- Q1. I'm imaging soft polymer films, and over time the contrast gets worse-how can I tell if the tip is contaminated, and what precautions can I take during scanning to prevent this?
- Q2. After scanning biological samples in fluid, the tip seems degraded-how can I reduce the chance of biofouling or sample transfer to the tip?
- Q3. What scanning parameters can I tune to reduce the risk of tip contamination when imaging adhesive surfaces like hydrogels or organic thin films?
- Q4. How can I determine if a drop in resolution is due to tip wear vs tip contamination, and what are the cleaning or replacement protocols?
- Q5. Can certain material types or surface treatments increase the likelihood of contaminating the tip? How can I prepare my sample differently to avoid that?
- Q6. I'm working with live cell cultures, and the tip seems to get contaminated during extended imaging-how can I minimize biological debris sticking to the tip?
- Q7. After scanning hydrated tissue sections, I see ghost features and blurred edges-what kind of tip care or material compatibility checks should I perform to prevent this?
- Q8. How do I choose tip material and coating to reduce contamination when imaging sticky biomolecules like DNA or unfolded proteins?
- Q9. I-m imaging fibrous extracellular matrix proteins, and resolution decreases with each scan-how can I prevent material buildup on the tip during repeated passes?
- Q10. When imaging soft protein layers in fluid, I notice a rapid drop in resolution-could this be due to tip contamination, and how can I reduce fouling during long scans?
- Q11. I'm imaging a lipid bilayer in fluid and notice fuzzy edges after a few scans-how can I reduce molecular transfer to the tip?
- Q12. What role does relative humidity play in promoting tip contamination while scanning hygroscopic samples like polysaccharides?

- Q13. I frequently switch between dry and fluid imaging sessions-how should I clean or condition the tip between sessions to prevent cross-contamination?
- Q14. How can I detect early signs of organic contamination on the tip before they significantly affect resolution?
- Q15. I'm working with hydrophobic surfaces-how does tip hydrophilicity impact contamination risk, and how should I choose the tip coating?
- Q16. Are there best practices for storing AFM tips to reduce contamination risk before imaging sensitive protein samples?
- Q17. Can using a closed fluid cell reduce contamination while imaging weakly bound soft matter in aqueous conditions?
- Q18. What are the recommended low-force settings to minimize tip embedding and contamination in soft hydrogel networks?
- Q19. How do I distinguish between phase image artifacts from surface features and those due to tip contamination?
- Q20. Is there a recommended plasma or UV-ozone cleaning protocol for restoring contaminated tips used in biological samples?
- Q21. When imaging samples functionalized with charged ligands, how does electrostatic interaction contribute to tip fouling, and how can I mitigate it?
- Q22. What rinsing steps should be followed when transferring protein-coated mica samples to the AFM to reduce tip fouling?
- Q23. For imaging high-resolution DNA origami structures, how often should I check for tip contamination and recalibrate?
- Q24. I observe gradual contrast loss across sequential tiles in a stitched image-could progressive tip contamination be the cause, and how can I validate this?
- Q25. What's the safest way to clean a tip in-situ during a long imaging session without fully retracting it or disturbing the fluid environment?

- Q26. I'm imaging a hard metallic surface and I notice the tip is skipping over topography-what feedback loop settings or scan parameters should I modify?
- Q27. Why does my AFM lose contact with the sample while scanning a soft material in tapping mode, and how can I maintain good tracking without damaging the surface?
- Q28. I see flat lines in my topography image that suggest tracking failure-what environmental or scanner factors could be contributing to this, and how do I address them?
- Q29. How does the scan speed or line rate affect the ability of the tip to track rough features, and what's the best strategy for balancing speed and stability?
- Q30. When I zoom into nanostructures on my sample, the AFM fails to follow sharp edges-what tip and mode settings improve tracking on high-aspect-ratio features?
- Q31. While imaging soft live cells in tapping mode, I see sudden height drops and tracking loss-how can I tune feedback parameters to avoid indentation or disengagement?
- Q32. I observe periodic loss of tracking on collagen-coated tissue samples-could this be due to topographical steepness, and how should I adjust my scan parameters?
- Q33. When zooming into nanometer-scale protein aggregates, my AFM fails to follow sharp topography-what tip specs and scanning parameters can ensure accurate tracking?
- Q34. I-m imaging hydrated cells on soft substrates, and feedback seems unstable-how does substrate compliance affect tracking, and what imaging mode would be more stable?
- Q35. During imaging of biofilms, the feedback loop can-t keep up with surface changes-how can I improve the tracking without increasing damage to the biological sample?
- Q36. While scanning porous scaffolds, the feedback loop fails near void edges-how can I improve tracking over these abrupt depth changes?
- Q37. On rough mineral surfaces, I experience repeated line skip artifacts-how should I adjust Z-piezo gain and PID parameters?
- Q38. How can cantilever spring constant affect tracking stability when imaging stiff samples in

intermittent contact mode?

- Q39. What imaging mode is best suited for tracking sharp nanoparticle clusters on a soft biological matrix?Q5. I'm imaging curved bacterial surfaces in contact mode-how should I optimize scan size and tip approach to maintain tracking?
- Q40. What are signs of scanner creep leading to tracking issues during long-term imaging of patterned substrates?
- Q41. I notice tracking failure only during upward topography changes-what scan direction or phase settings should I revise?
- Q42. What precautions should I take when switching from low-viscosity to high-viscosity fluid imaging environments to prevent tracking instability?
- Q43. During high-magnification scans of fibrillar proteins, feedback becomes noisy-how can I enhance tip-sample coupling without increasing normal force?
- Q44. My sample is mounted at a slight tilt-how can uneven loading lead to lateral tip displacement and tracking defects?
- Q45. When imaging a cross-section of a multilayer film, the AFM fails to track abrupt material transitions-what scan angle or force setpoint strategies can help?
- Q46. What is the effect of scanner thermal drift on long, slow scans, and how can I compensate for it to maintain tracking?
- Q47. I'm scanning live cells under a perfusion system-how do flow-induced vibrations impact tracking, and how can I stabilize the setup?
- Q48. When scanning soft-aligned nanofibers, I see tracking loss along the fiber axis-should I change scan direction or engage mode?
- Q49. My sample shows anisotropic stiffness-how can I adapt feedback gains to compensate for lateral compliance variation across the field?
- Q50. When scanning a sample with periodic nano-patterns, I notice inconsistent tracking across similar features—could scanner resonance or feedback bandwidth be affecting performance, and how can I fine-tune the system to maintain consistent tracking?