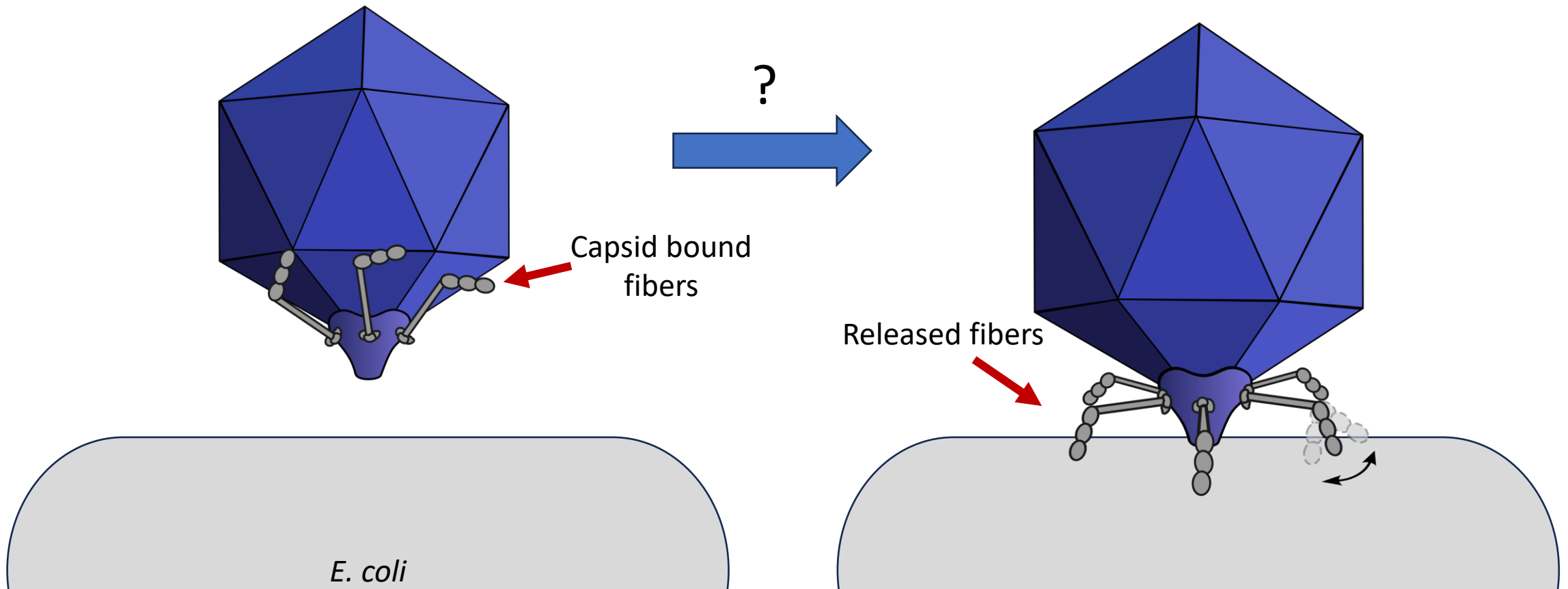


How do fibers transition into extended/infective state?

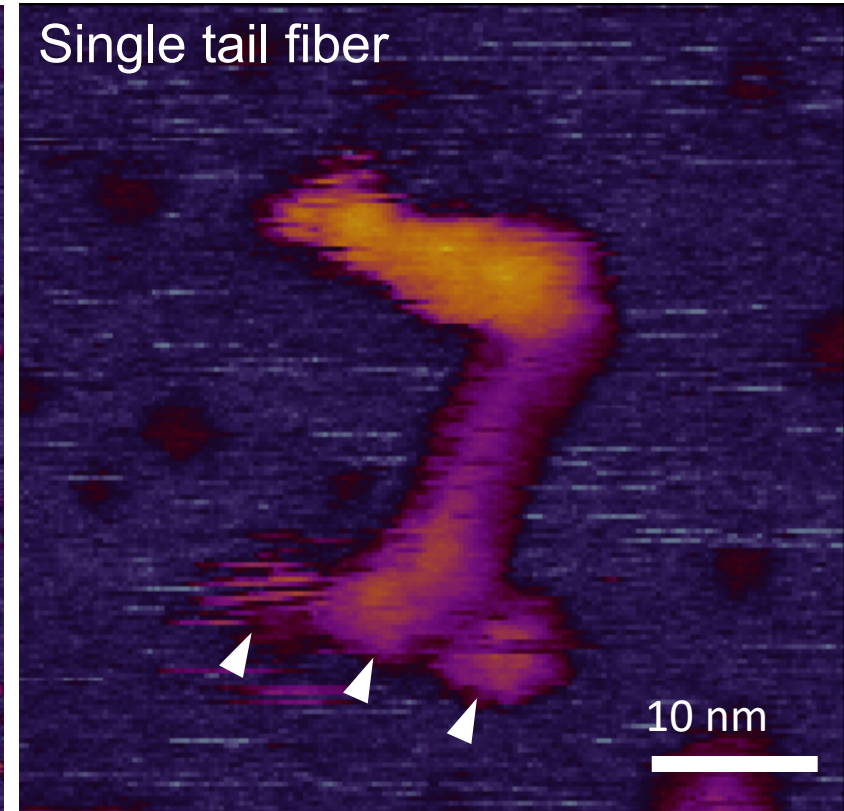
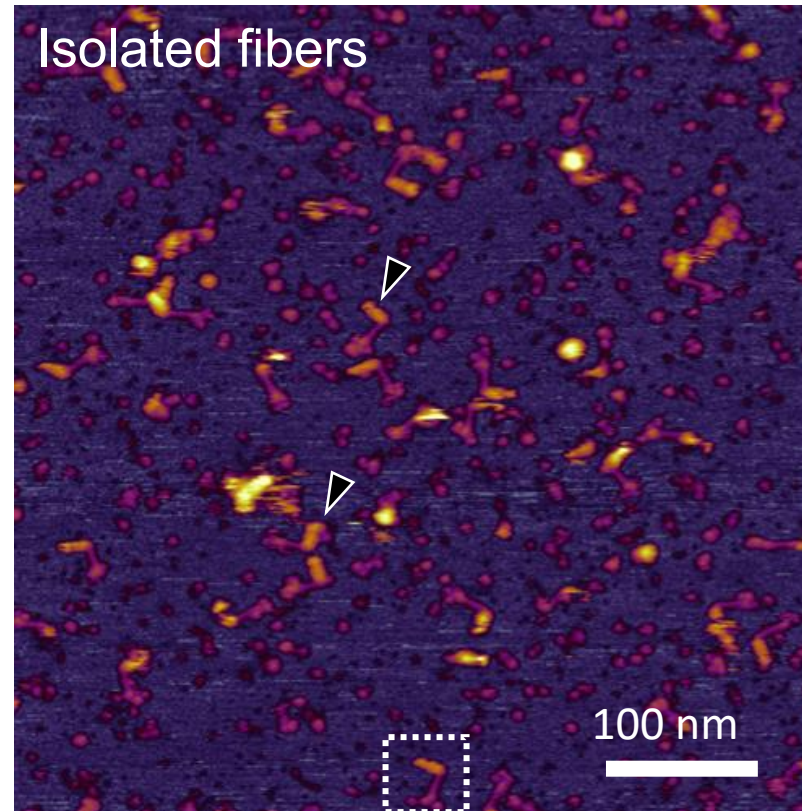
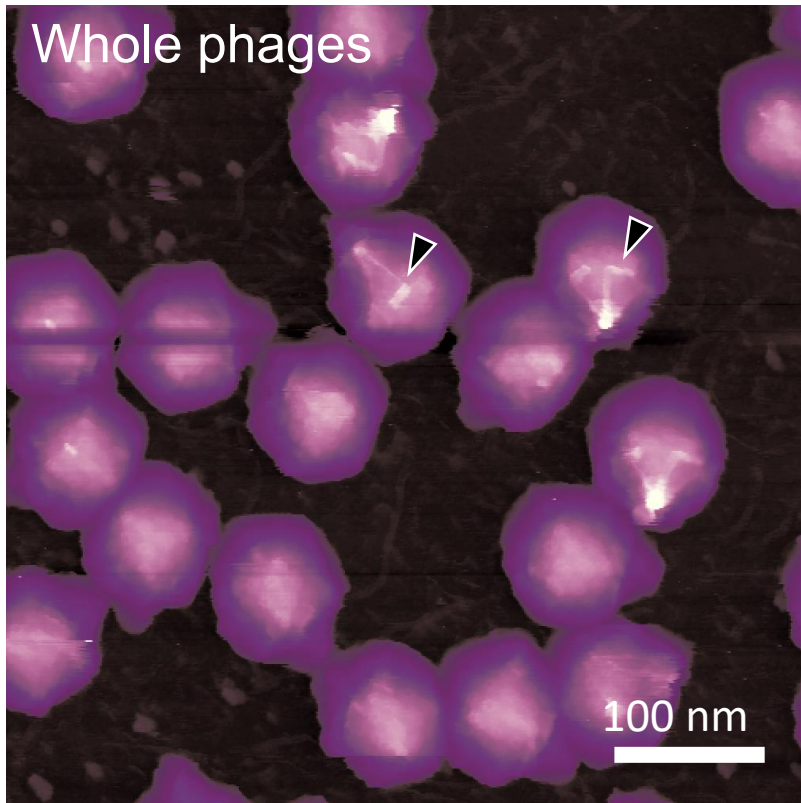
Is “walking” possible on the bacterial surface?

Unbound state

Infective state

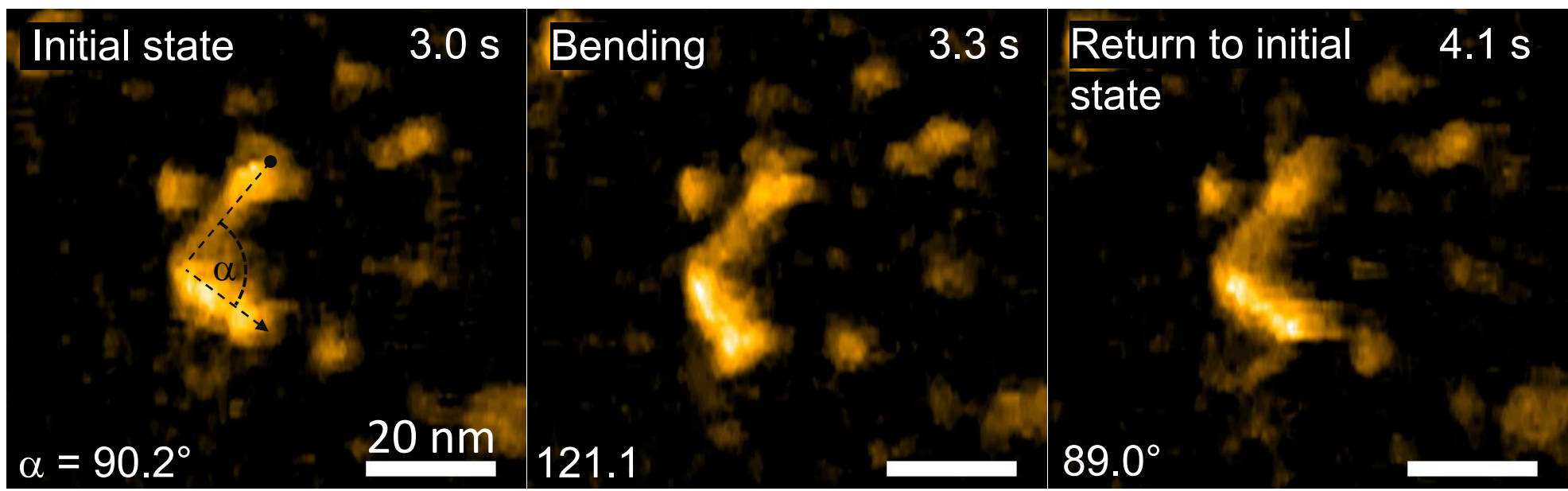
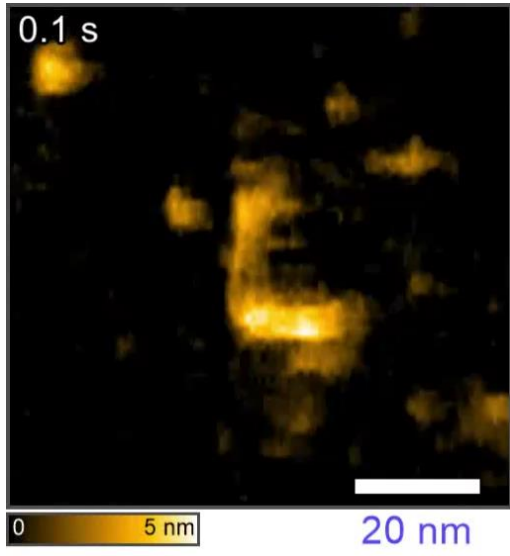


Isolated fibers of T7 bacteriophage

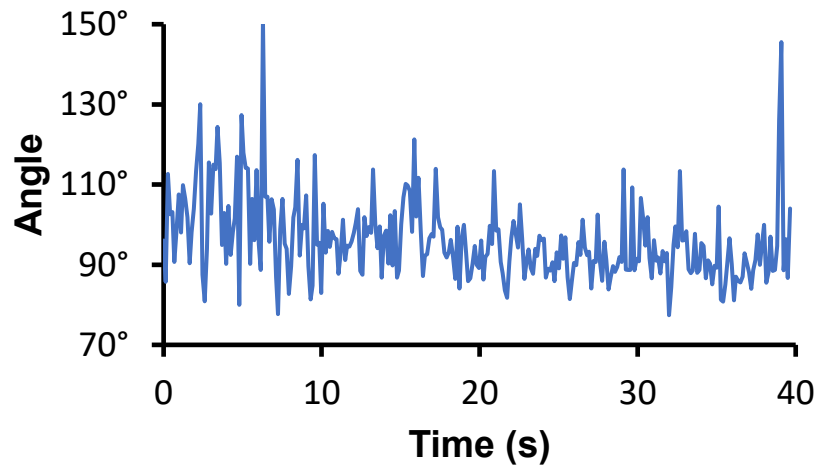


Fibers have a wide bending angle distribution

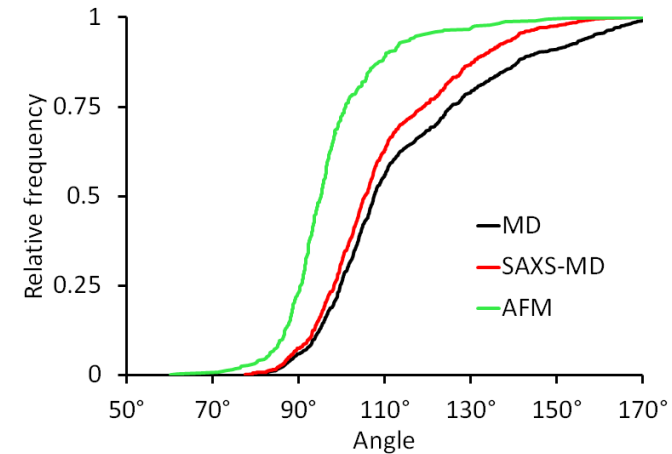
Hinge movement



Bending angle over time

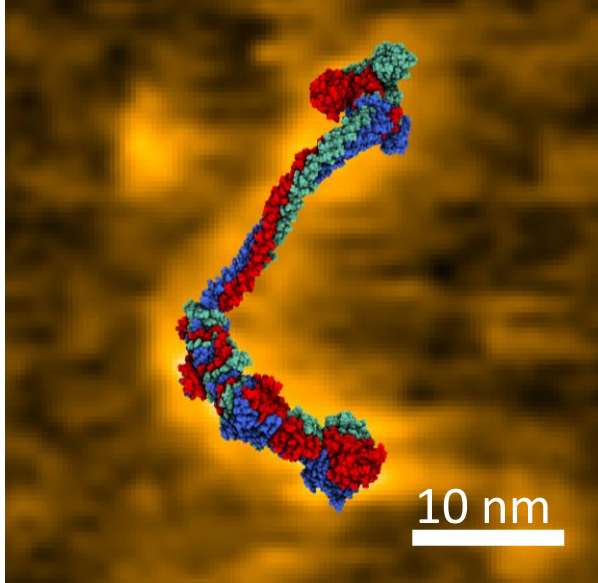


Bending angle distribution

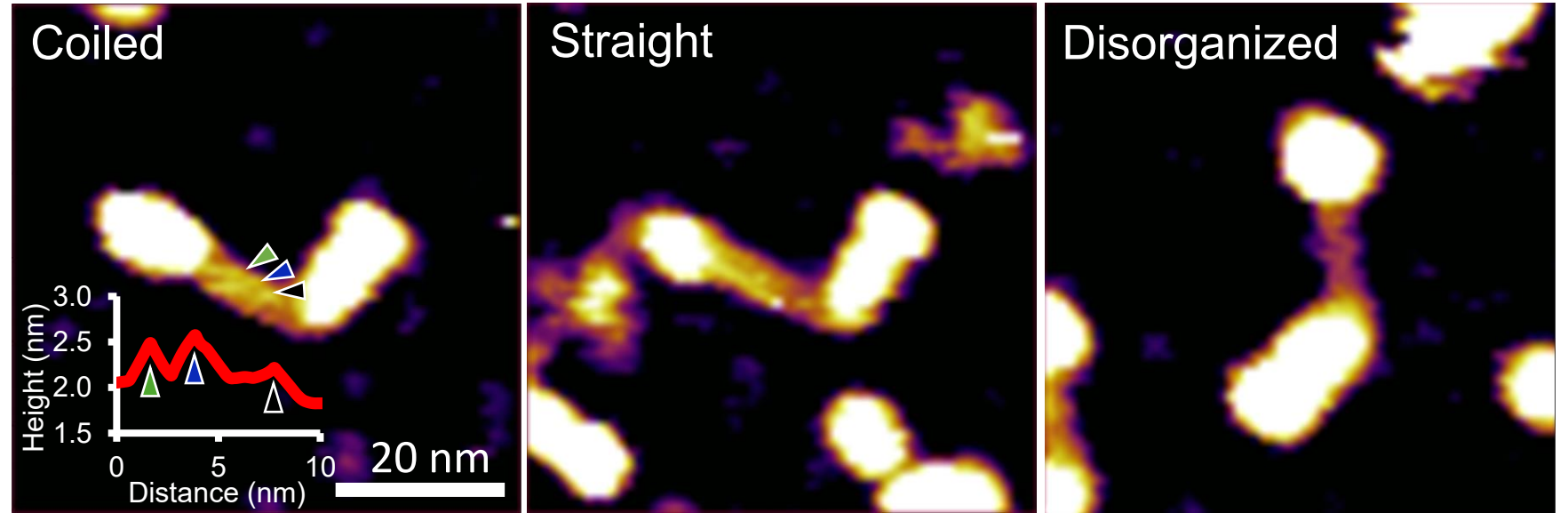


Fiber proximal section detailed structure

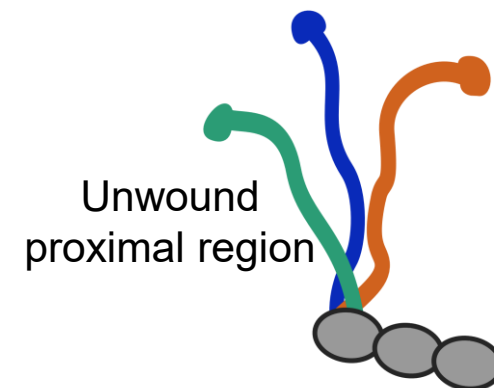
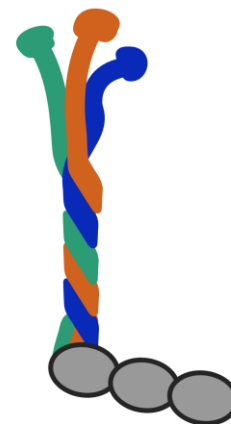
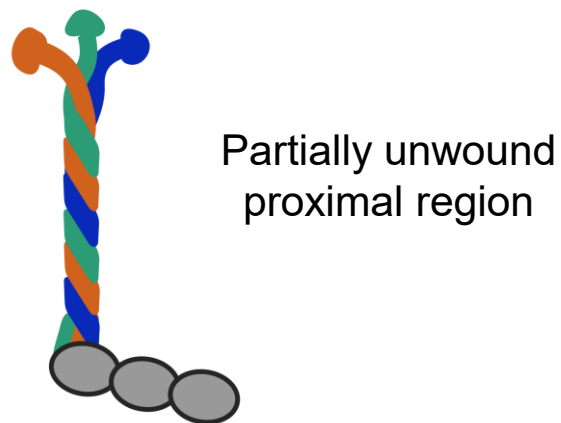
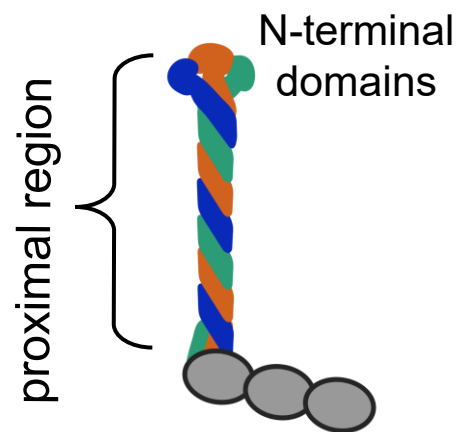
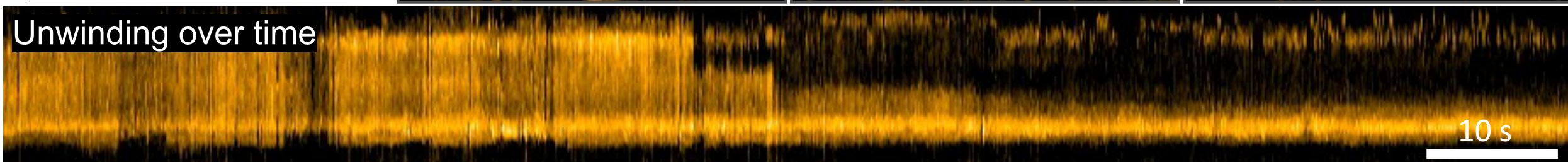
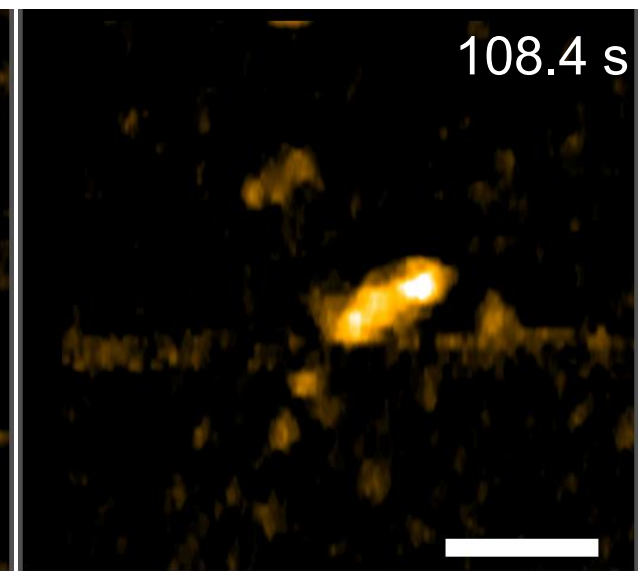
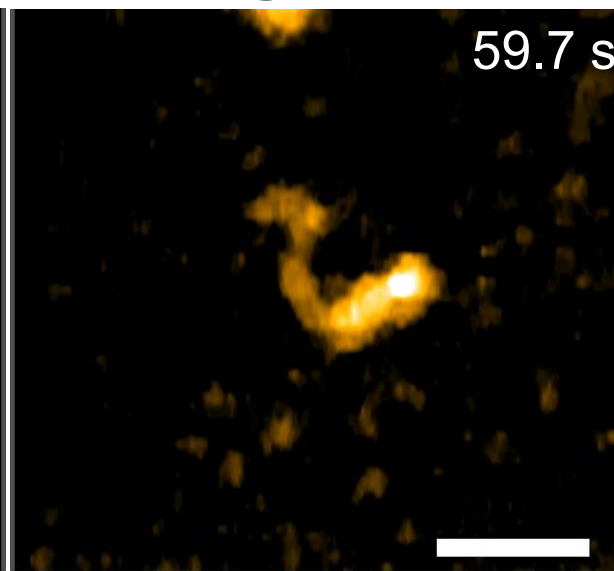
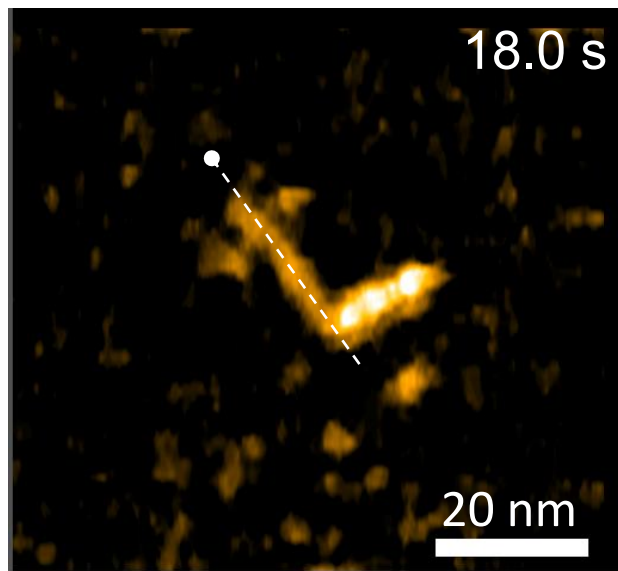
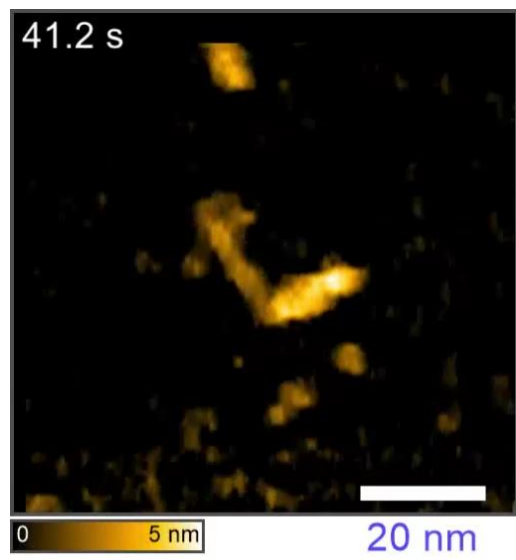
Structural fitting (NMFF)



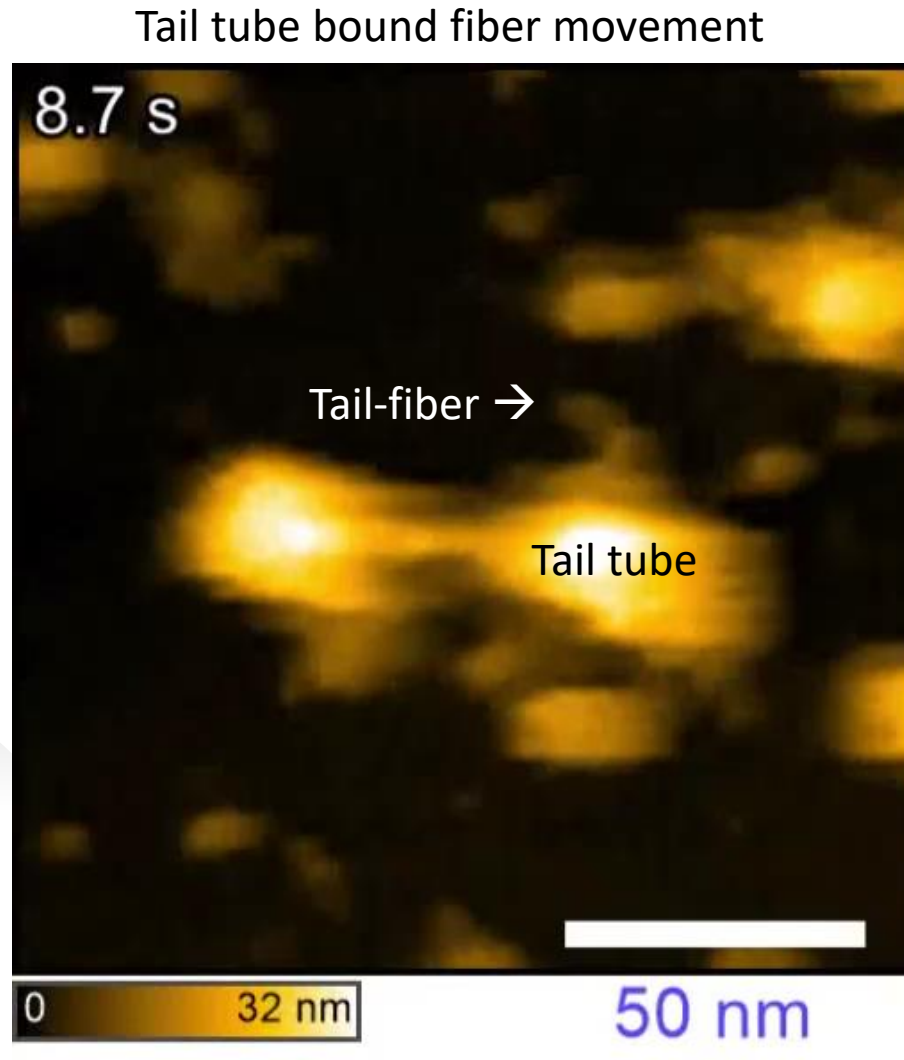
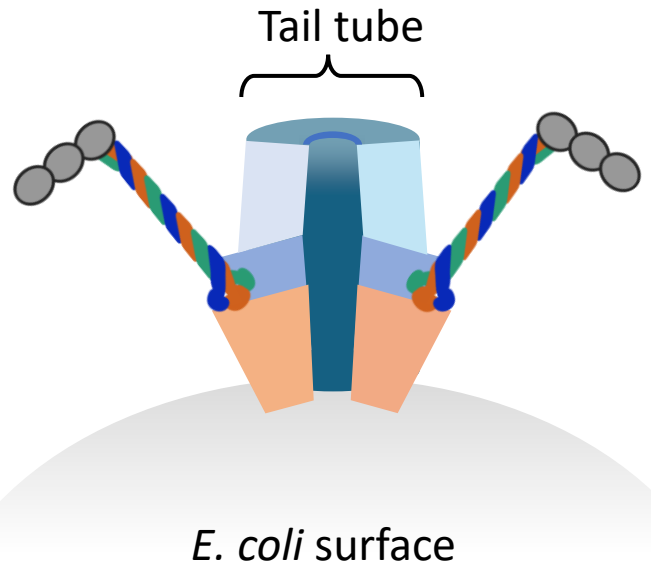
Fiber proximal section is a dynamic coiled-coil



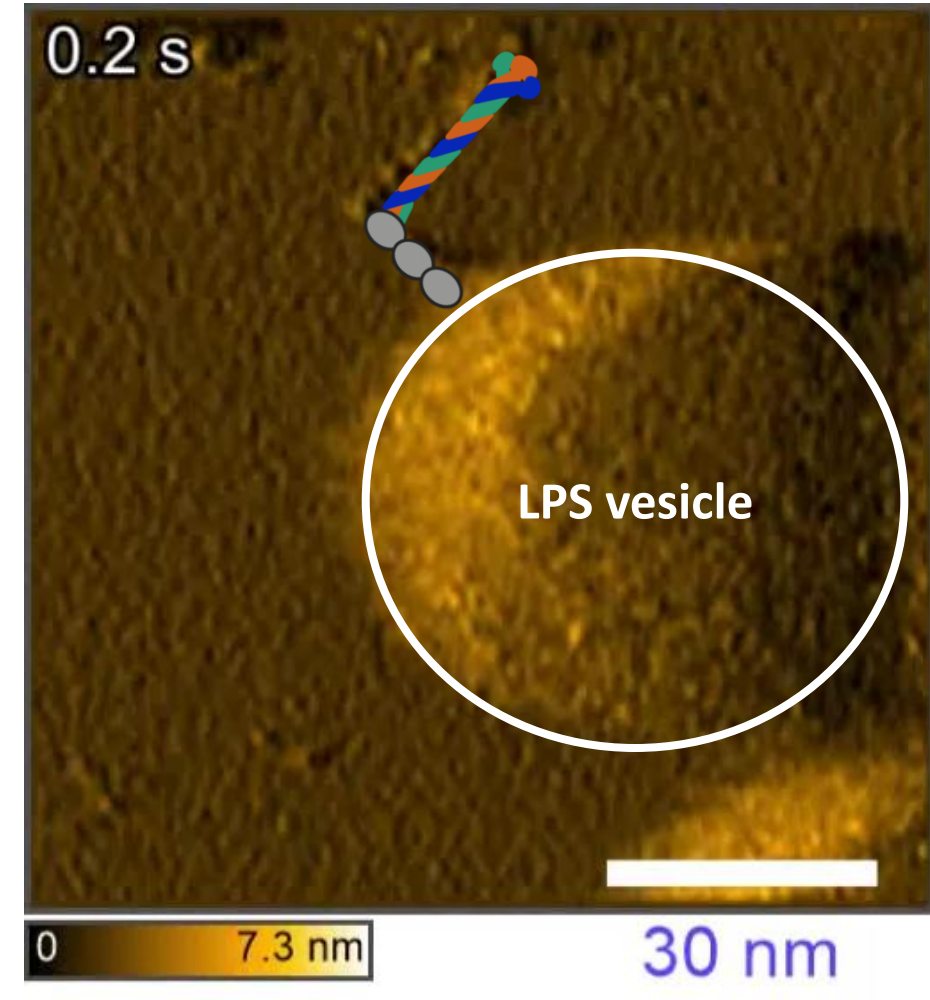
Fiber proximal section unwinding



Fiber binding and motion in more “native” states

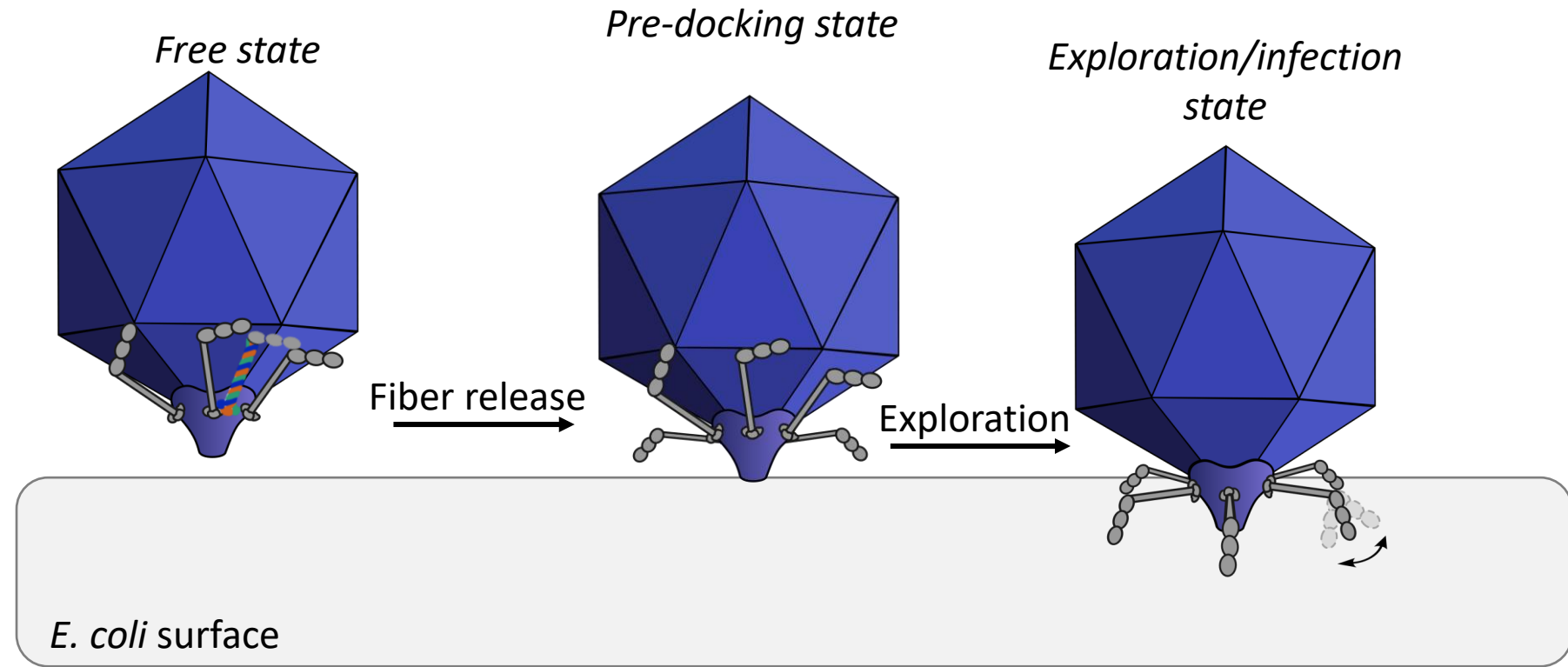


Fiber “jumping” on lipopolysaccharide vesicles

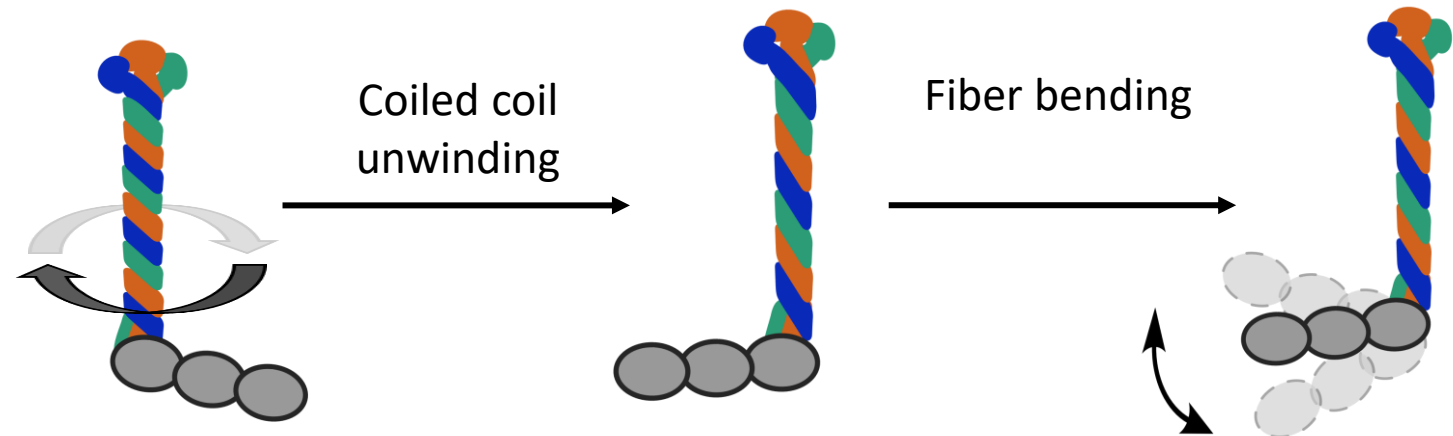


Role of fibers in target recognition and docking

Fiber bending:
Internal hinge region



Fiber rotation:
Dynamic coiled coil



Future plans...

- High-speed AFM imaging of whole tail fiber complexes
- Imaging fibers and tail-complexes on living bacteria
- Measuring virus-host interaction forces with optical tweezers

Special thanks to my students and colleagues:



Luca Kosik
PhD student



Miklós Cervenak
Student researcher



Dominik Sziklai
Postdoc
researcher

And funders:

