# Single-Molecule Recognition and Manipulation Studied by Scanning Probe Microscopy

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**MTAN-inhibitor** 

# **Scanning Probe Microscope (SPM) : Basic aspects** $\chi +$ χ- $\forall +$ motion even-COOH vs. even-COOH tip surface **m1**( atmosphere **m2** <del>\$</del>

# Scanning



•Principle of piezo element. The applied voltage makes the element longer or shorter.



• The combination of three piezo elements makes it possible to move the STM tip in the X-, Y-, and Zdirections.



•In most modern scanning probe microscopes, one uses a tube geometry. .

### **Reciprocal Space vs Real Space**

Low Energy Electron Diffraction

•Scanning Probe Microscopy

### **Periodic system**



VC(100)



Local structures without periodicity can be understood only by scanning probe microscope.

# Image: Construction of the second second

Instrument	depth resolution	lateral resolution
Mechanical Profiler	0.5 nm	0.1-25 μm
Optical Profiler	0.1 nm	0.35 - 9 μm
Atomic force microscope (AFM)	0.01nm	0.1nm
Scanning tunneling microscope (STM)	0.001nm	0.1nm

# Resolution

# **Scanning Probe Microscope (SPM)**

# General experimental approach-

Monitor a physical quantity which is dependent on separation between a probe tip and a sample surface.  $\rightarrow$  Use Current (I) and Force (F)

- Scanning Tunneling Microscope (STM)
- Atomic Force Microscope (SFM)

# SPM techniques provide unique information-

Local - probes topographic and electronic structure on an atomic scale Real space - information is collected through a direct measurement

# STM-monitors a tunneling current. I $\propto$ Ue<sup>-(kd $\sqrt{\Phi}$ )</sup>

I-tunneling current U-sample bias k -transmission coefficient d -tip sample separation Φ-average work function

- conductive sample
- tunneling current : pA~nA
- lateral resolution : 0.1nm
- depth resolution : 0. 001nm

# **PRINCIPLE of STM**









# STM Tip









PVBA/Pd(111)



## Molecular Orientation and Chiral Recognition of a Single Chiral Molecule



### Adsorption Preference of PVBA on Pd(111)

(Submitted to PRL, July 2003)





### Model Potential V(r) between a Carbon Atom and fcc(111)

V(r) = 0 at a top-site, V(r) = 1 at a hollow site (Fourier Expansion of Surface Reciprocal Lattice)

V(**r**) = 2 / 3 - 2 / 9  $\sum_{n=0}^{2} \cos \omega_n \cdot k\mathbf{r}$ **r** : position coordinate,  $\omega_0 = (0,1)$ ,  $\omega_1 = (-\sqrt{3}/2, -1/2)$ ,  $\omega_2 = (\sqrt{3}/2, -1/2)$ ,  $k = 4\pi/\sqrt{3}a$ .

### **Binding Energy**

Summation of each potential value for all atoms of two rings as a function of angle for each chirality( $\chi$ )



### **Orientation and Chiral Recognition of PVBA on fcc(111)**



# AFM-Monitors interfacial forces $F \propto (1/d)^{x}$

F-interfacial force d-tip sample separation x-power law of acting force

Interfacial forces include:

repulsive forces (contact AFM) van der Waals forces (noncontact AFM) electrostatic forces (EFM) magnetic forces (MFM) chemical forces (CFM)

- interfacial forces : pN -nN
- lateral resolution : 0.1-10nm
- depth resolution : 0.01 0.1nm

# **Atomic Force Microscopy (AFM)**



AFM has made it possible to view living biomolecules with atomic resolution in physiological environment

# **AFM Tip and Cantilever**

### •Tip radius : 4 nm





# Surface Forces on Switchable Bioactive Surfaces Studied by Interfacial Force Microscope



**Probing of Biomolecular Interaction** 

Thermally-Activated Surfaces: Tethered PNIPAM Films





# Force-Distance Curve





# Single Molecular Force Spectroscopy Enzyme-inhibitor interaction

### **Bomb Fishing**



Hook and Bait Fishing



# 5'-methylthioadenosine/S-adenosylhomocysteine nucleosidase (MTAN)



S-Adenosylhomocysteine (SAH)





MTAN complexed with an inhibitor MT-ImmA

Homocystinyl Immucillin A (HIA)

This enzyme-inhibitor pair is an important one, owing to its potential for antibiotic development.

# **MTAN-inhibitor interaction**



→ Epoxy group forms a covelanet bonding with -NH2 group

Single molecular interaction (enzyme-inhibitor)

# Force Induced Dissociation : Evans & Ritchie (1997)



→ Statistical analysis enables to probe intermediate states between an enzyme and an inhibitor for a new drug design

# Temperature Controlled Reversible Switching in Tethered pNIPAM films



# **PNIPAM+Microhotplate = Reversible Protein Trap**

PNIPAM coating on \_ silicon nitride membrane (200µm wide)



Hot plate can be programmed to heat PNIPAM above transition

Heating promotes protein adsorption. Cooling promotes protein desorption



T = 0.0 sec

T = 0.8 sec

T = 1.2 sec

# Variable Temperature Studies Reveal Sharp Transition for PNIPAM

-ODTS Coated Tip



**Behavior of PNIPAM under water** 

- All surfaces experience long-range repulsion → consistent with "anti fouling" behavior
- 2) The repulsion collapses at a sharp transition temperature
- 3) Above the transition, PNIPAM surface become sticky after contact is made, consistent with protein adsorption.

# **Tip-PNIPAM Interactions**

Below Transition Temperature



Above Transition Temperature



Chain Hydration(2 nm thick)

- 1) Promotes swelling
- 2) Inhibits adhesion

**Disruption of Hydration Layers** 

- 1) Allows collapse
- 2) Promotes adhesion

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