

# Optimization on Scanning Ion Conductance Microscopy

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- Fundamental Principle of SICM
- Fast SICM Model & Applied Techniques\*
- Ultra-high Resolution on HSQ Imaging\*
- Future Work



# Introduction

## SICM System Sketch

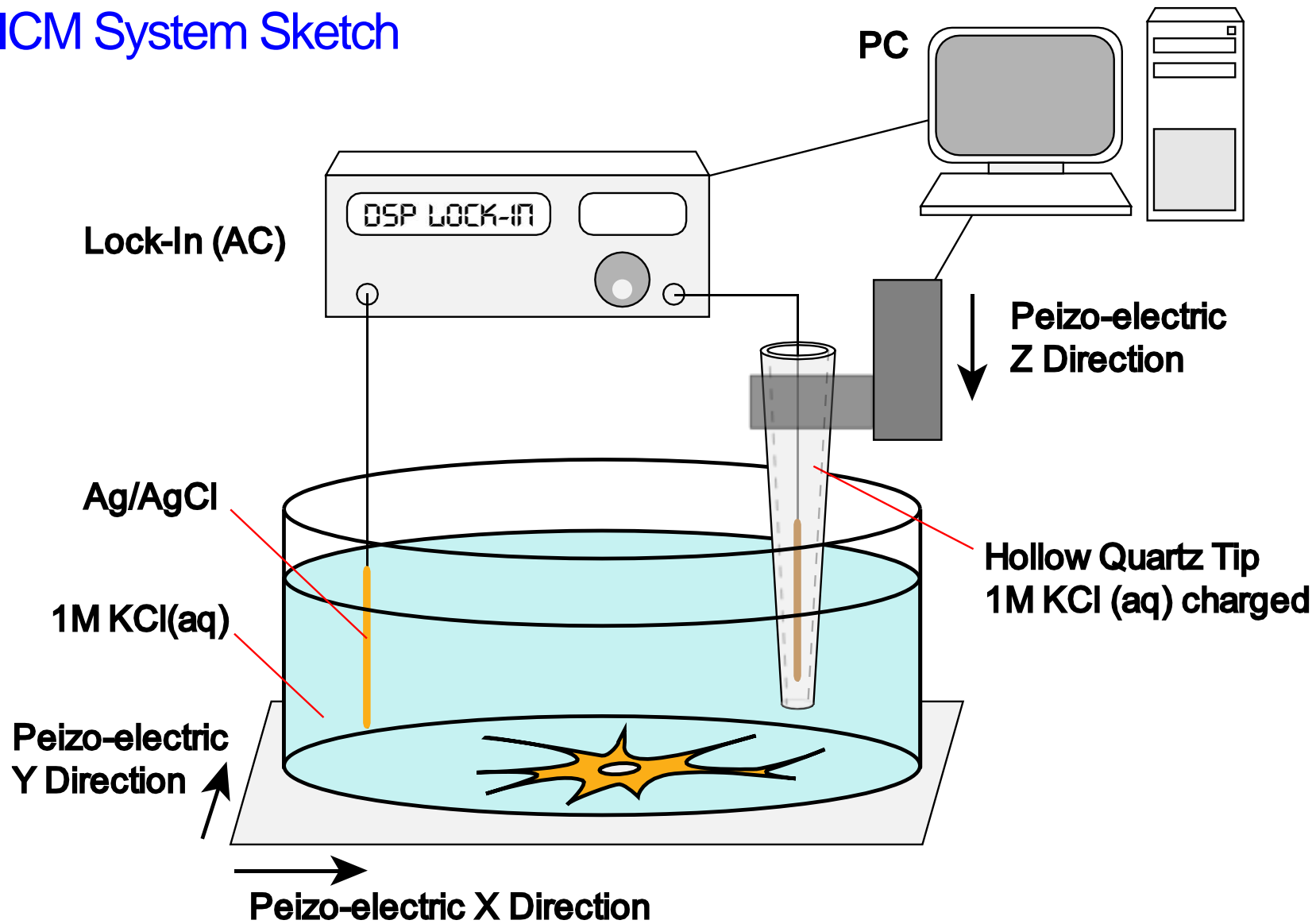


Fig. 1-1. SICM Simple Sketch (Digital device not shown)

# Introduction

## SICM Principle

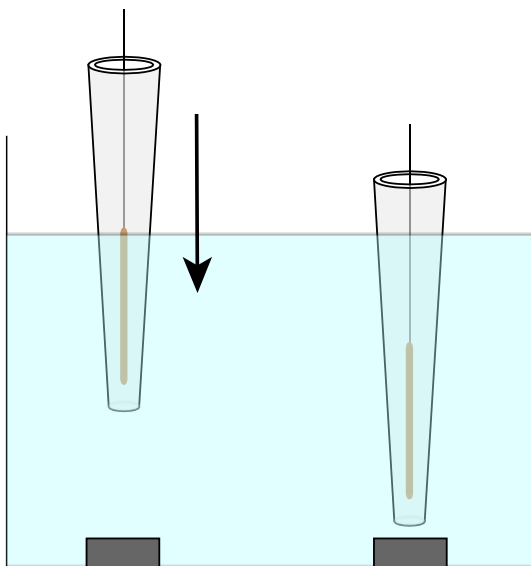


Fig. 1-2. Downward Motion

## Sudden Back

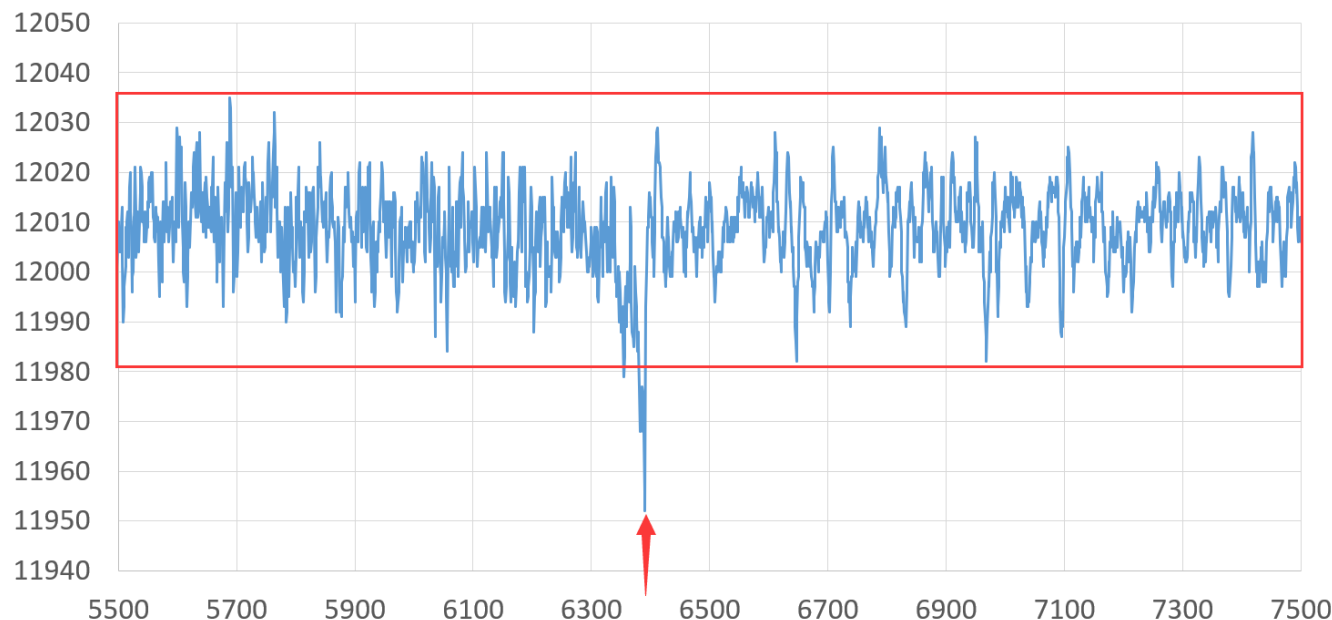


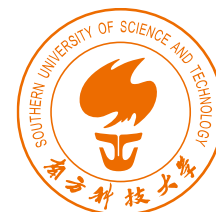
Fig. 1-3. Current Decline Feedback

General displacement current decline expression:

$$I(d) = \frac{V_0}{R_p + R_a^{\frac{r_o}{r_i}}(d)} = \frac{R_p + R_a^{\frac{r_o}{r_i}}(d \rightarrow \infty)}{R_p + R_a^{\frac{r_o}{r_i}}(d)} I_0 = \frac{R_p + R_a^{\frac{r_o}{r_i}}(d \rightarrow \infty)}{R_p + R_a^{\frac{r_o}{r_i}}(d)} \times \frac{\sigma V_0 r_i}{\frac{1}{\pi \tan \alpha} + \frac{1}{4}}$$

Where,  $R_p$  is the resistance at inner side of probe, and  $R_a$  is the outside resistance,  $r_o$  is the outer radius of tip,  $r_i$  is the inner radius,  $\alpha$  is the angle of pipet tip.

Johannes Rheinlaender, Tilman E. Shaffer. *Anal. Chem.* **2017**. 89. 11875-11880



# Introduction

## SICM Control Subsystem

**Servo Off** for analog control, no micro-adjustment occurs to raise accuracy and speed.

FPGA is **real-time** and can generate a voltage **-10V ~ 10V** interval with 65536 amplitude resolution range.

CPU is a **preemptive** processor, which can be interrupted by multitasking traps.  
Not appropriate for real-time control.

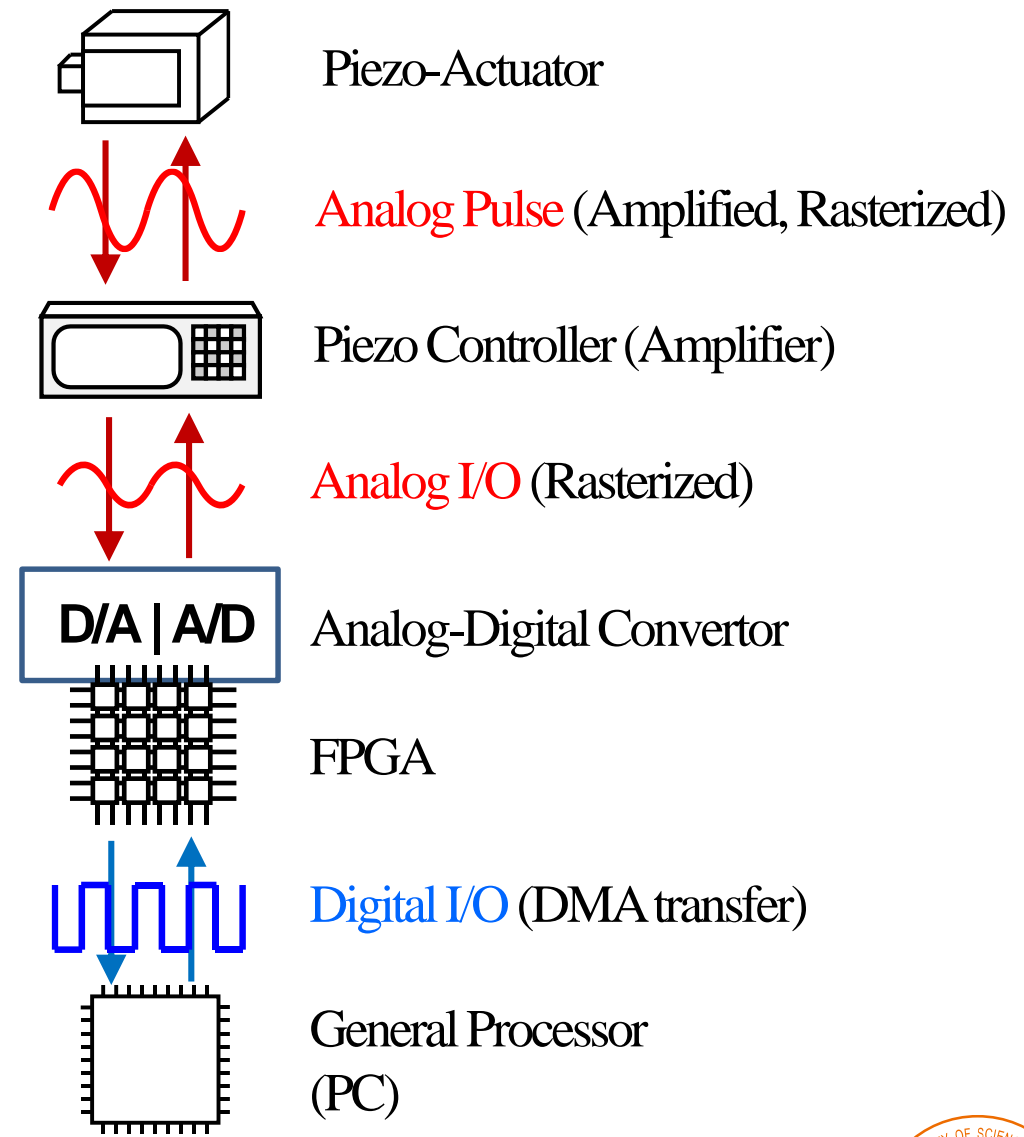
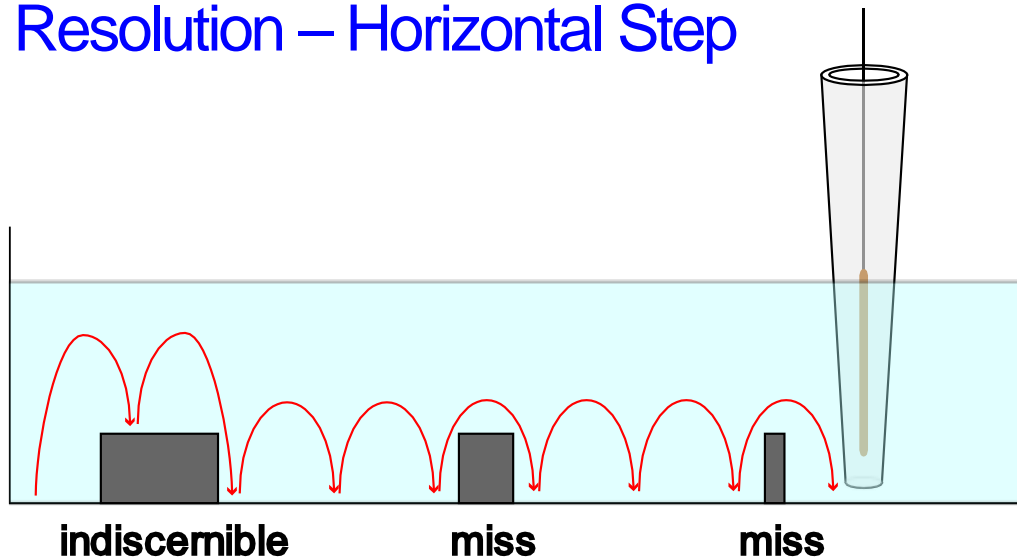


Fig. 1-4. Control Subsystem



# Introduction

## Resolution – Horizontal Step



1 indiscernible and 2 missed.  
**Poor** x-y resolution

Fig. 1-5. Hopping Model with Large x-y Step

2 discernible and 1 indiscernible.  
**Relative good** x-y resolution

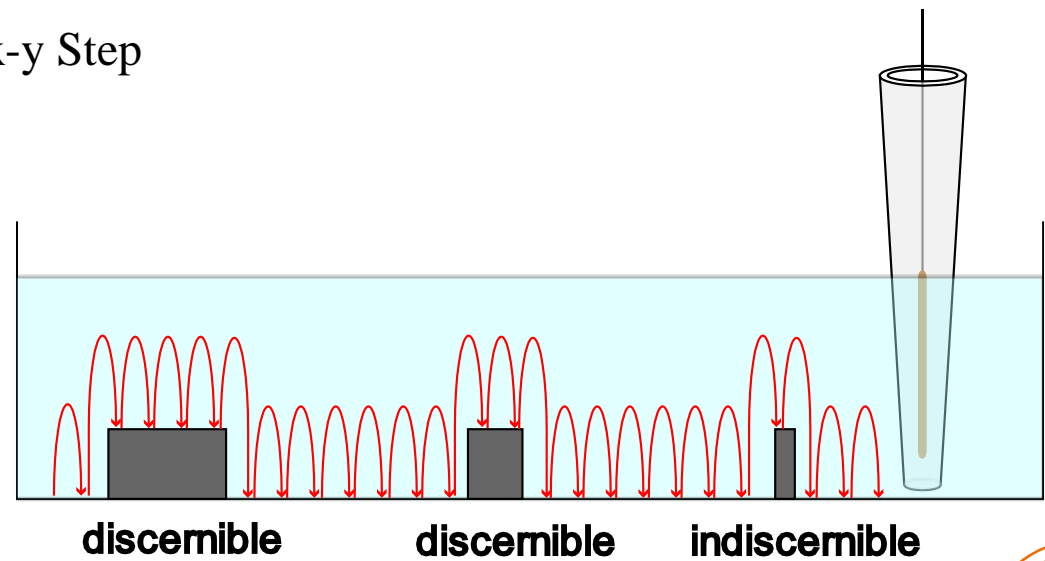


Fig. 1-6. Hopping Model with Small x-y Step



# Introduction

## Resolution – Tip Aperture

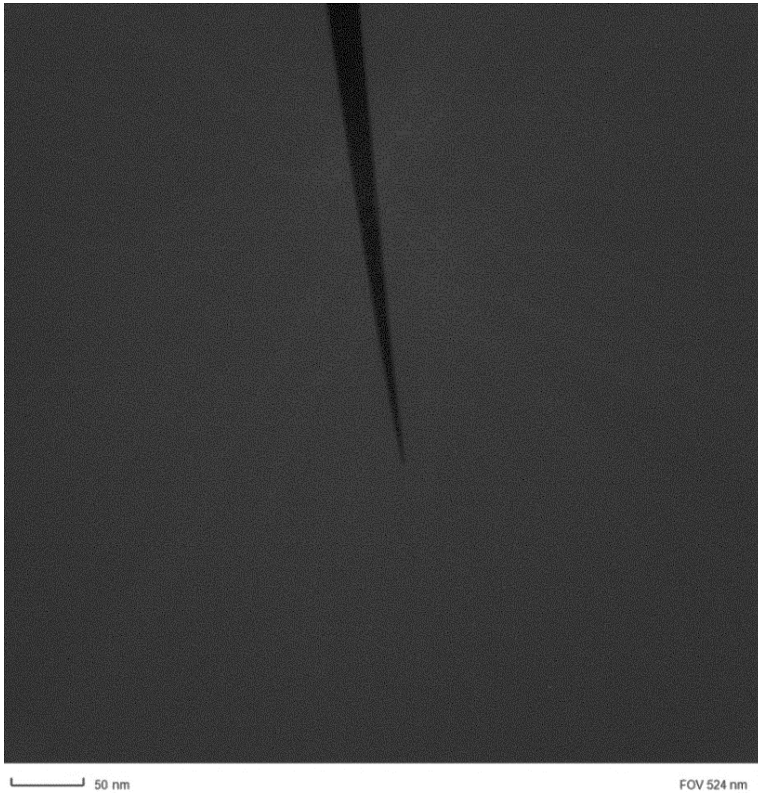
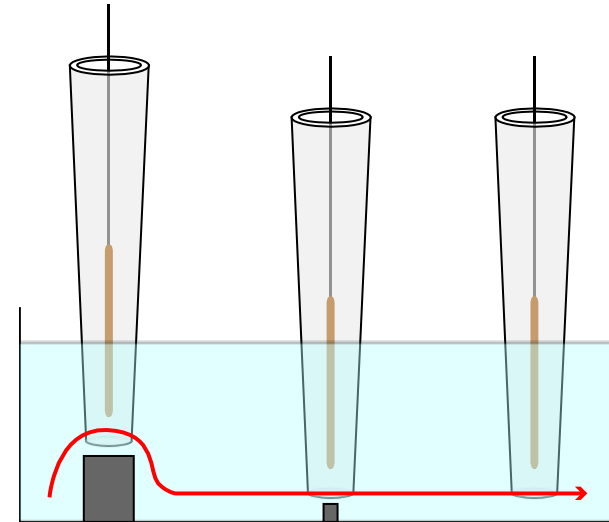
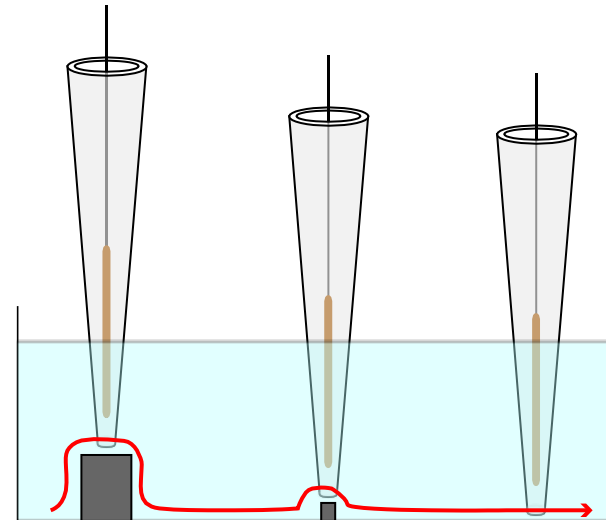


Fig. 1-7.  $r_i \leq 5 \text{ nm}$  Tip under TEM



**discernible    indiscernible    reference**

Fig. 1-8. Tip with Large Aperture



**discernible    discernible    reference**

Fig. 1-9. Tip with Small Aperture



# Fast Hopping Model

## Result

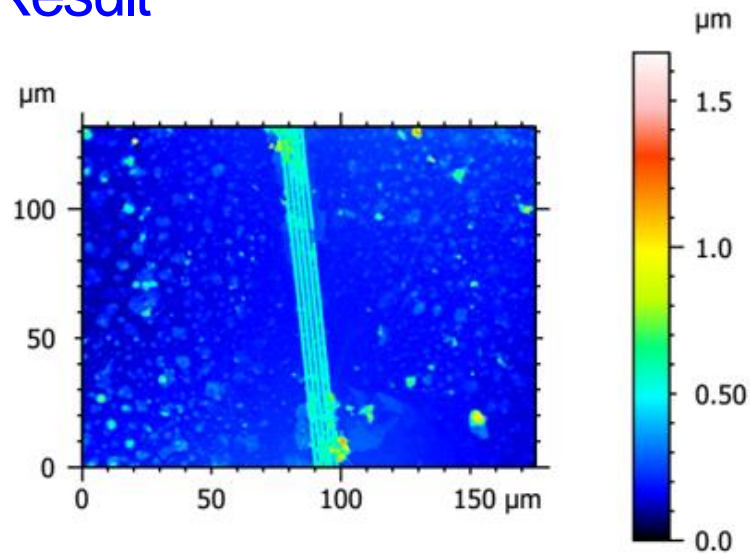


Fig. 2-1. ITO Confocal Microscopy Imaging



Fig. 2-2. Quartz Tip TEM Size Scaling

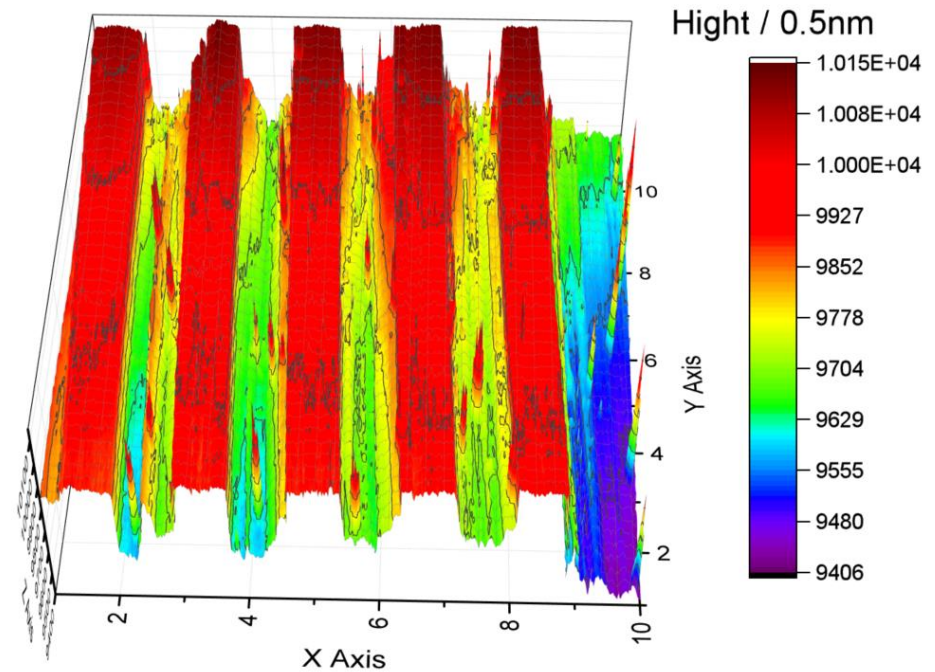


Fig. 2-3. SICM Imaging

Area:  $10\mu\text{m} \times 10\mu\text{m}$  region,  $500 \times 500$  steps

Time Elapsed: 124 min = 2.07 hr (29.8 ms / point)

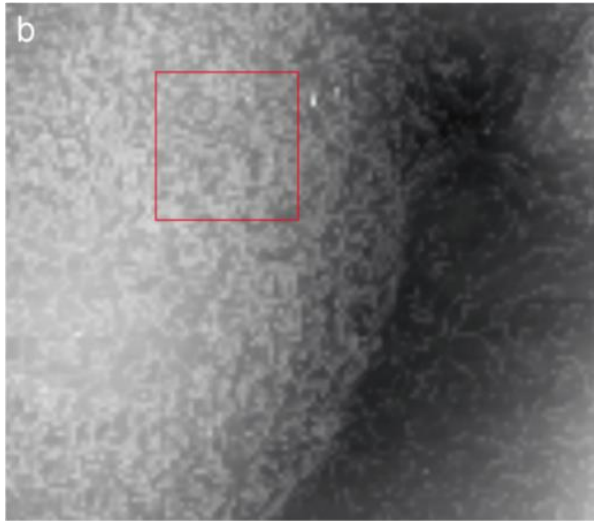
Probe Aperture: 15 nm





# Fast Hopping Model

## Comparison



Area: **40 $\mu$ m  $\times$  40 $\mu$ m region, 128  $\times$  128 steps**

Time Elapsed: **25 min (91.6 ms / point)**

Probe Aperture: **80 nm**

Fig. 2-4. A6 Kidney Epithelial Cell SICM Hopping.

hopping mode and fast SICM. Due to the huge number of points being scanned it is possible to resolve features in fast SICM that are not possible to resolve in the hopping mode in a reasonable time scale for a dynamic, live cell surface. A hopping mode image obtained to the highest resolution possible with the hopping mode software, 512  $\times$  512 pixels would take around 5 h to complete compared to around 10 min for the 1024  $\times$  600 pixel image in fast SICM.

David Klenerman, Yuri Korchev. *Ultramicroscopy*. 2012. 10. 1016-1023



# Fast Hopping Model

## Strategy – Nonlinear Scanning

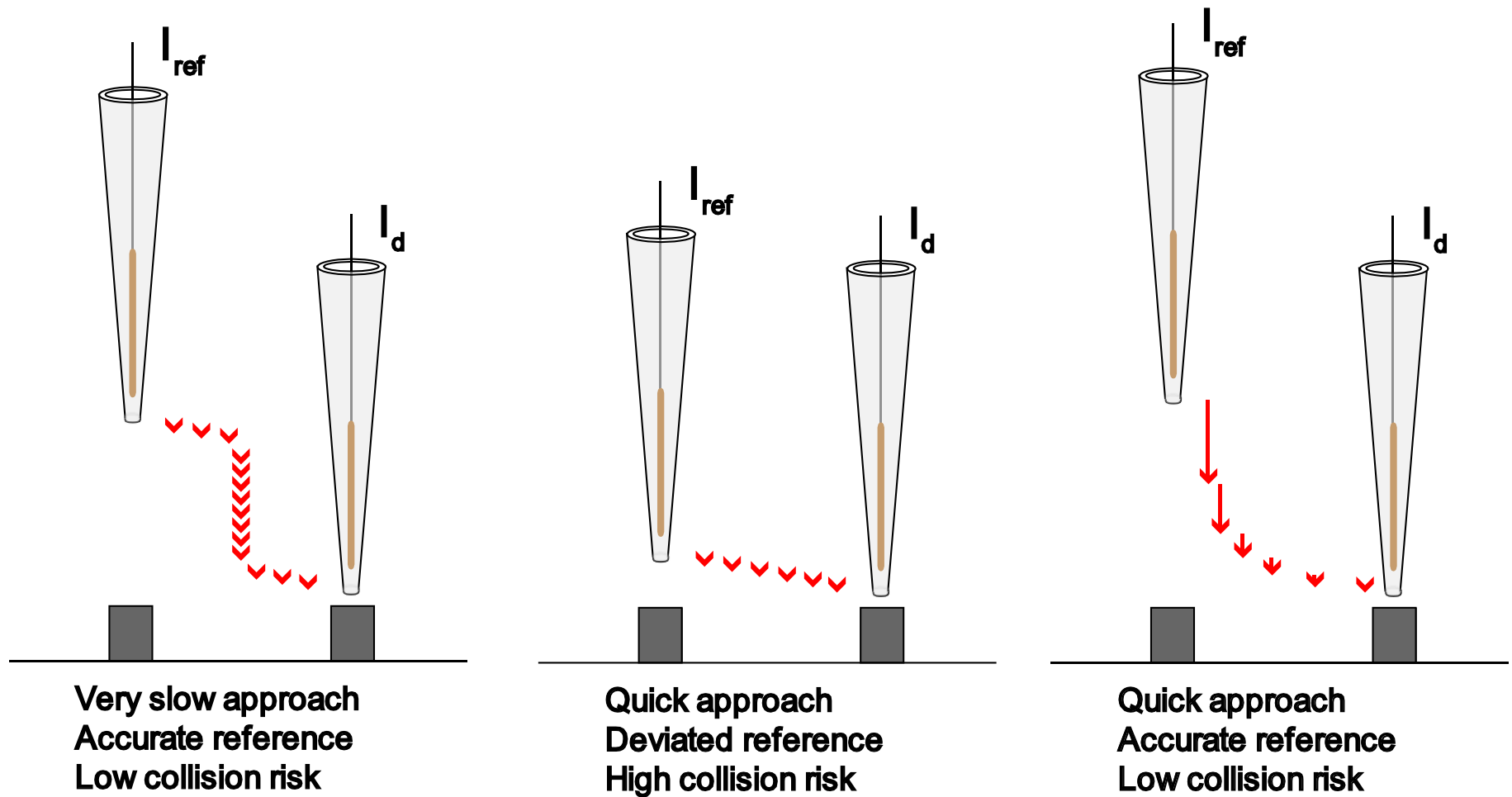


Fig. 2-5. Three Strategies of Single Point Scanning



# Fast Hopping Model

## Strategy – Region Classification

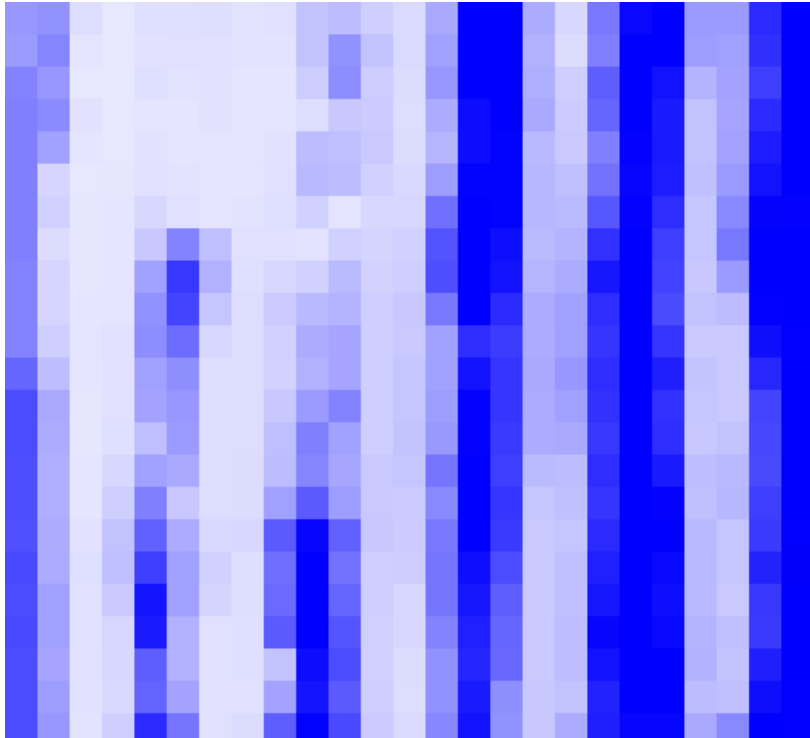


Fig. 2-6. Rough ITO Scanning

$25 \times 25$  steps in  $12 \times 12 \mu\text{m}$  square

Mask Matrix

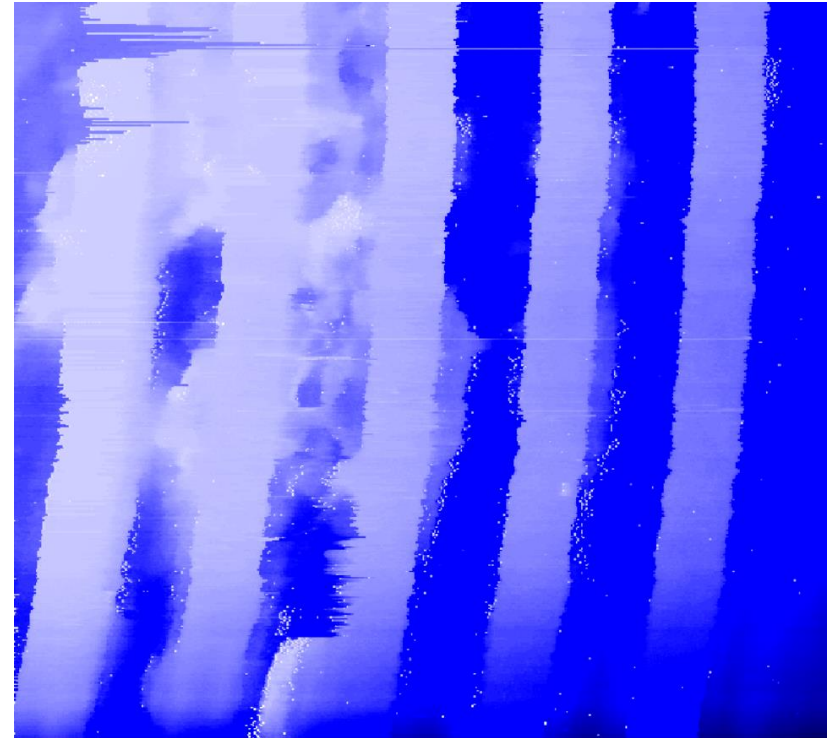


Fig. 2-7. Precise ITO Scanning

$500 \times 500$  steps in  $12 \times 12 \mu\text{m}$  square



# Fast Hopping Model

## Strategy – Dynamic DMA FIFO

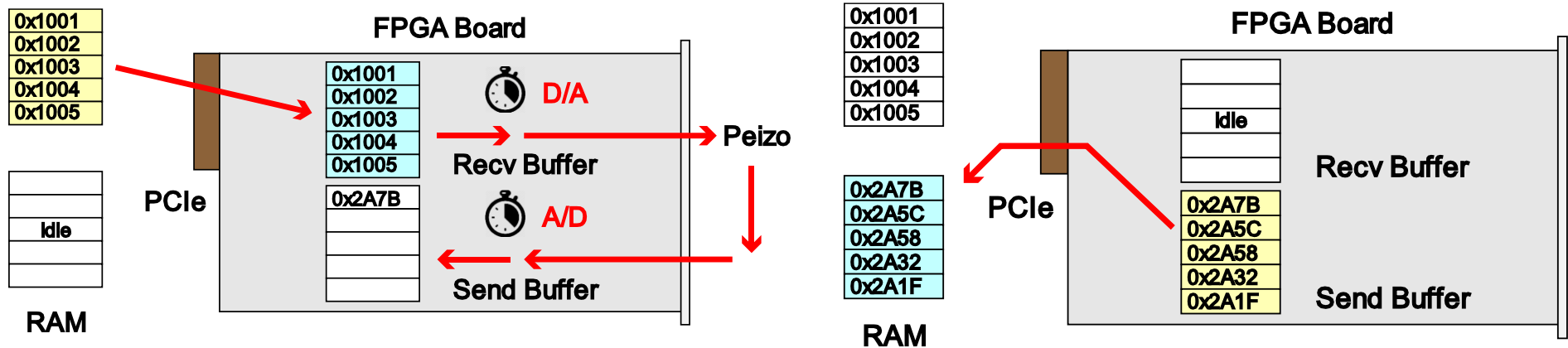


Fig. 2-8. Data Batch Processing and Transmission (Low Efficiency)

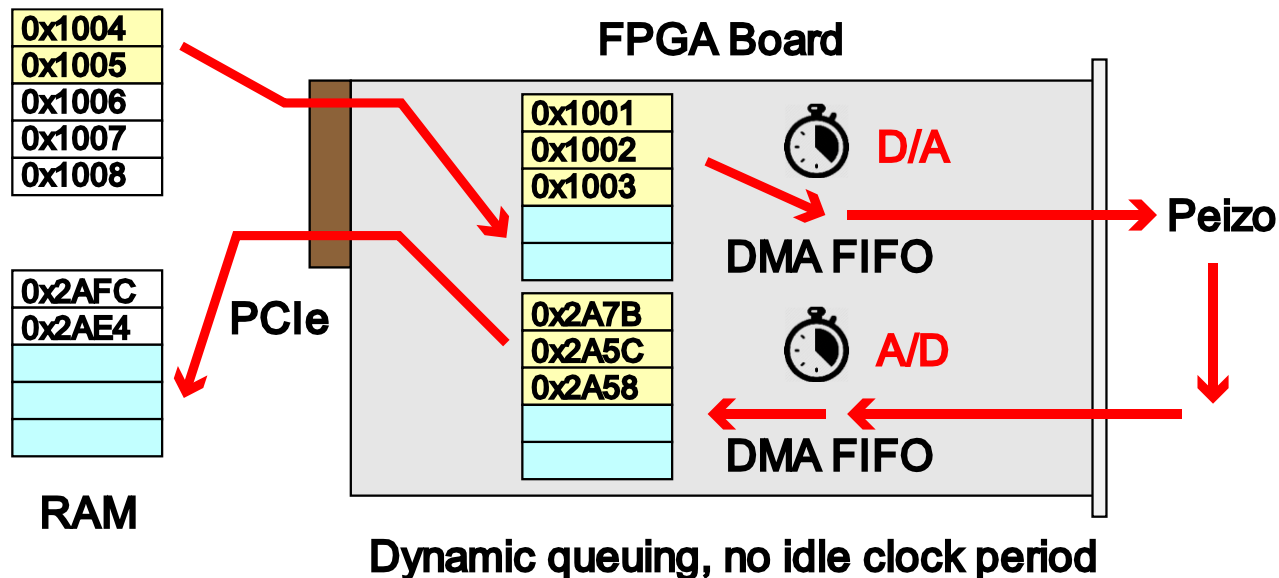


Fig. 2-9. Dynamic DMA FIFO (High Resources Occupying Efficiency)



# Ultra-high Resolution Imaging

## Result

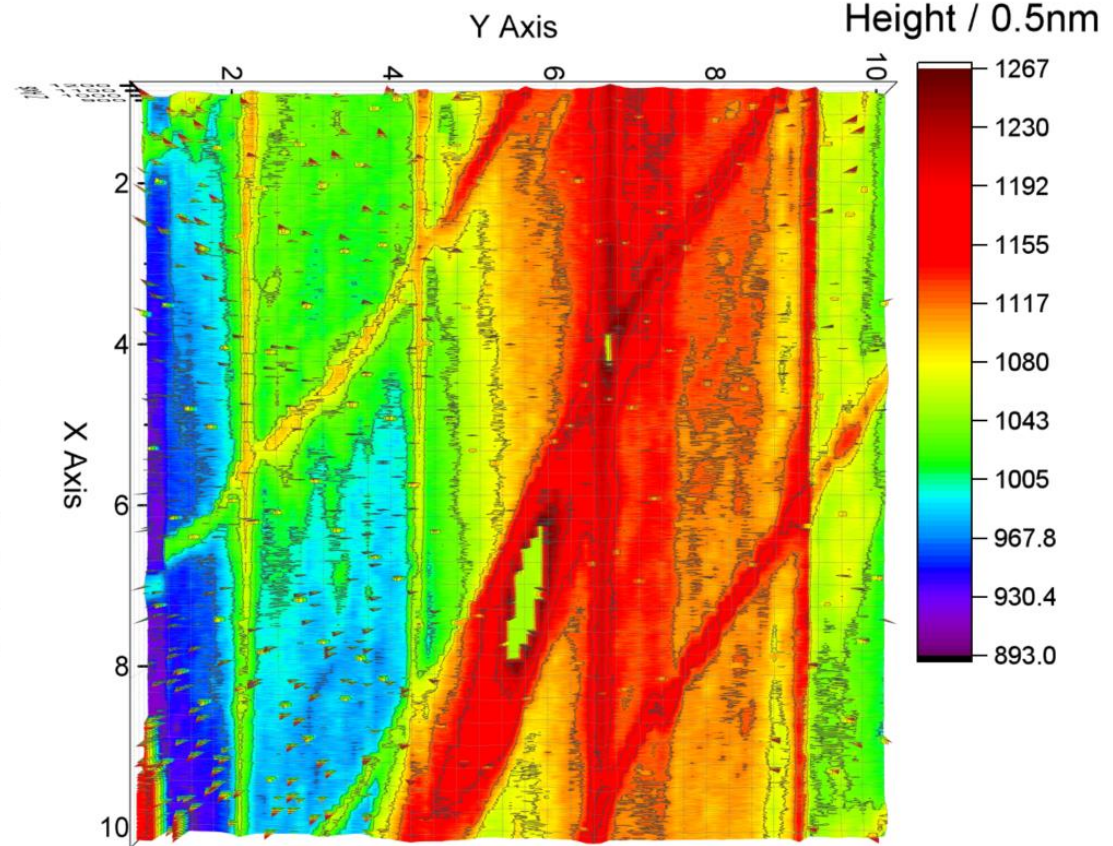
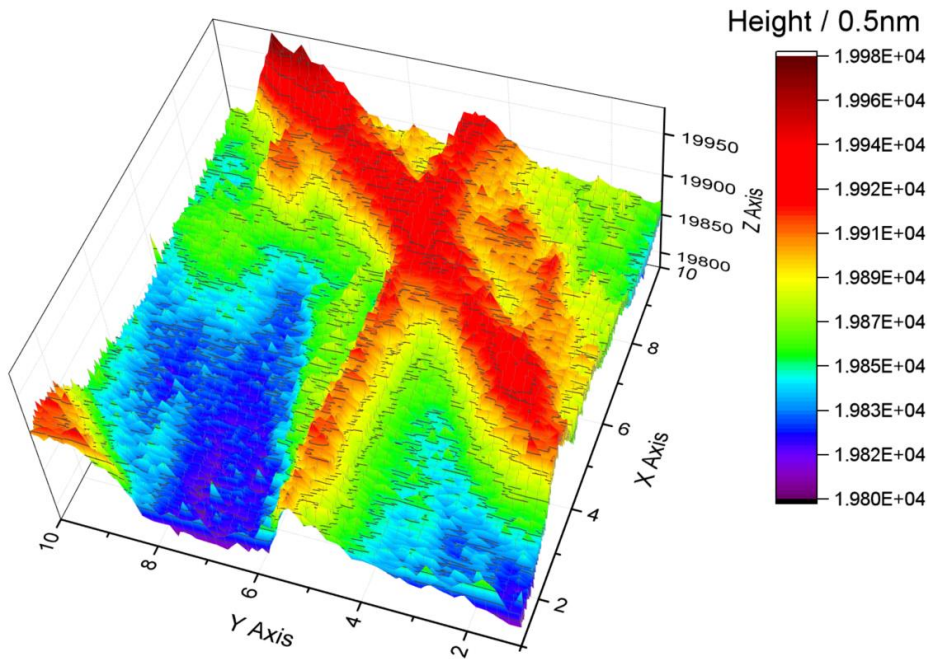


Fig. 3-1 and 3-2. Inclined-crossed HSQ Nanobeam

3-1.

Area: **300nm × 90nm region, 600 × 45 steps**

Time Elapsed: **30.15min (67 ms / point)**

Discernibility: **20nm width, 14nm height beam.**

Probe Resolution: **Below 10nm.**

3-2.

Area: **1.65μm × 1.5μm region, 1100 × 100 steps**

Time Elapsed: **2.75 hr (90 ms / point)**



# Ultra-high Resolution Imaging

## Result

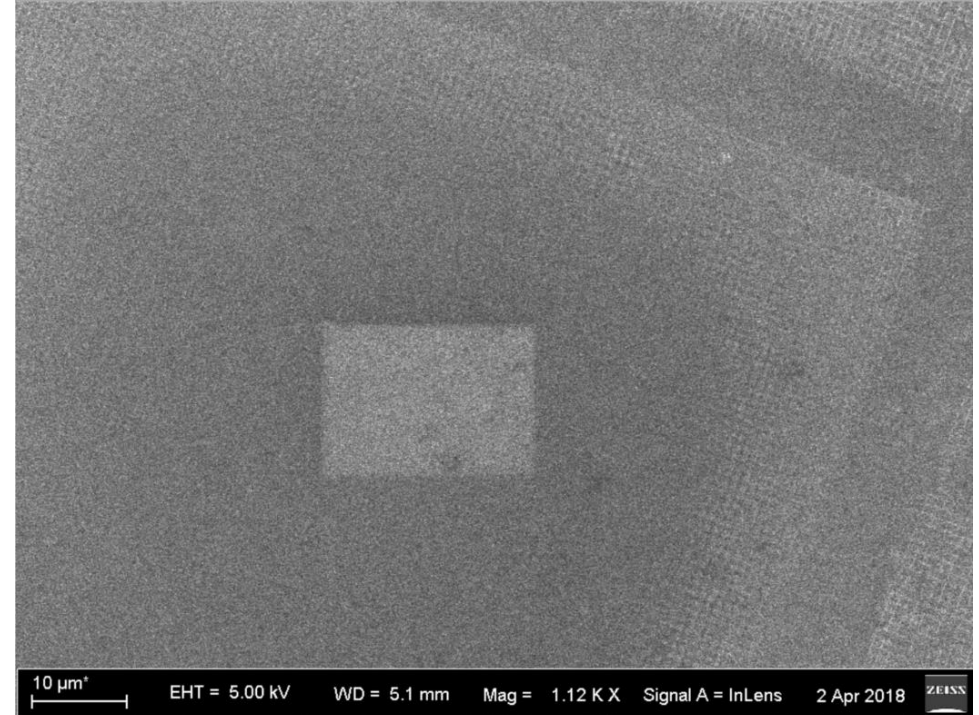
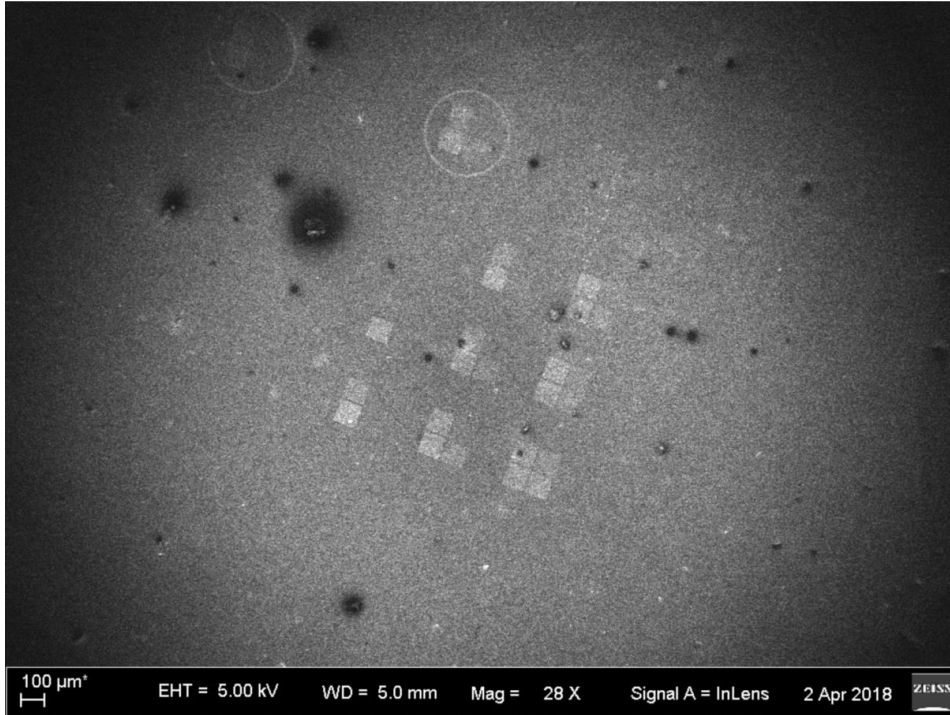


Fig. 3-3 and 3-4. SEM Images of HSQ Beams

**It is a pity that this became a pending case till now!**



# Future Work

## Ion Channel Scanning

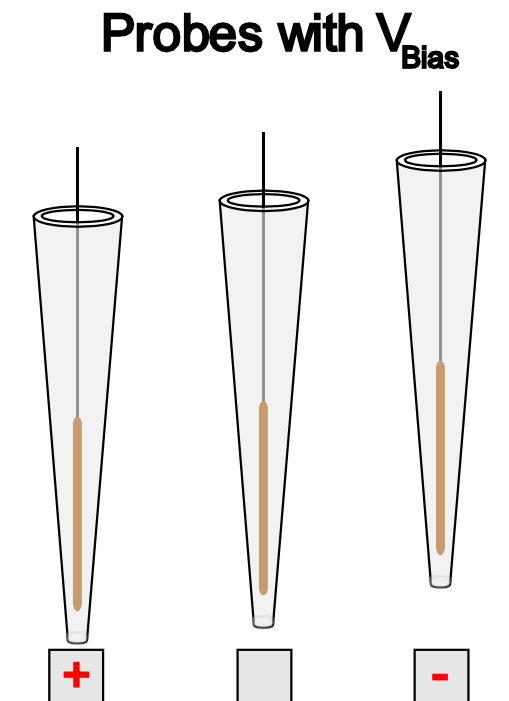


Fig. 4-1. Ion Current Rectification of SICM Scanning Charged Sample

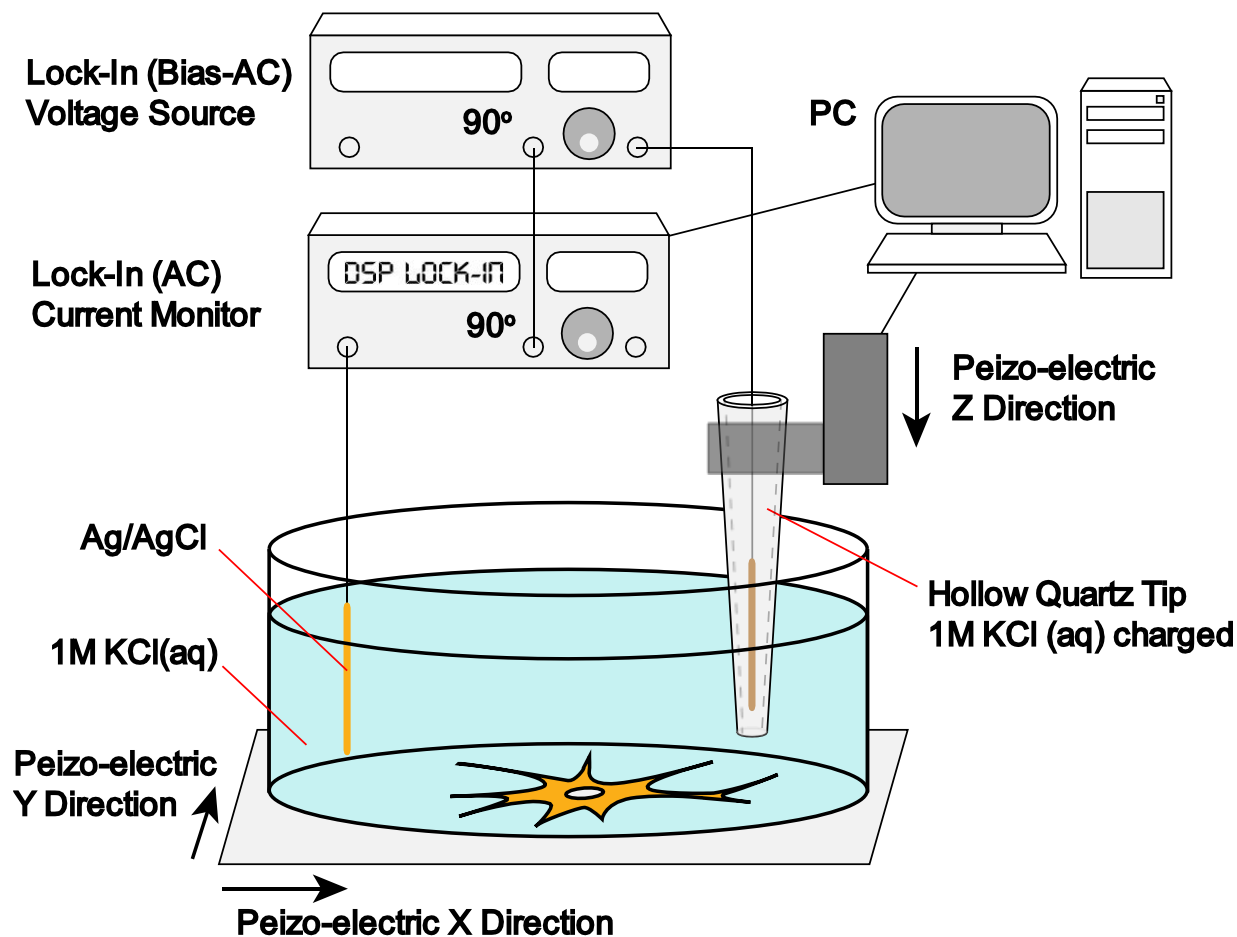


Fig. 4-2. Double Lock-In System Making a Sensitive Bias Tip

David Perry, Rehab AI Botros, Dmitry Momotenko, Sophie L. Kinnear, and Patrick R. Unwin.  
*ACS. Nano.* **2015.** 10. 1021-1032.



# Future Work

## Ion Channel Scanning

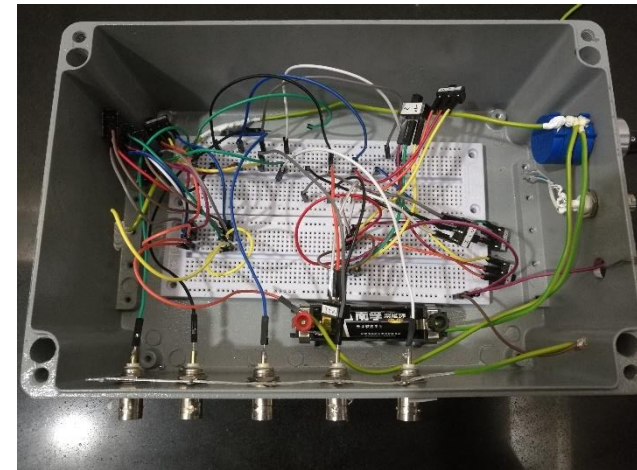
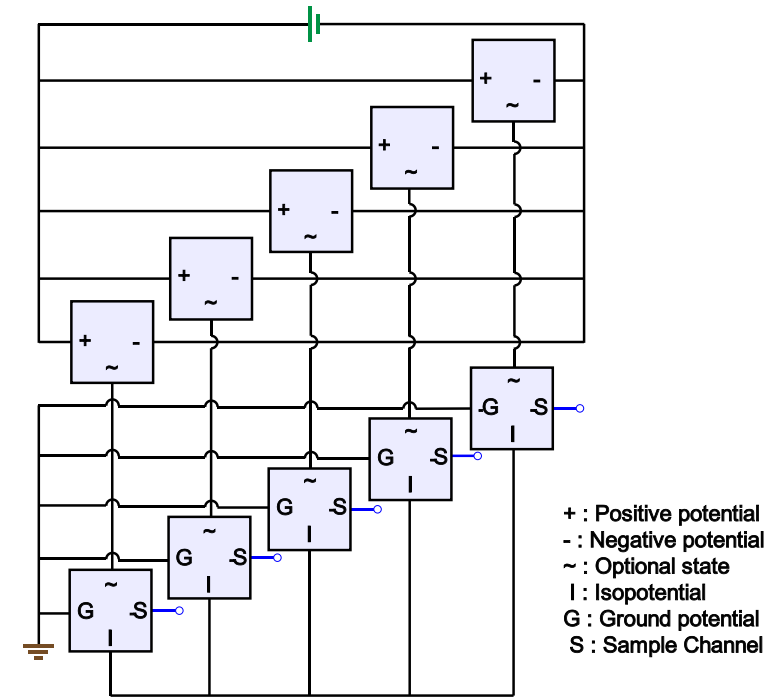
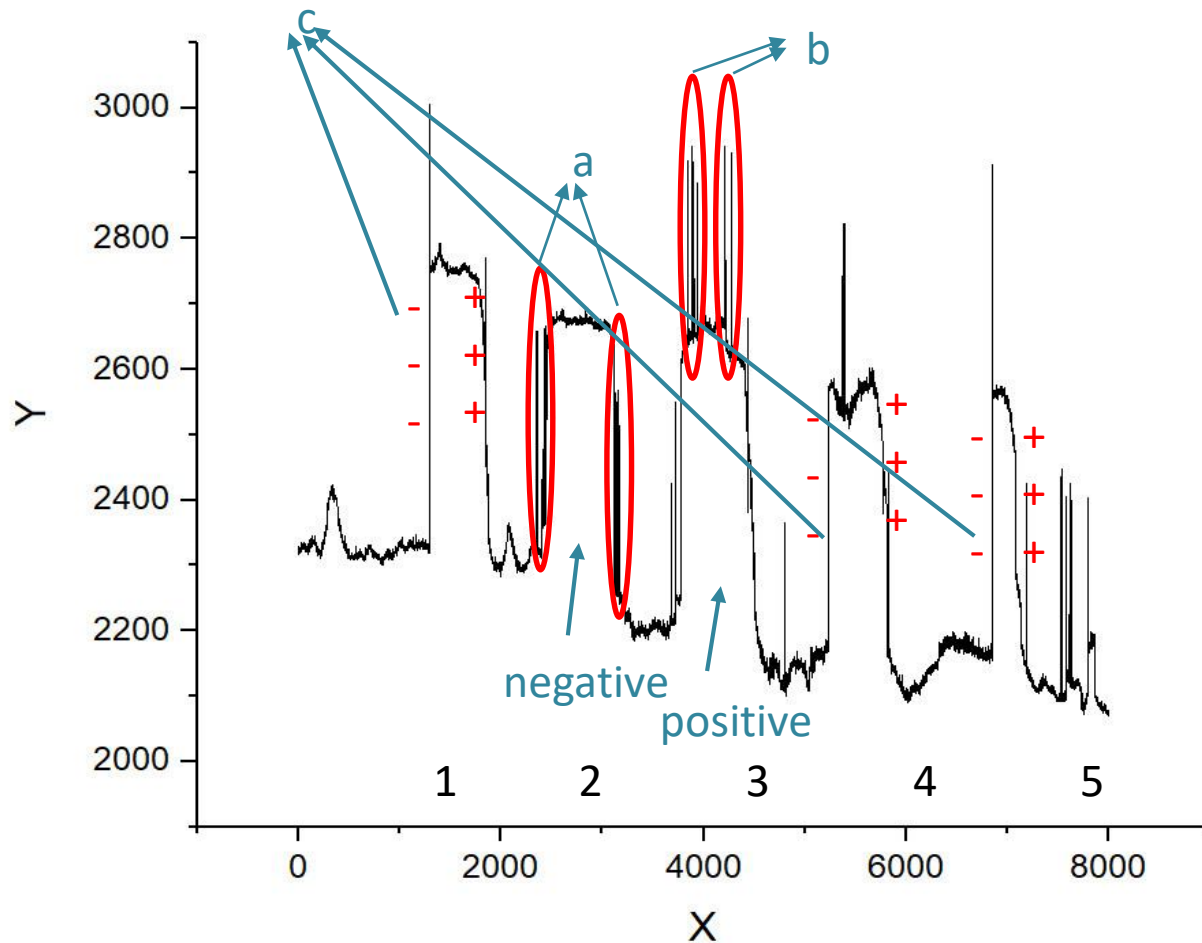


Fig. 4-3. Impact of Charged Sample on SICM Tip



# Future Work

## Ion Channel Scanning

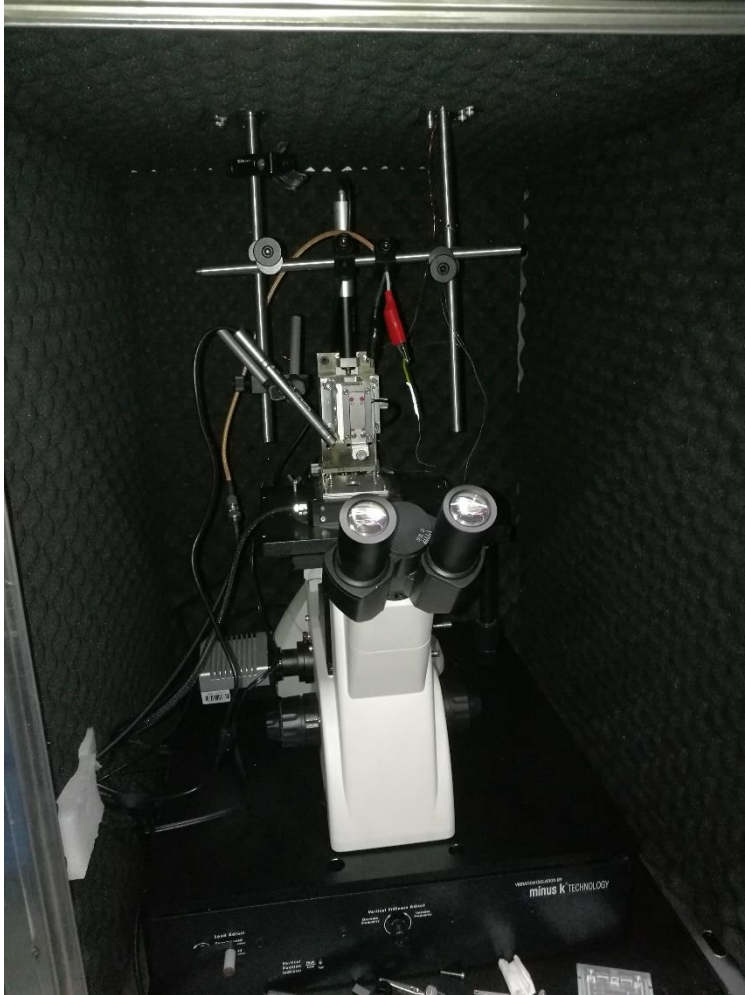
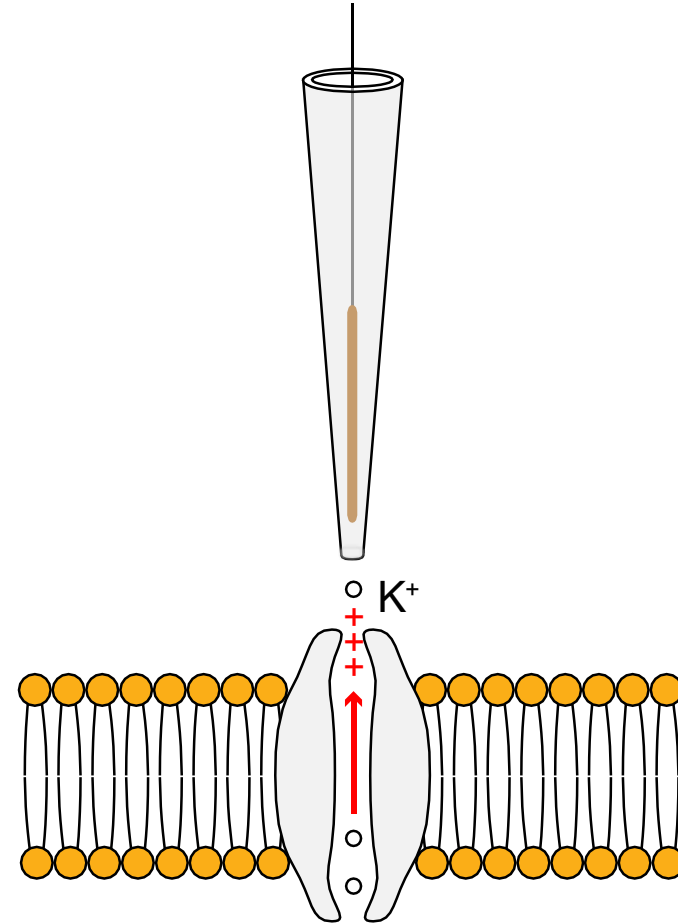


Fig. 4-4. SICM for Cell Imaging



# References

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THANKS FOR LISTENING!