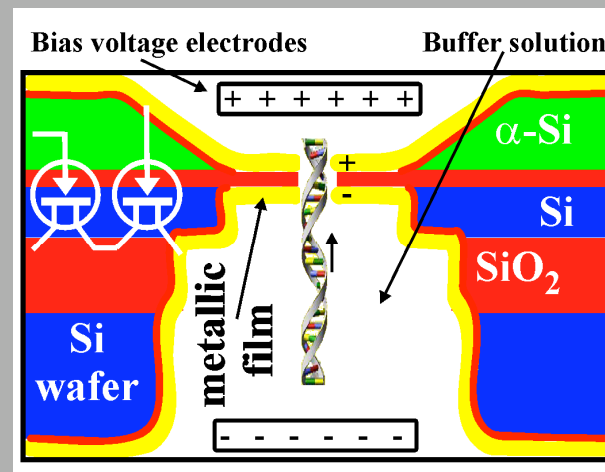


# Semiconductor Nano-Pores Tunability for DNA Sequencing\*

Jean-Pierre Leburton,

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University of Illinois at Urbana-Champaign, Urbana, IL 61801, USA

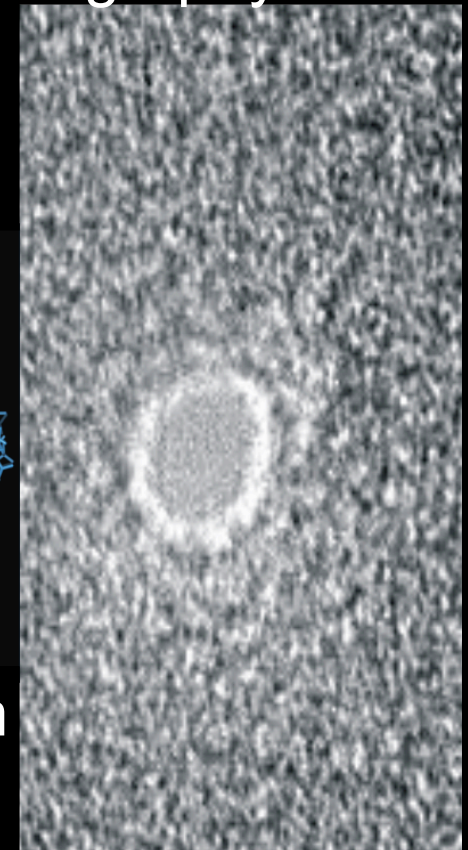
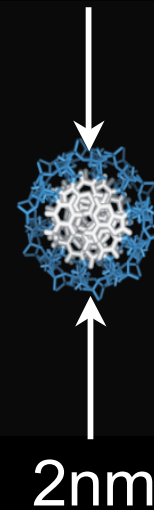
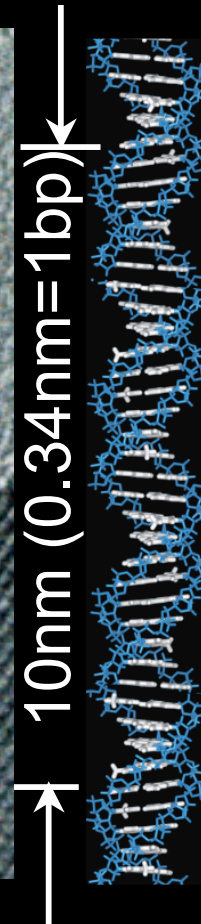
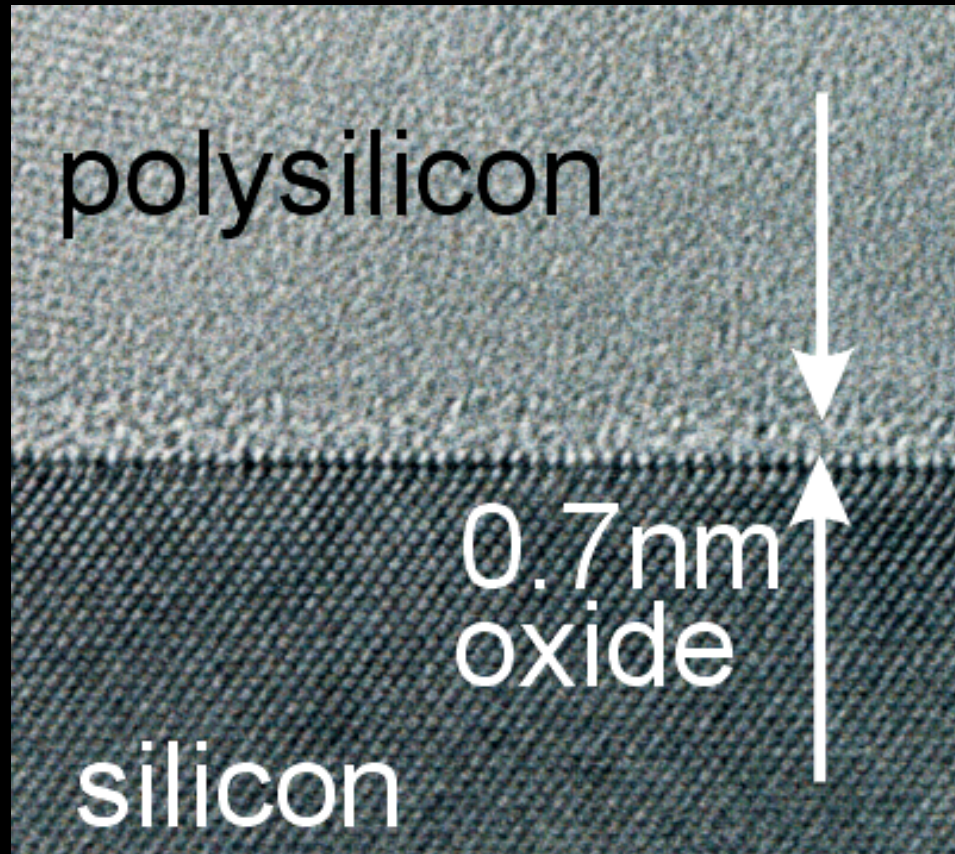


\*Work done in collaboration with M. Gracheva, A. Aksimentiev and G. Timp and supported by NIRT-NSF and NIH

# Silicon Nanotechnology for Sequencing DNA

- ultra-thin membranes

- sub-nm (sub  $\text{\AA}$  Batson)  
bright e-beam for  
lithography

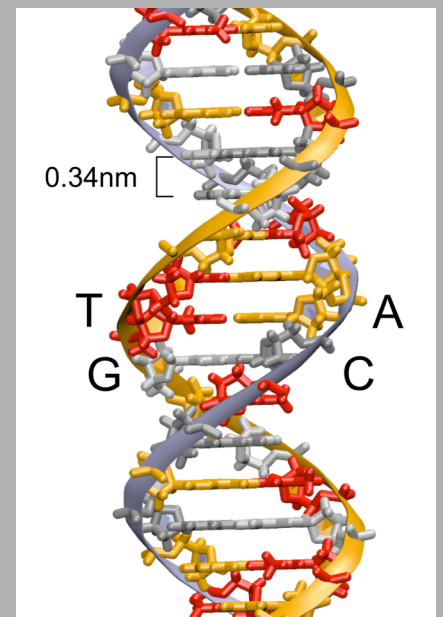
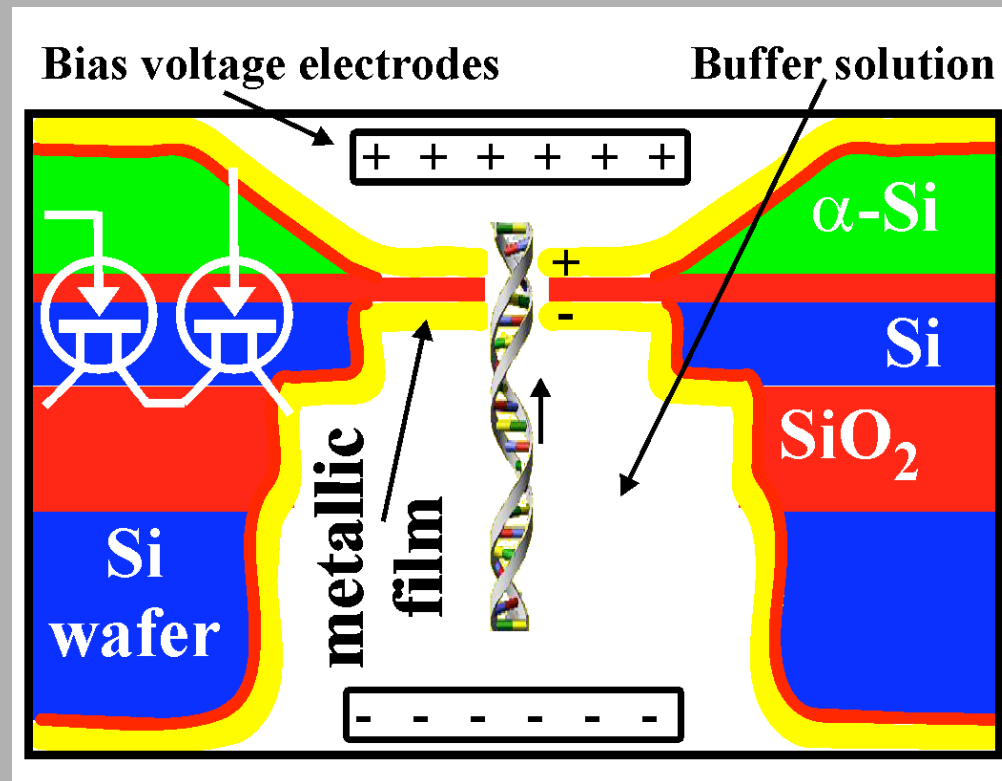


TEM X-section through a gate

DNA

TEM (top-down  
projection)

# A Challenge: Electronic Detection of DNA Molecular Sequences



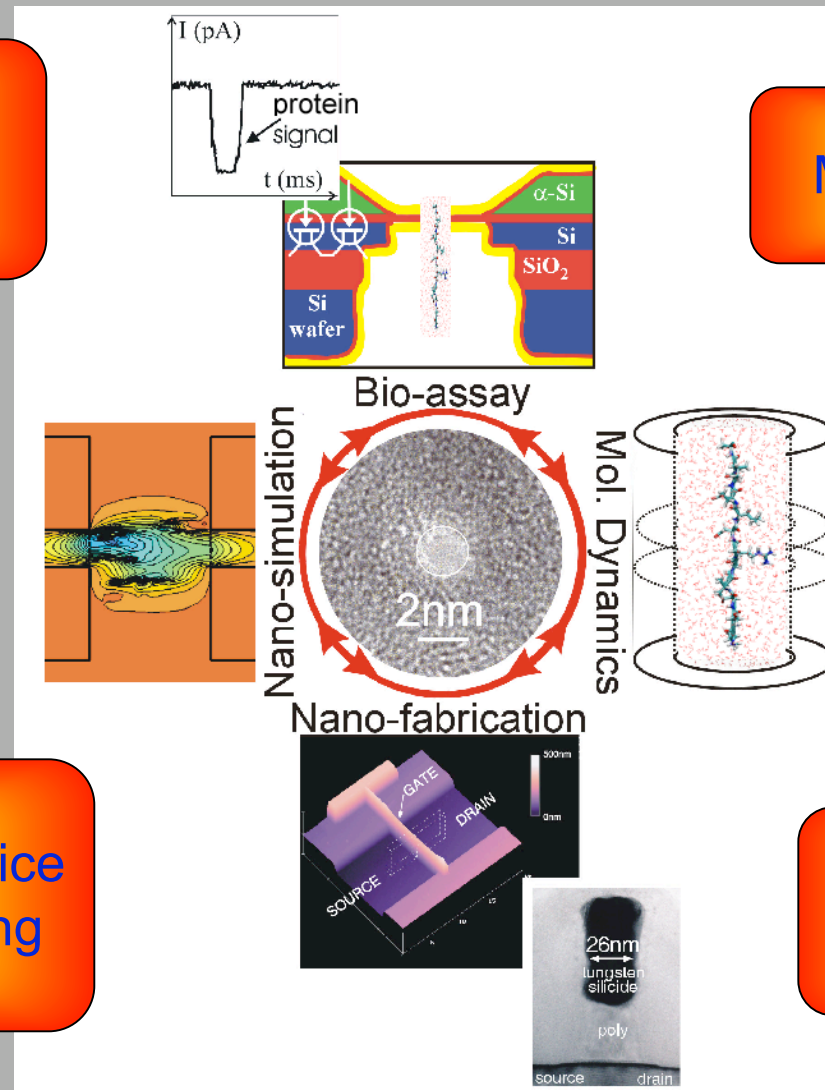
# Merging Biology and Nano-Electronics: An Integrated Approach.

DNA translocation experiments

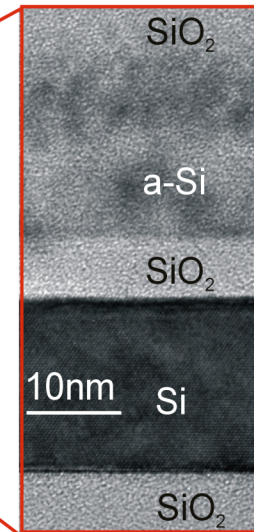
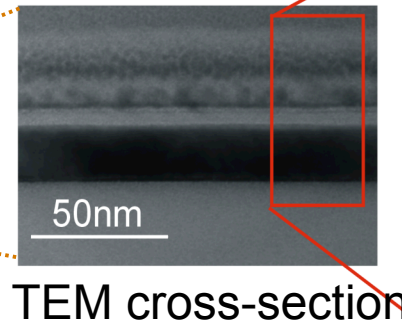
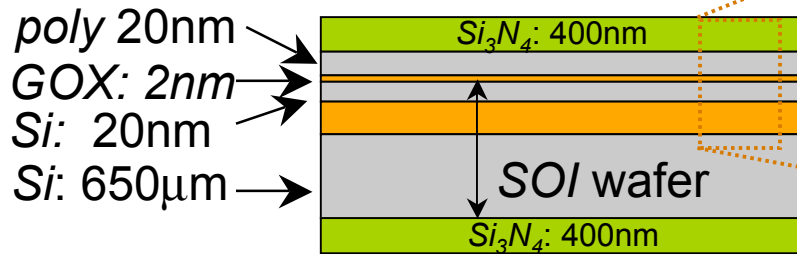
MD simulation

Self-consistent device and circuit modeling

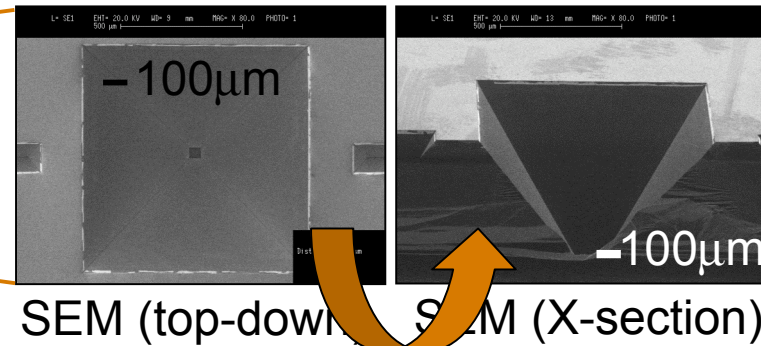
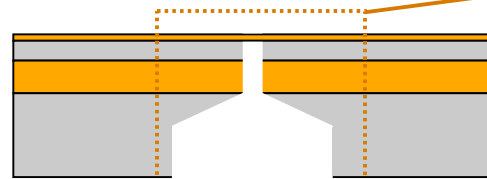
Nano-pore fabrication



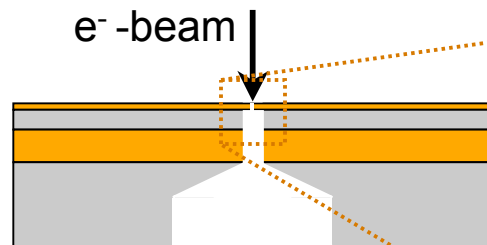
# Nanofabrication of Nanopores/Membranes



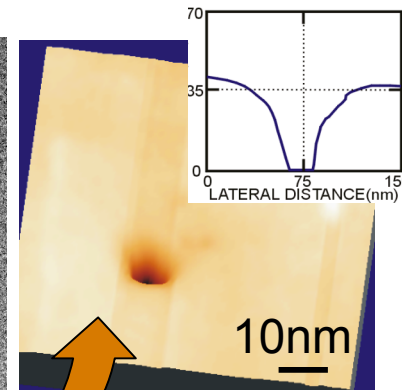
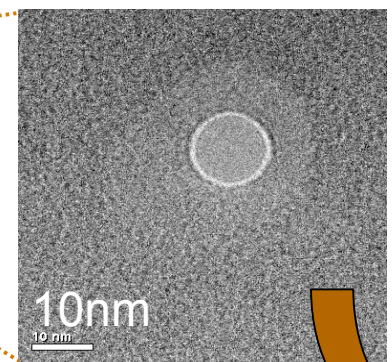
poly/device layer etching RIE+ wet etch



fabricate single pore via e-beam decomposition/ sputtering

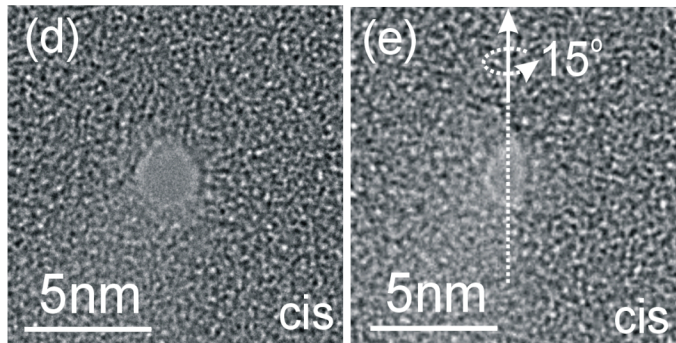
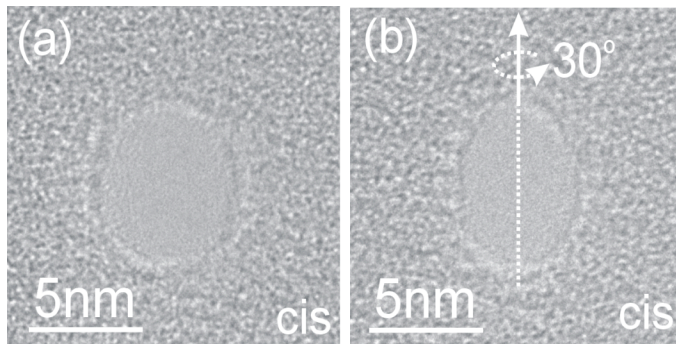
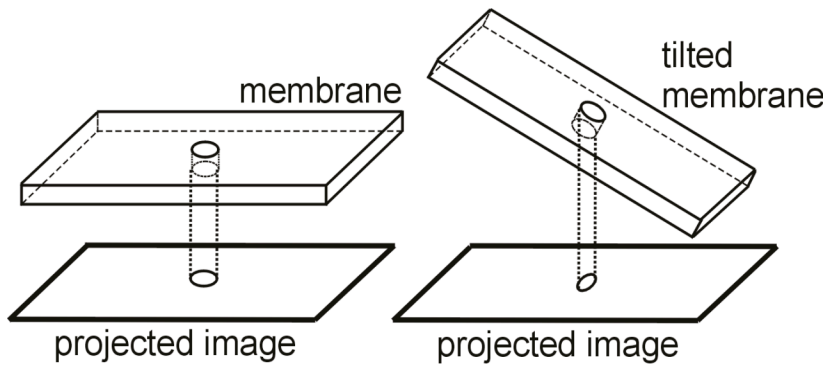


( $Si_3N_4$  membrane)

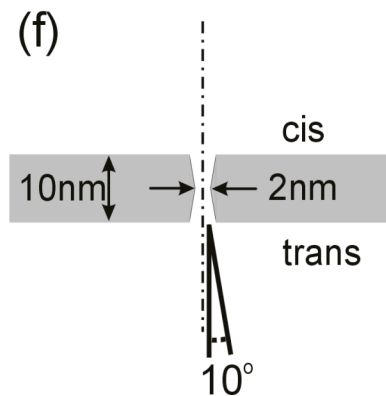
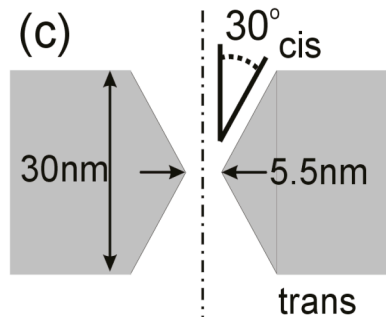


# Nanometrology of Nanopores using TEM

(artificial nanopore in  $Si_3N_4$  membrane)



TEM (top-down)    single tilt-TEM

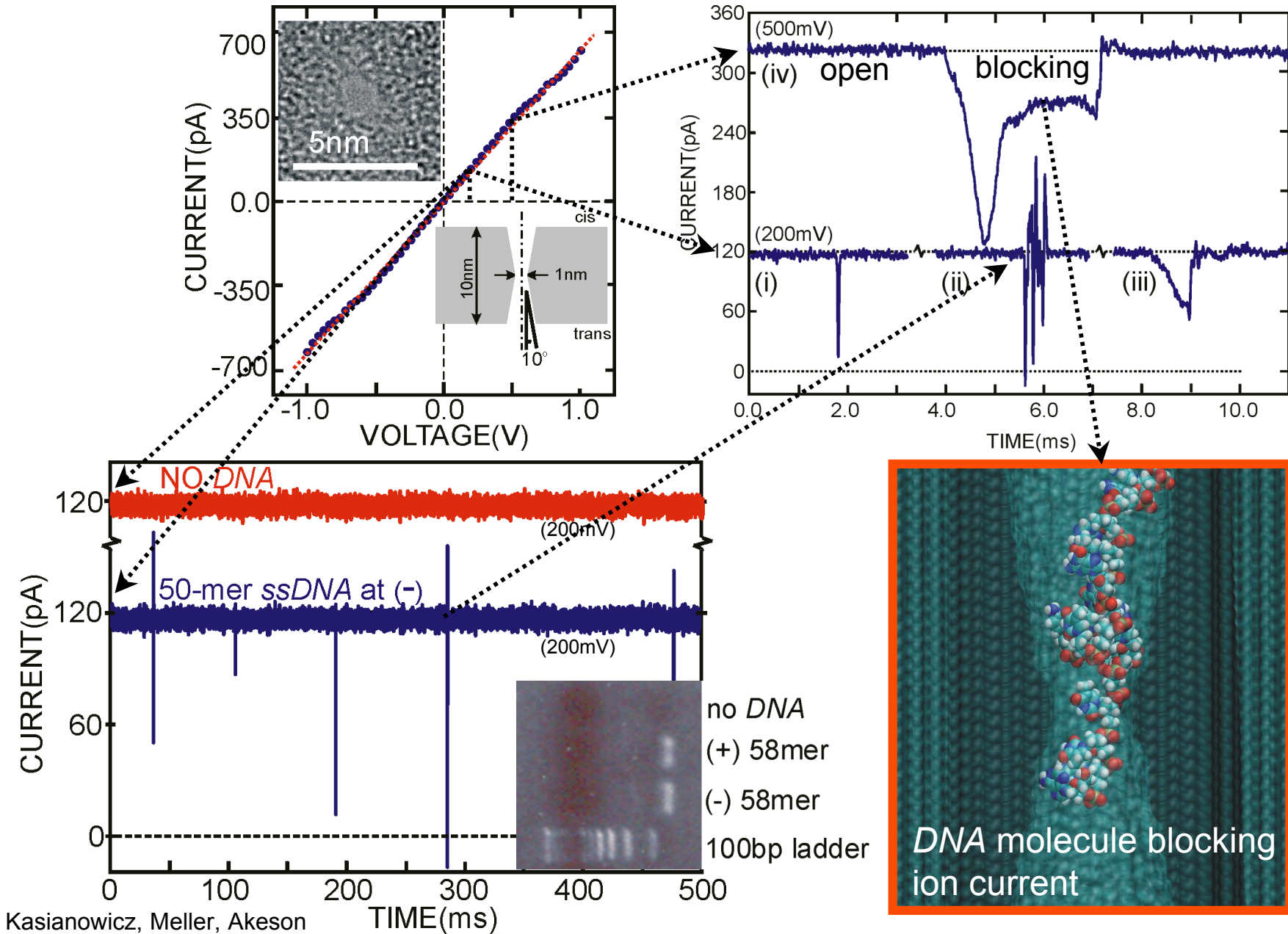


schematic

• artificial nanopores have a non-uniform cross-section

# DNA Translocations through a Nanopore

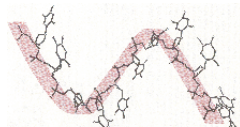
(measurements using a 1nm diameter nanopore like a molecular Coulter counter)



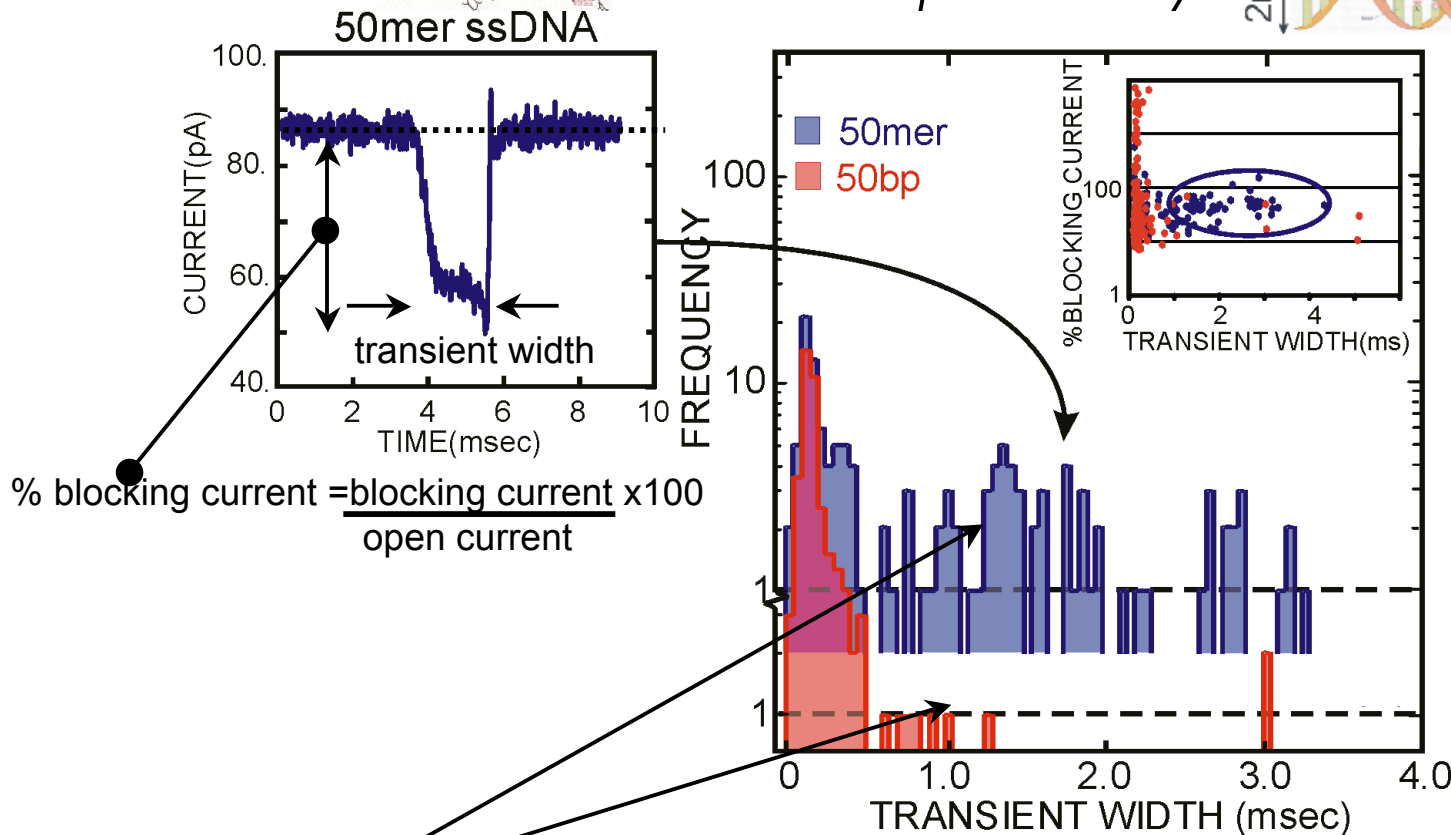
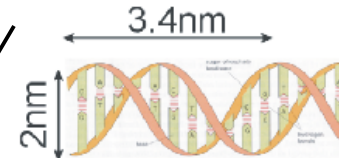
# Sorting DNA polymers

(measurements using a ~1nm diameter nanopore as a stochastic sensor)

50mer *ssDNA*: poly(dT) 0.34nm/nucleotide  
50mer=17nm p~4nm(persistence)



50bp *dsDNA*: 0.34nm/bp  
50bp=17nm p~50nm



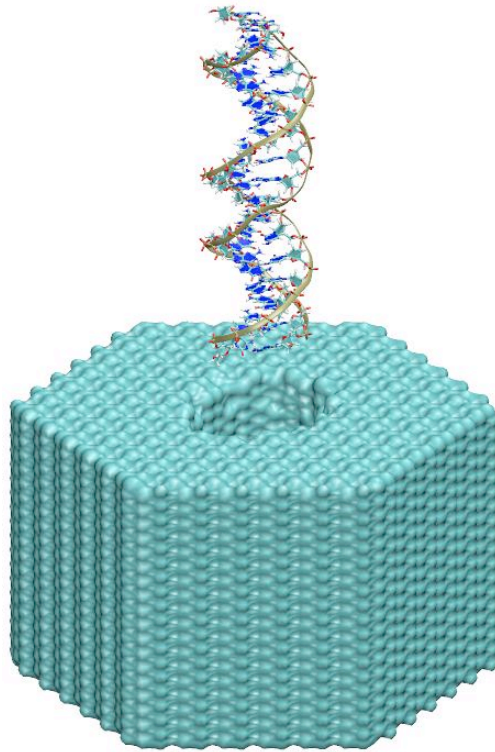
distribution of transient widths may be used to discriminate between *ssDNA*/*dsDNA*



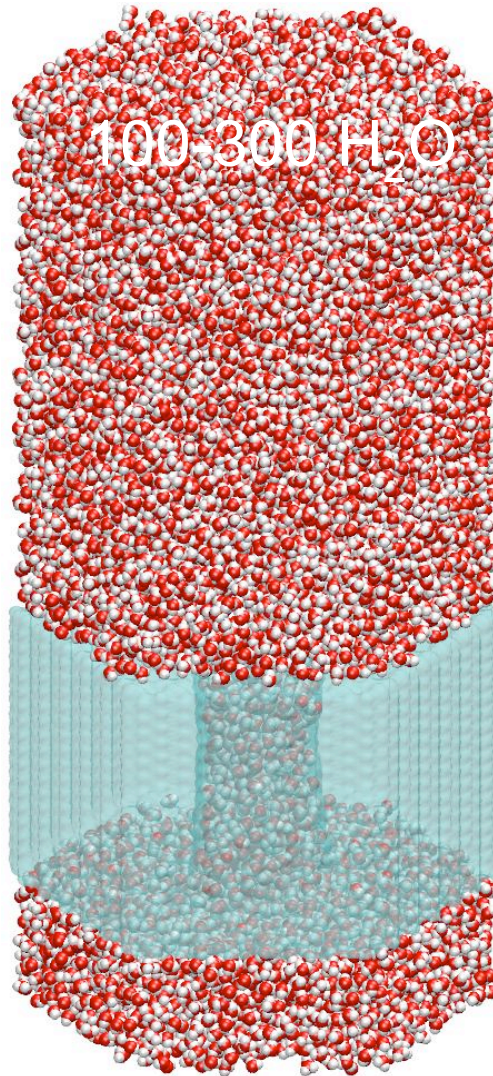
# Molecular Dynamics of DNA in a Nanopore

(force fields: CHARMM27: DNA,  $Si_3N_4$ , TIP3  $H_2O$ , ions; AMBER95: DNA)

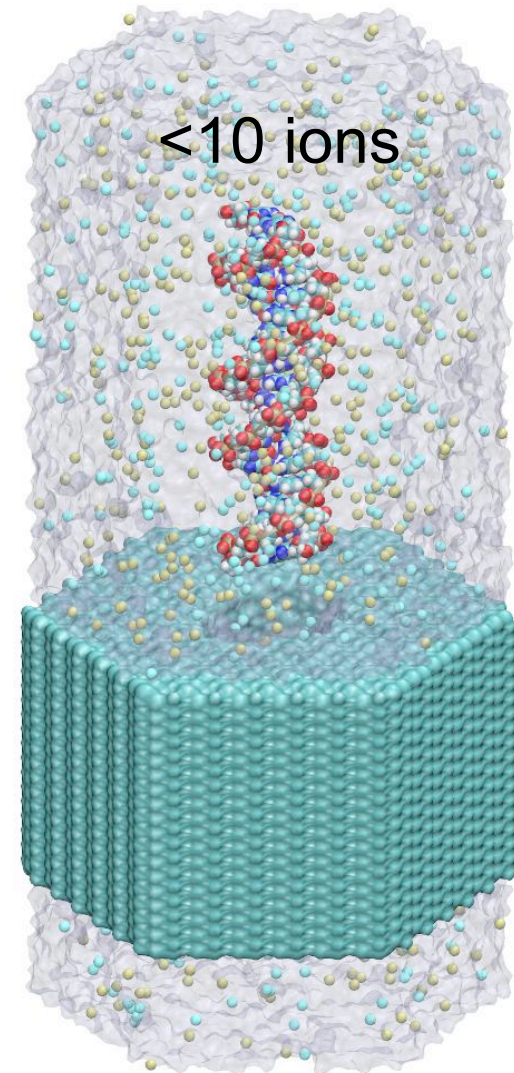
1 DNA molecule



• nanopore+DNA



• nanopore+DNA+  
water

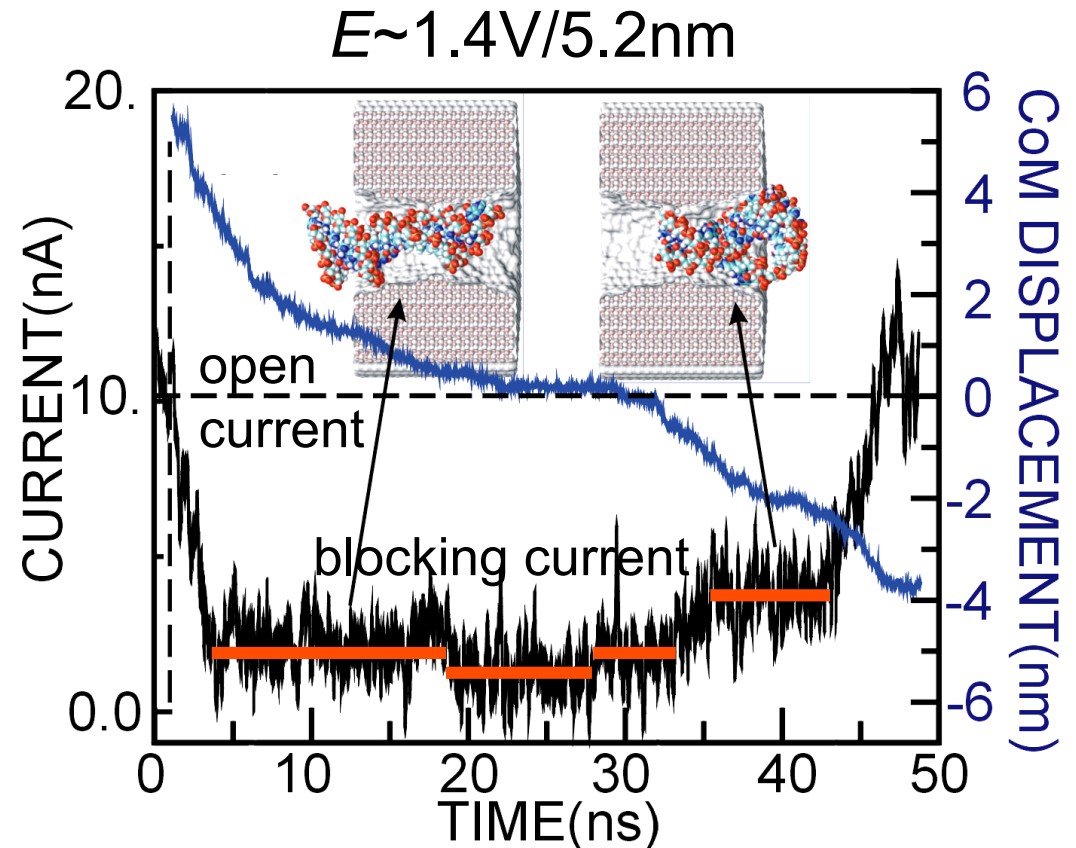


• nanopore+DNA+  
water+ions

# Molecular Dynamics of DNA in a $\text{Si}_3\text{N}_4$ Nanopore

2.4nm

QuickTime™ and a decompressor are needed to see this picture.



- translocation time: 10nsec-3 $\mu$ sec depending on field and pore interactions
- % blocking current correlated with molecular velocity
- (• large fields cause the *DNA* to denature)

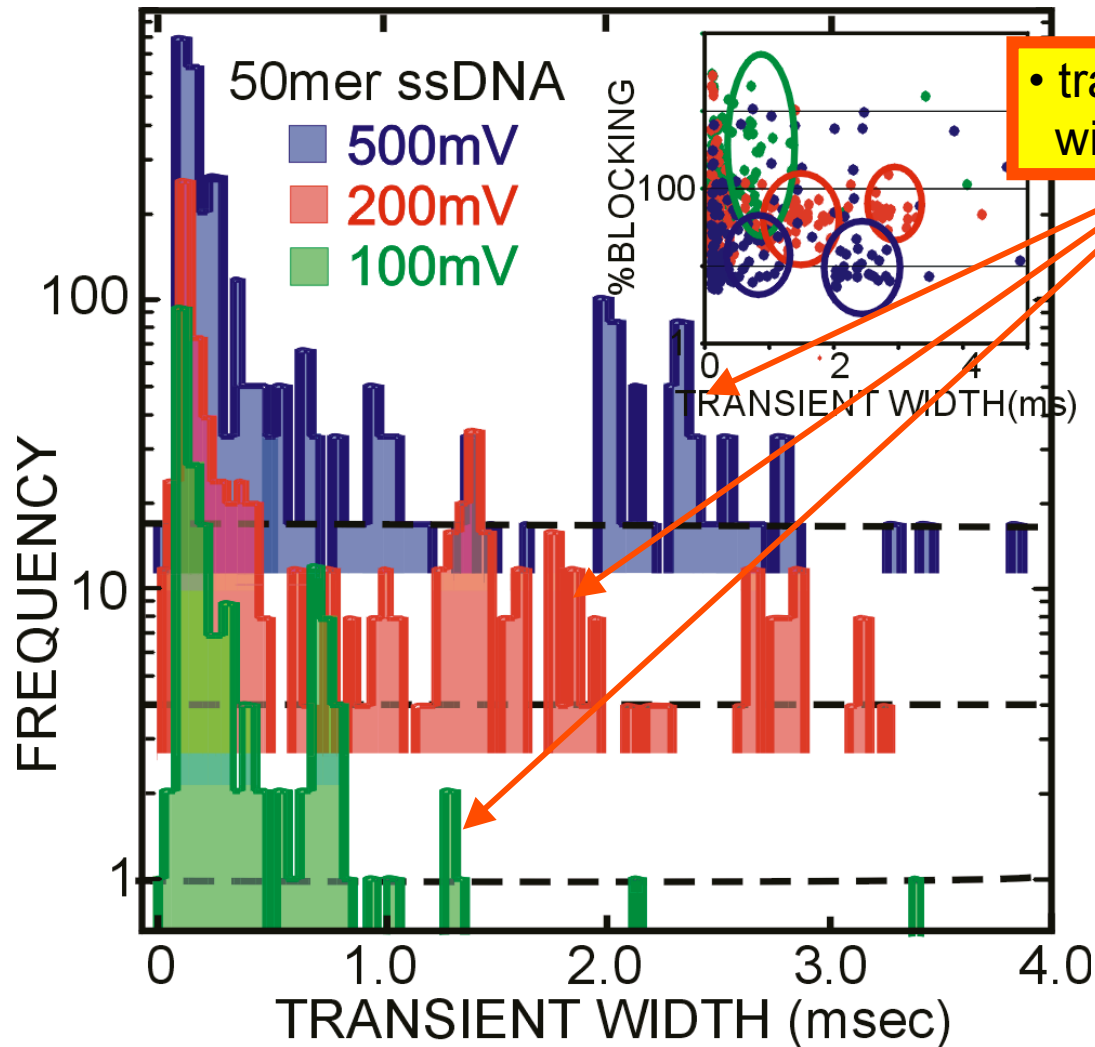
- Simulations: 1.4V/5.2nm  $\rightarrow F \sim 400\text{pN}$
- DNA sequence is CCCCCCCCCCCCCCCCCC
- Only those ions that are within 1nm of either DNA or pore are shown; some ions may appear or disappear.

*K. Schulten and A. Aksimentiev*

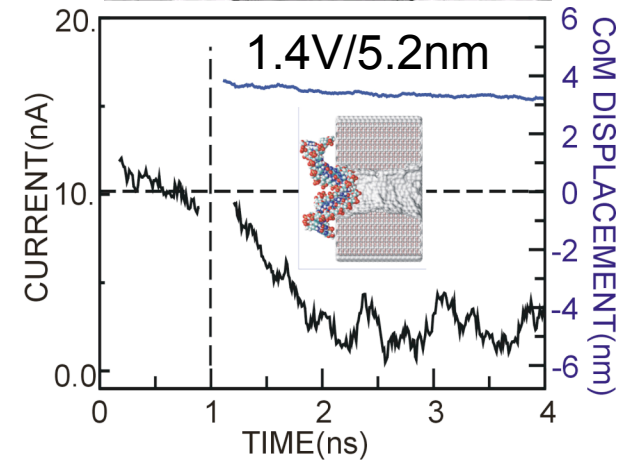
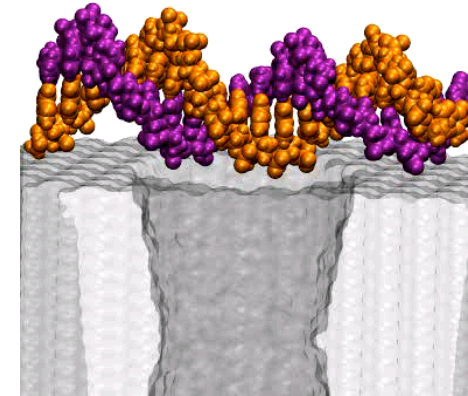
# Voltage-dependence of Blocking Current

(measurements using a 1.5nm diameter pore in a 10nm thick  $Si_3N_4$  membrane)

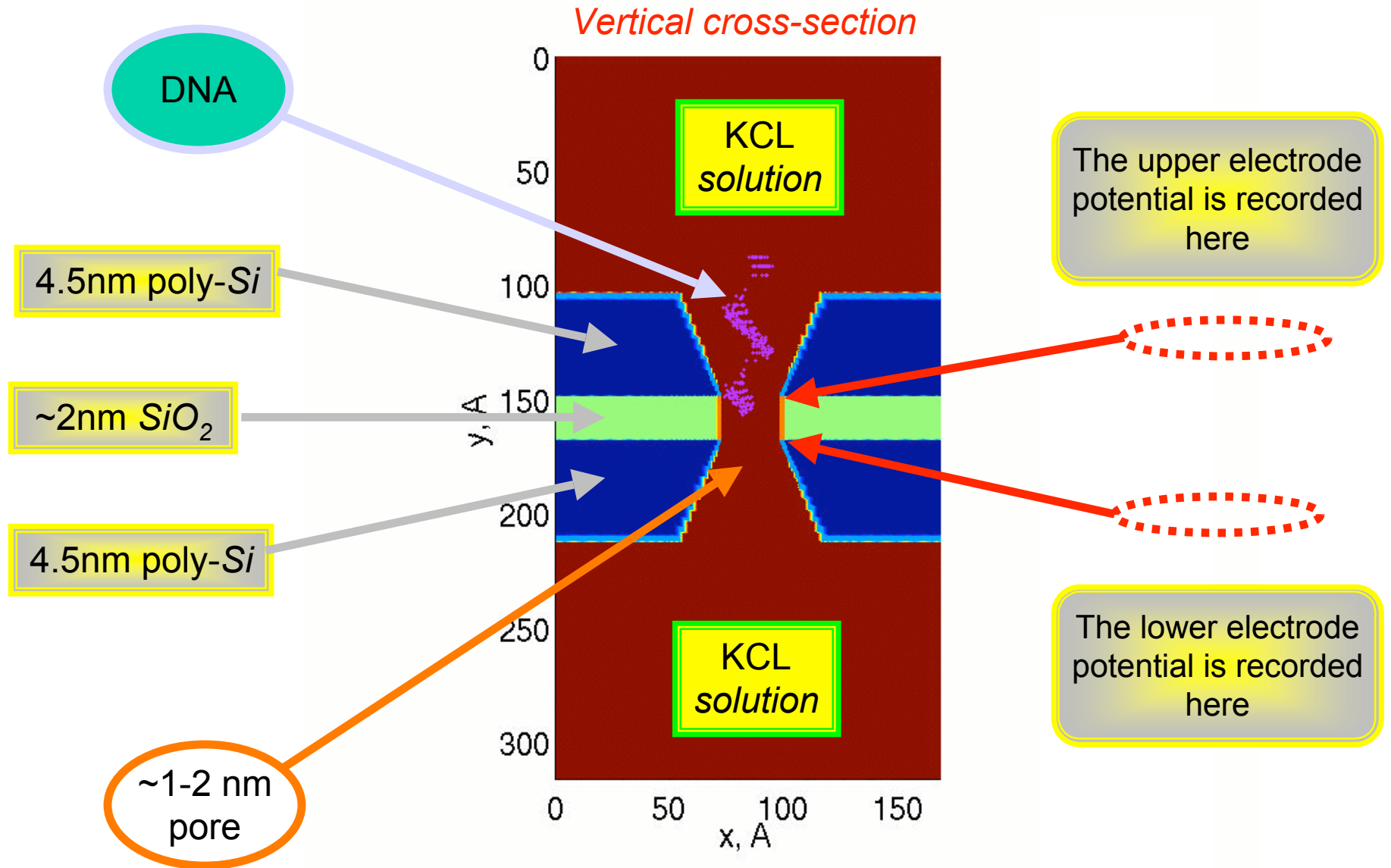
(ssDNA) poly(dT) 0.34nm/nucleotide  
50mer=17nm p~4nm(persistence)



• transient width distribution broadens with increasing voltage



# The Electronic Device Aspect



Gracheva et al. Nanotechnology 17(3), 622-633 (2006)

# Nanopore-Membrane Electrostatic Modeling

Poisson Equation:

$$\vec{\nabla} \cdot (\epsilon(\vec{r}) \vec{\nabla} \phi(\vec{r})) = -\rho(\vec{r})$$

Charge density:

$$\rho_{solid-state}(\vec{r}) = q \{ N_d^+(\vec{r}) - N_a^-(\vec{r}) + p(\vec{r}) - n(\vec{r}) \}$$

$$\rho_{solution}(\vec{r}) = q \{ [K^+](\vec{r}) - [Cl^-](\vec{r}) \} + \rho_{DNA}(\vec{r})$$

$$n(\vec{r}) = N_c \frac{2}{\sqrt{\pi}} F_{1/2}(\eta_c(\vec{r}))$$

$$\eta_c(\vec{r}) = \frac{E_f - E_c(\vec{r})}{kT}$$

$$[K^+](\vec{r}) = [K^+]_0 \exp(q\phi(\vec{r}) / kT)$$

$$p(\vec{r}) = N_v \frac{2}{\sqrt{\pi}} F_{1/2}(\eta_v(\vec{r}))$$

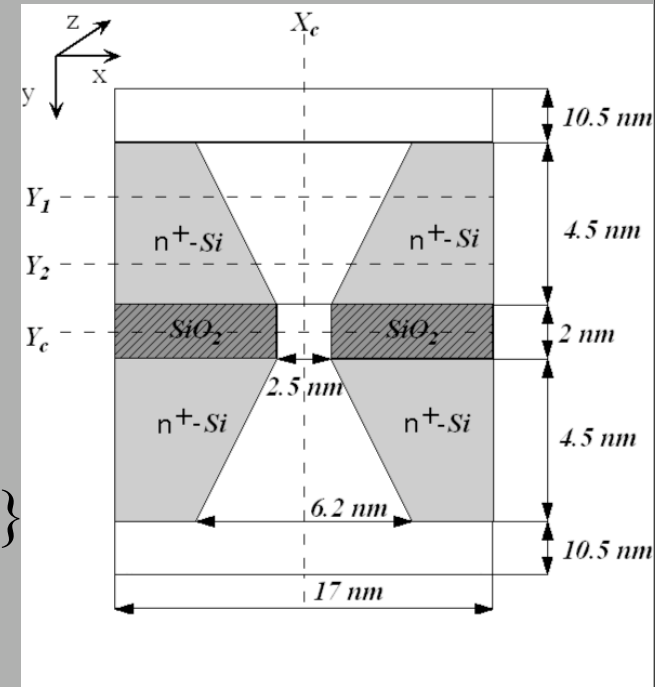
$$\eta_v(\vec{r}) = \frac{E_v(\vec{r}) - E_f}{kT}$$

$$[Cl^-](\vec{r}) = [Cl^-]_0 \exp(-q\phi(\vec{r}) / kT)$$

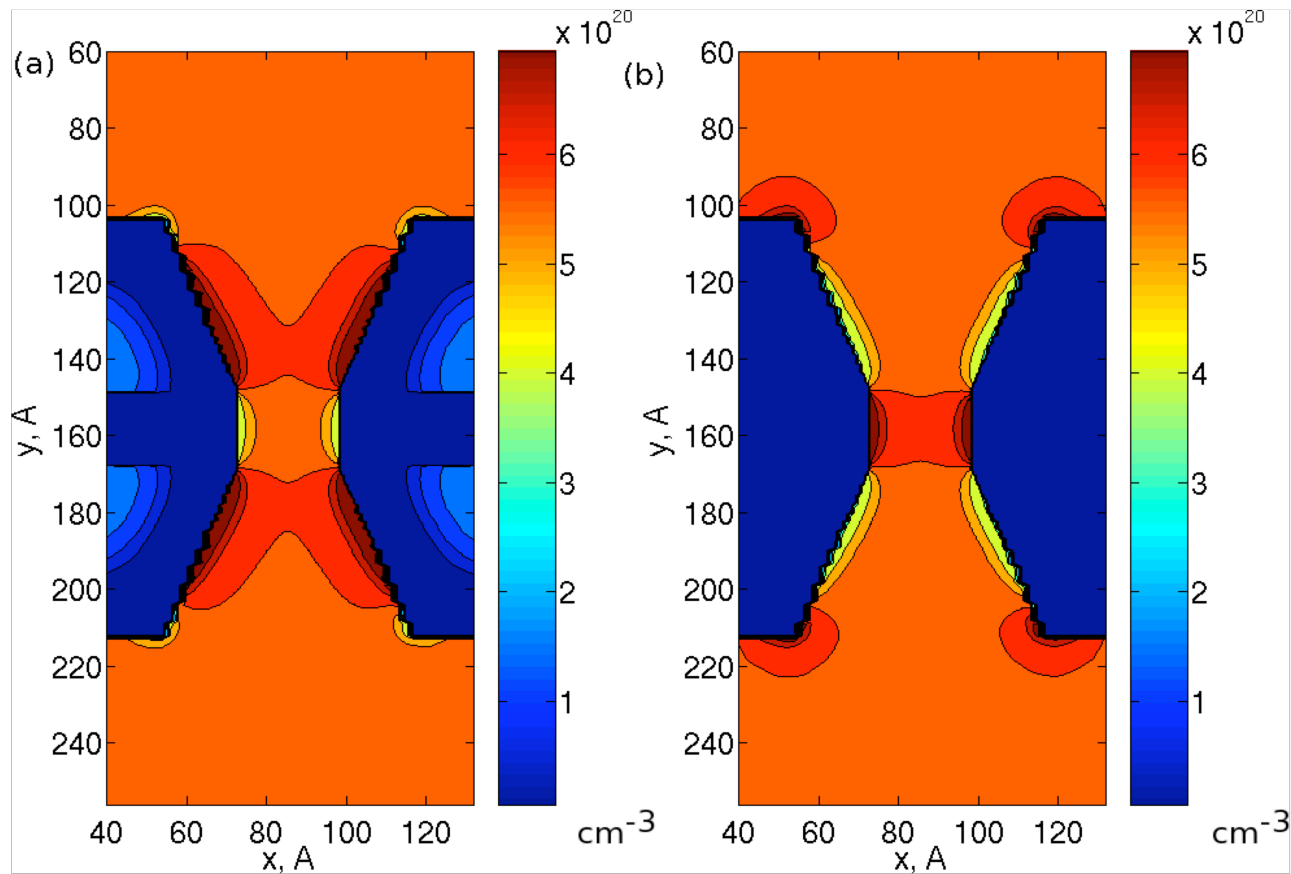
$$[K^+]_0 = [Cl^-]_0 = c$$

$$E_c(\vec{r}) = -q\phi(\vec{r}) - E_g$$

$$E_v(\vec{r}) = -q\phi(\vec{r})$$

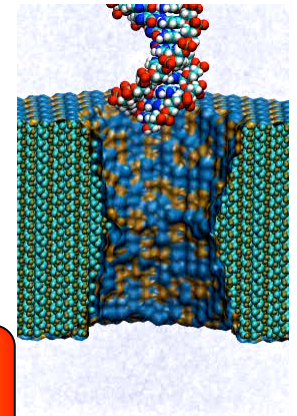


# Empty pore negative (left) and positive (right) charge in the structure and solution



# Multi-Scale “Self-consistent” Poisson Solver

(used in conjunction with “snapshots” provided by MD)



**Nanopore structure design**

Nanopore structure design

Electrolytic and semiconductor charge distribution

**MD simulation**

Molecular dynamics simulation

DNA charge distribution “snap-shot”

**Semiconductor device simulation**

Solve self-consistent 3-D Poisson equation

Updated voltage signal

Final time step

Next time step

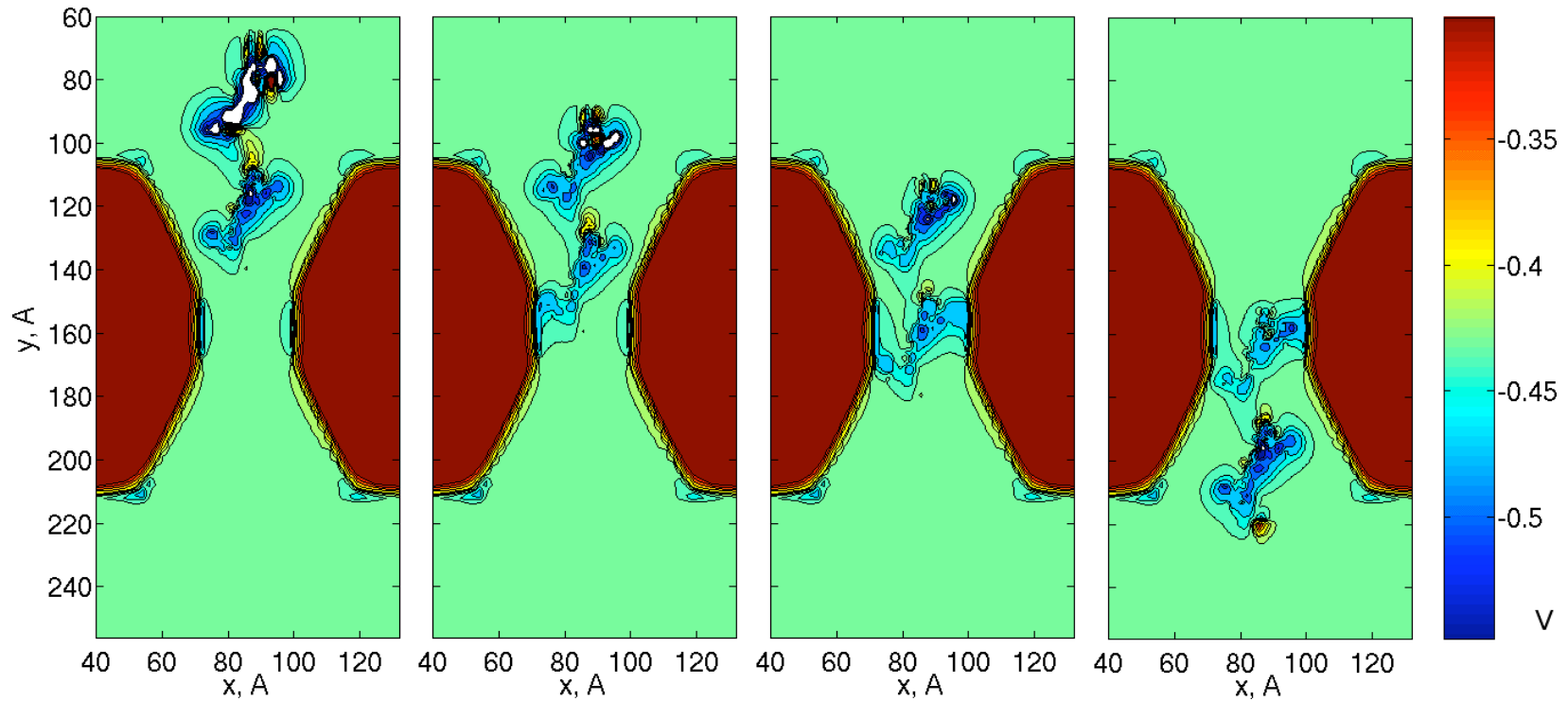
Exit

$$\nabla(\epsilon\nabla\phi) = -\rho$$

$$\rho = \begin{cases} [K^+] - [Cl^-] & \text{in electrolyte} \\ [K^+]_0 \exp(q\phi/kT); [Cl^-] = [Cl^-]_0 \exp(-q\phi/kT) \\ q(p - n + N_d^+ - N_a^-) & \text{in semiconductor} \end{cases}$$

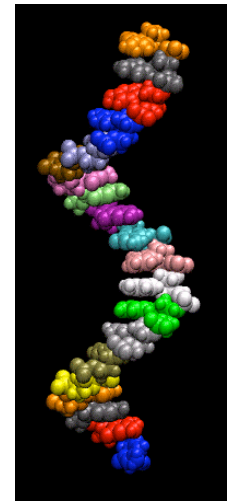


# Simulation of ssDNA Translocation Through a 2.5nm Nanopore

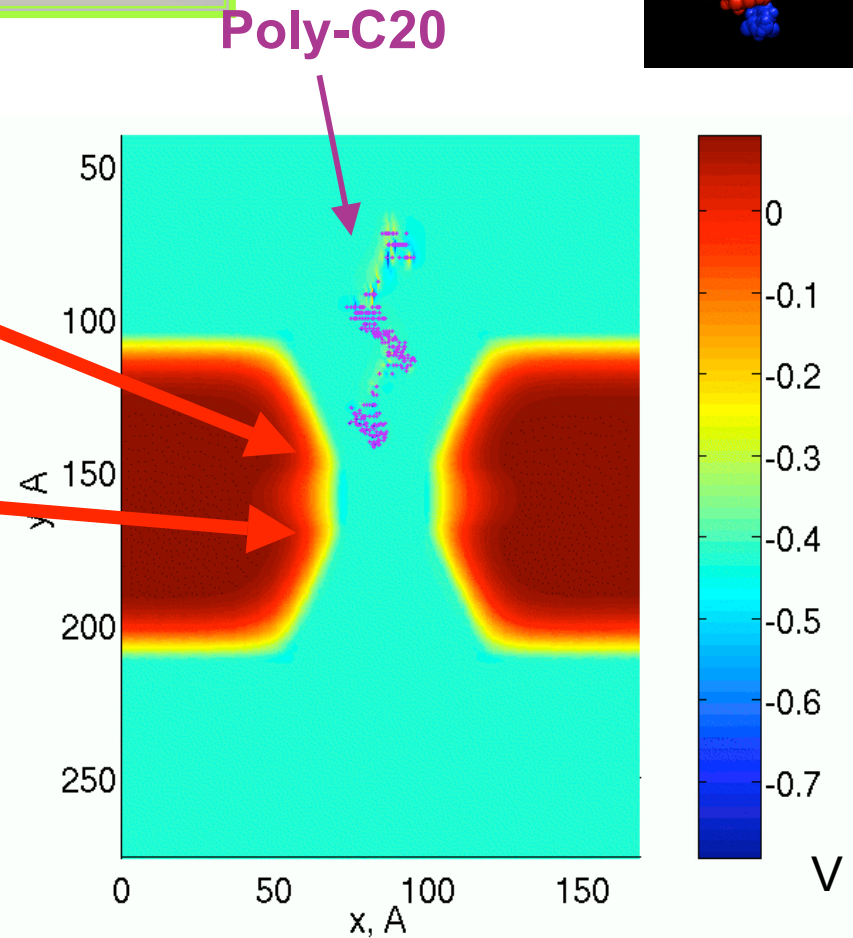
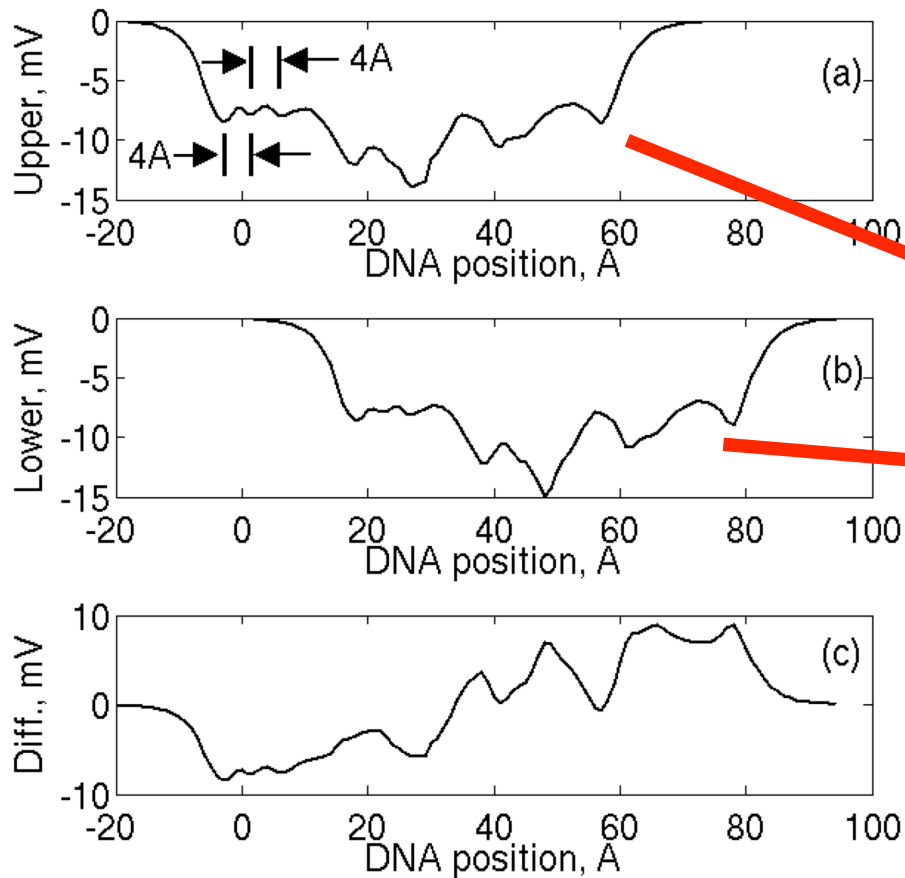




# Simulation of unstretched ssDNA Translocation Through a 2.5 nm Nanopore

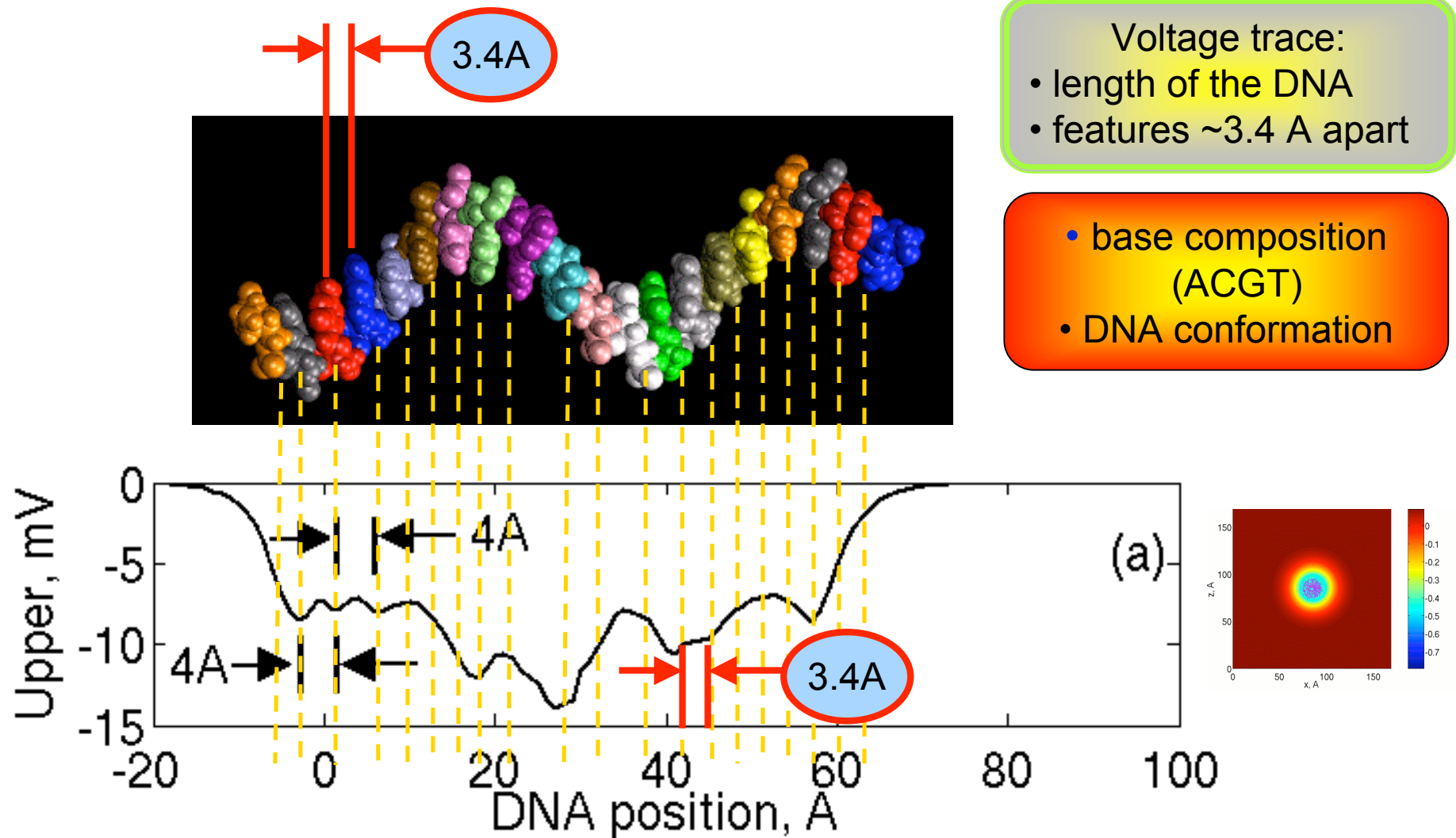


Electrode Voltage is sensitive to DNA charge and conformation



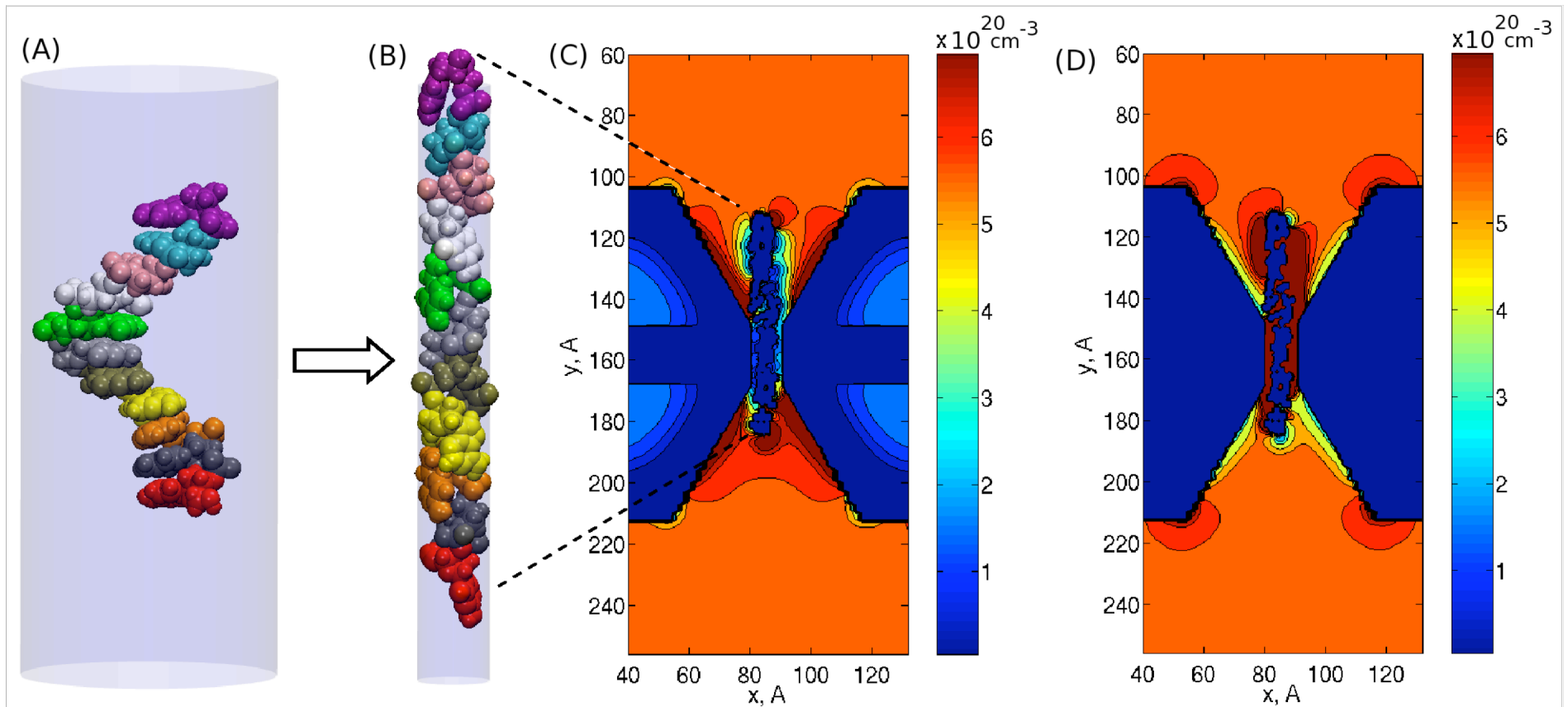
# Decoding DNA sequence and configuration

DNA sequence: C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C-C (ssDNA polyC20)



# Simulation of Stretched ssDNA Translocation Through a 1.0nm Nanopore

11 bases

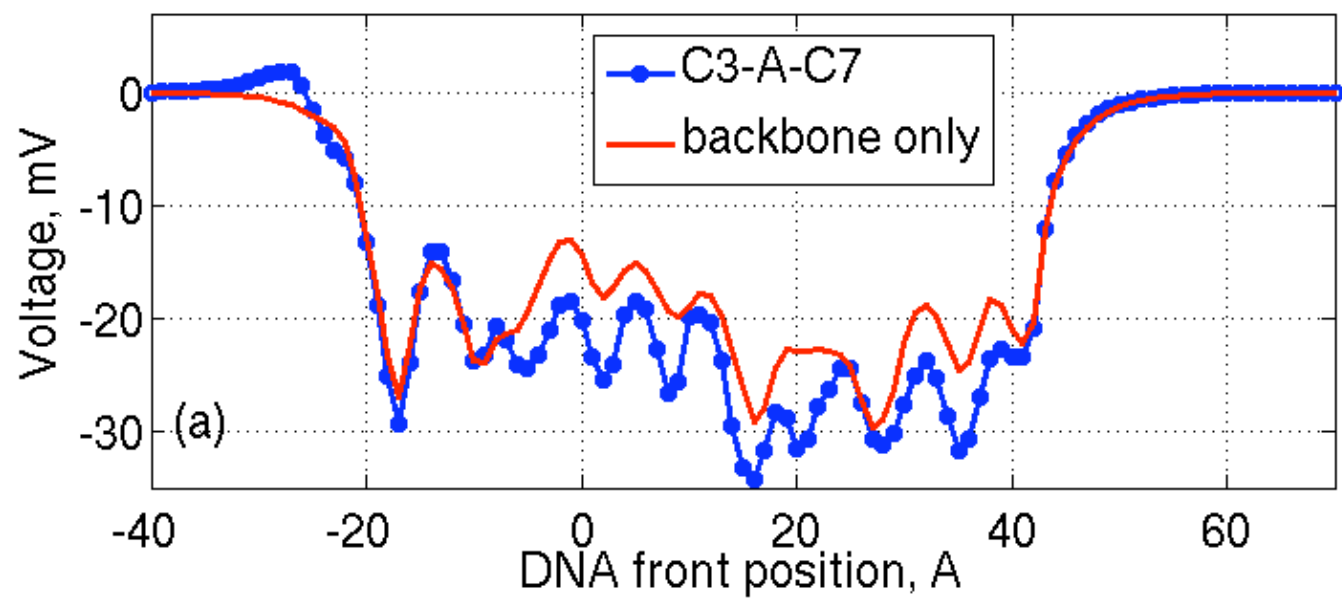


2.5nm  
pore

1.0nm  
pore

Gracheva et al Nanotechnology 17(13), p.3160 (2006)

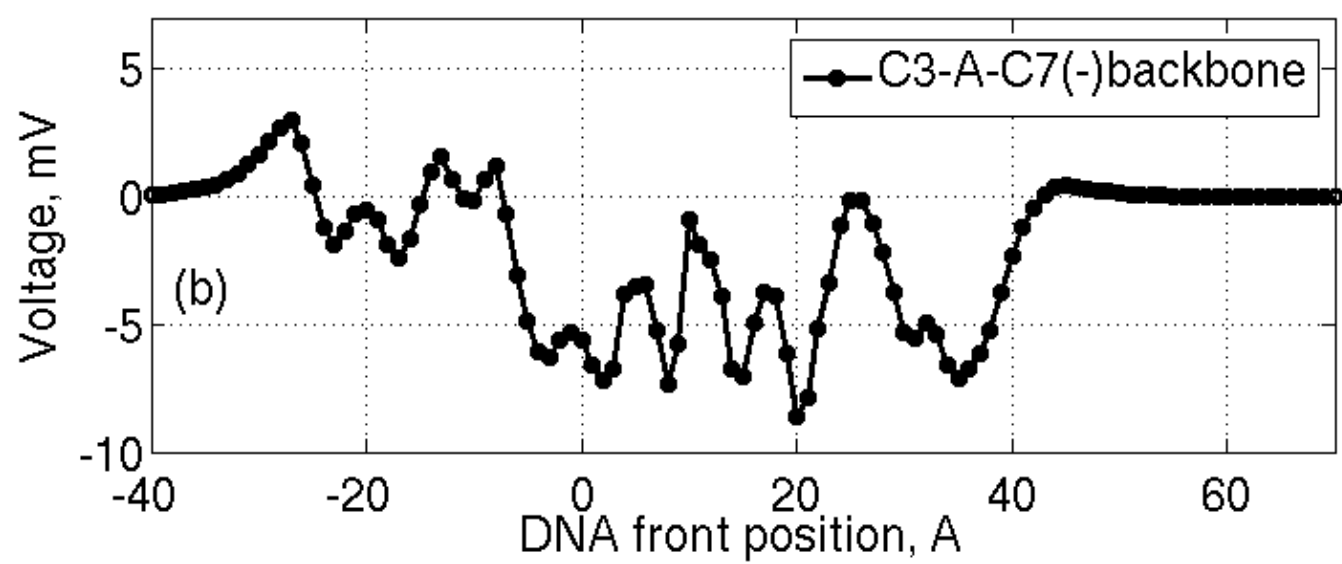
Upper electrode voltage for  $C_3AC_7$  and its backbone



11 bases

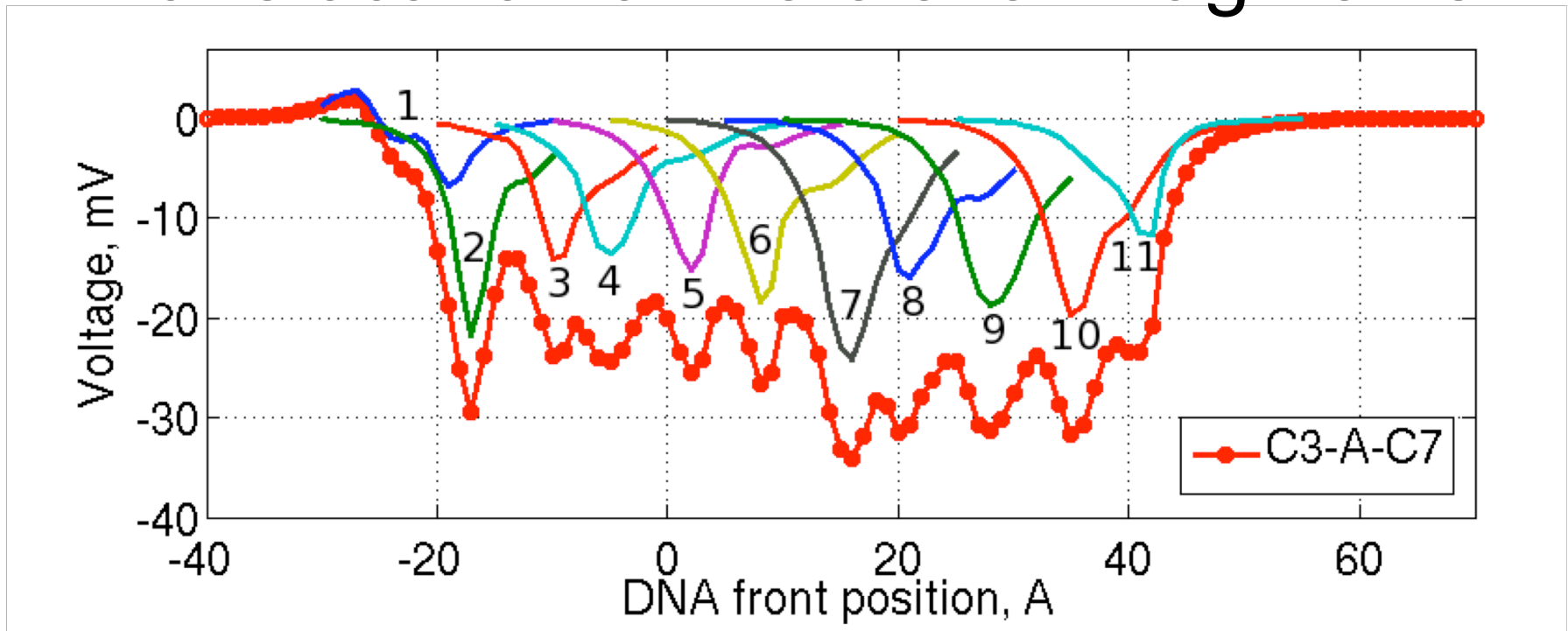
C3-A-C7

Difference of  $C_3AC_7$  and its backbone (for upper electrodes)

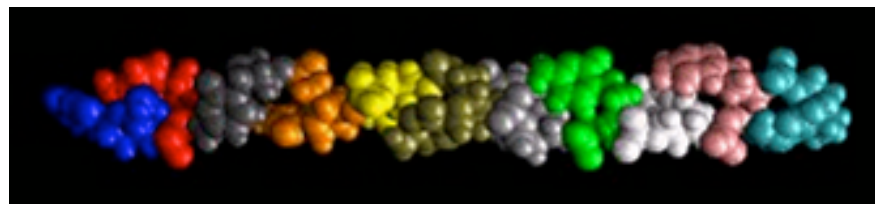


Backbone signal  
~30mV  
Bases ~10mV

# The whole DNA translocation and translocation of the eleven fragments

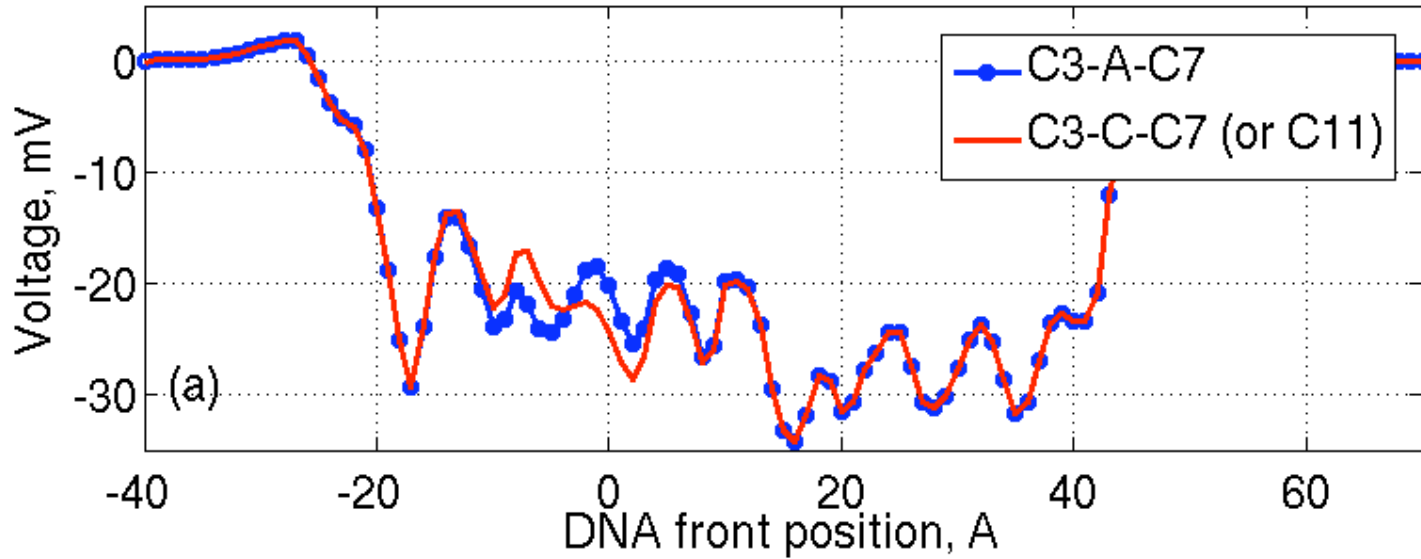


C3-A-C7  
and 11  
backbone  
segments  
with  
bases



Look how wide  
the signal  
of one base  
can be!!!

Upper electrode voltage for  $C_3AC_7$  and  $C_{11}$

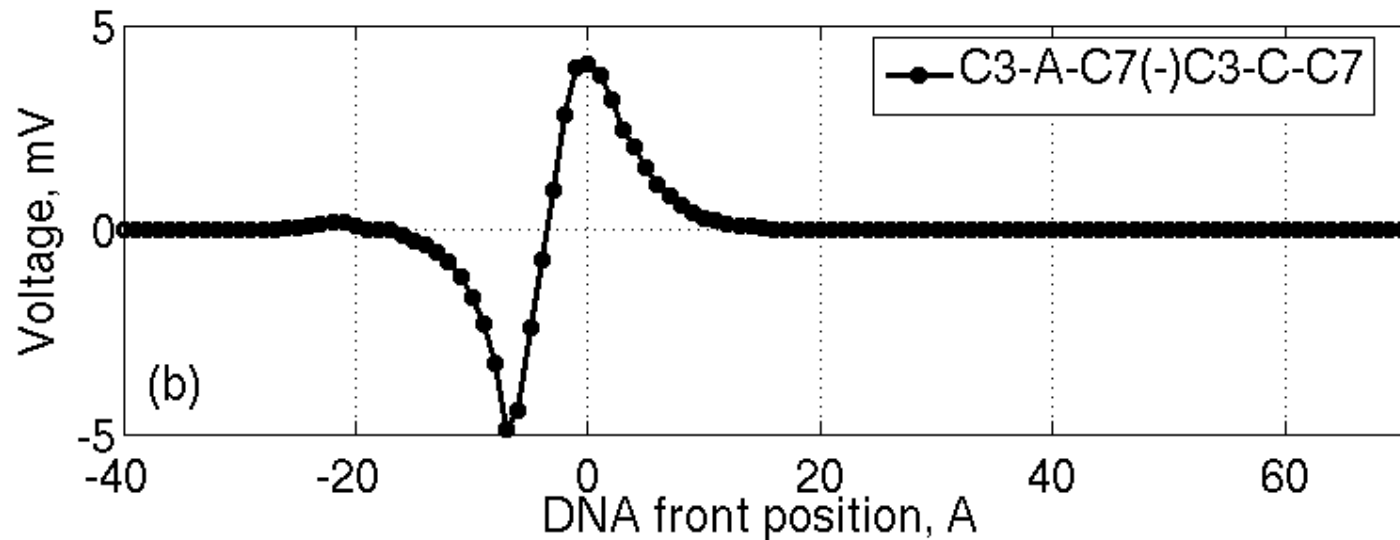


**In the same conformation!!!**

C3-A-C7

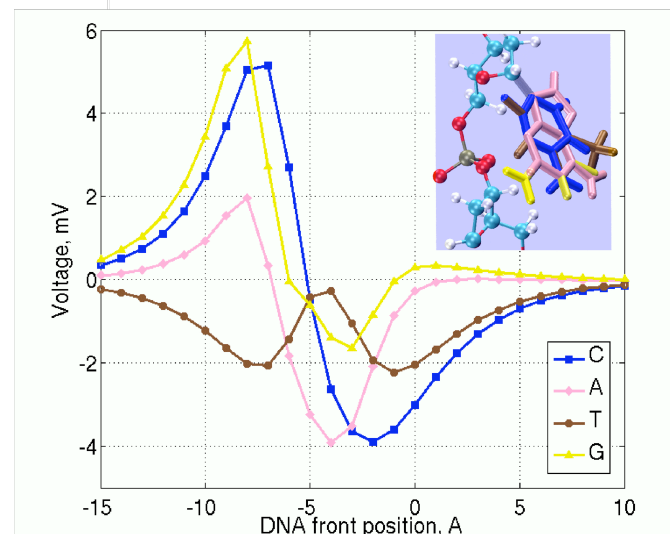
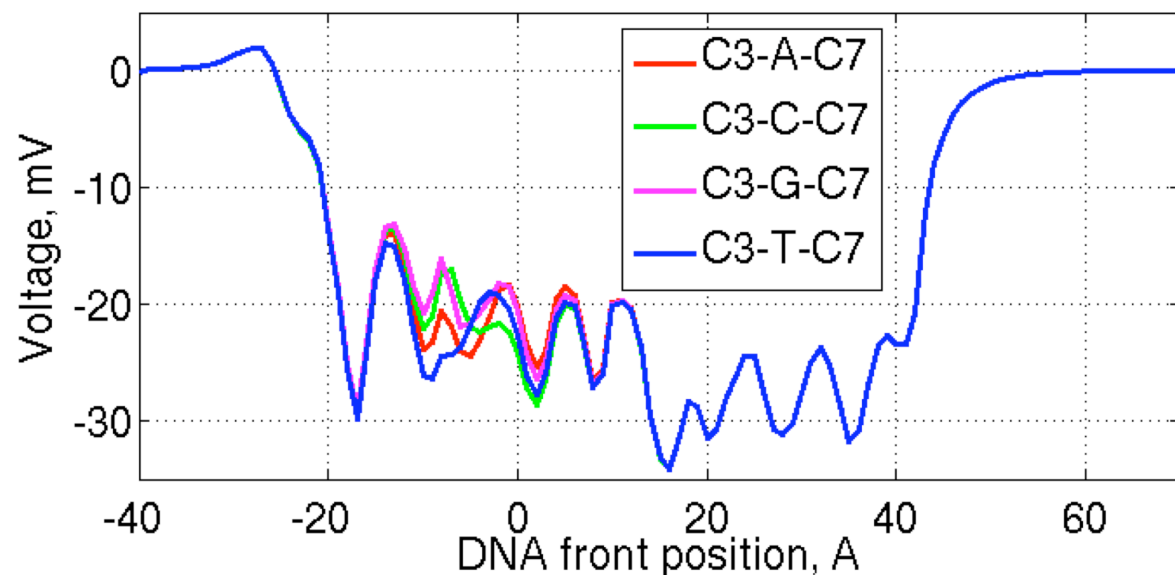
C3-C-C7  
or  
C11-mutant

Difference of  $C_3AC_7$  and  $C_{11}$  (for upper electrodes)



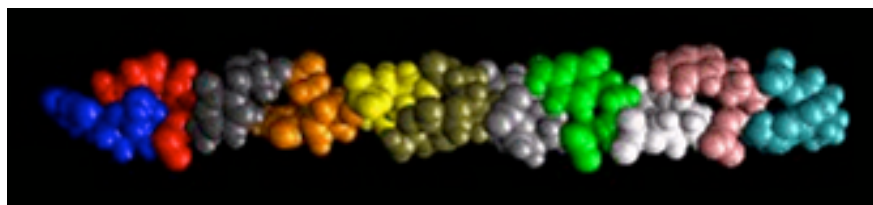
Difference  
between A and C  
~5mV

# The whole DNA translocation with one base mutation



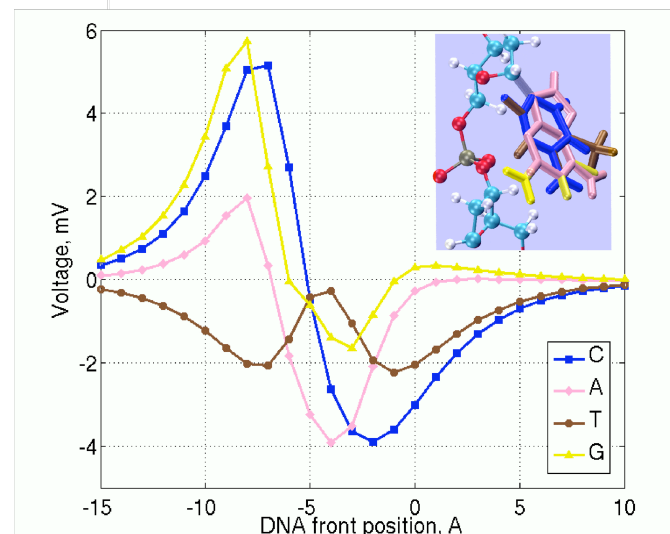
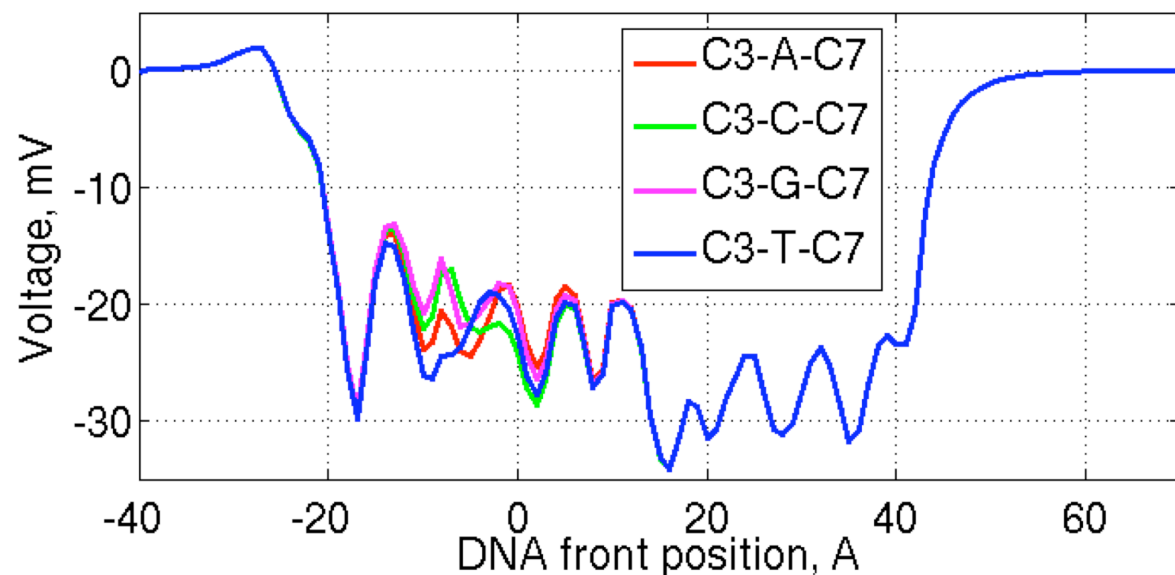
Mutated  
base #4

In the same  
conformation!!!



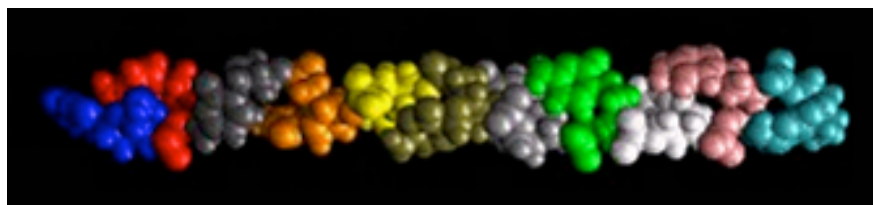
The signal  
difference  
is localized in  
the vicinity of  
base #4

# The whole DNA translocation with one base mutation



Mutated  
base #4

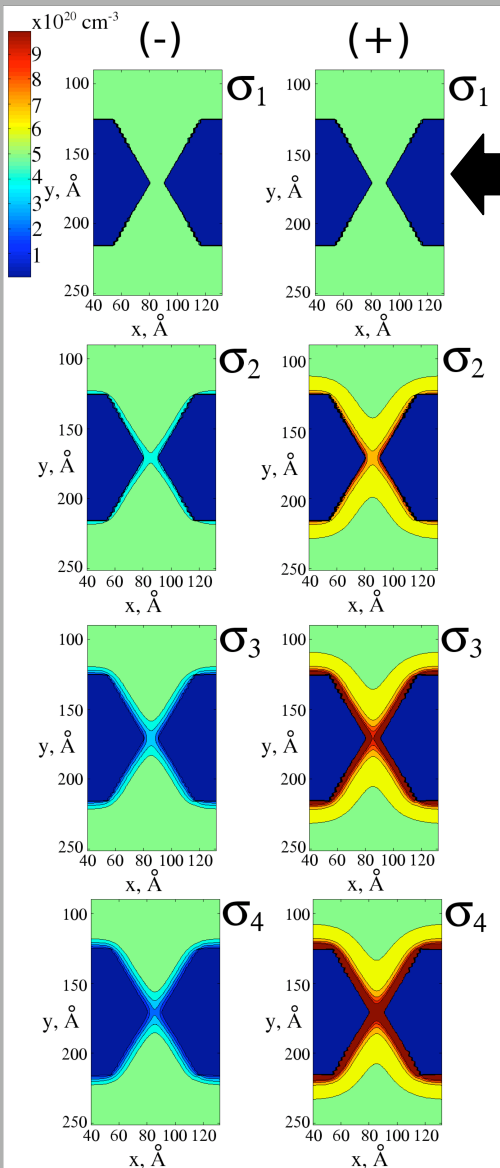
In the same  
conformation!!!



The signal  
difference  
is localized in  
the vicinity of  
base #4



# Semiconductor Membrane: Double Layer Engineering



$\text{SiO}_2$ : dielectric material  
no carriers of either sign

$n\text{-Si}$ : heavily doped Si  
semiconductor material  
Fixed positive charge  
 $N_d^+ = 2 \times 10^{20} \text{ cm}^{-3}$ ,  
electrons, some holes

Surface charge:

$$\sigma_1 = 0$$

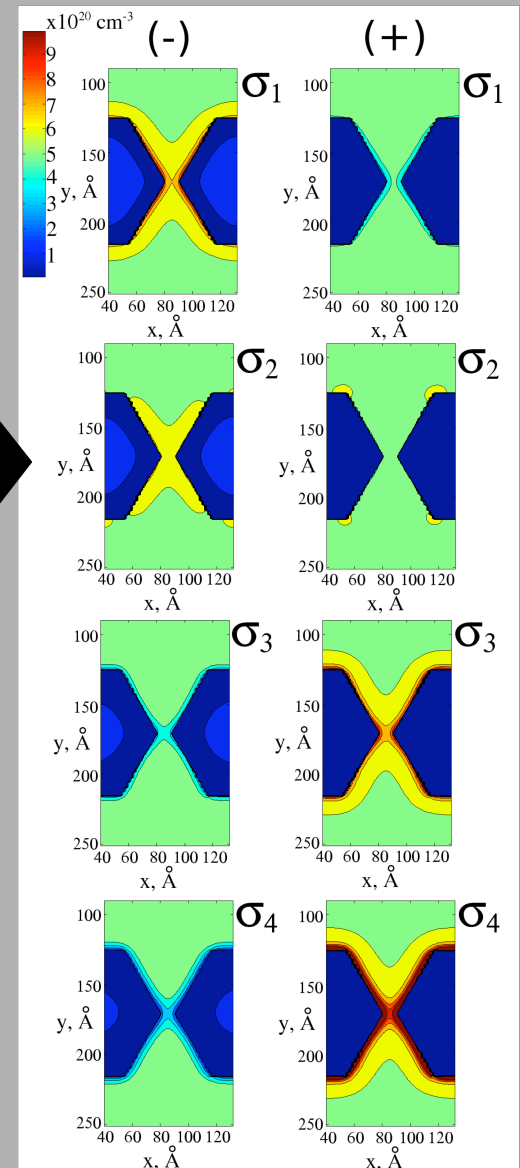
$$\sigma_2 = -0.0064 \text{ Cm}^{-2}$$

$$\sigma_3 = -0.019 \text{ Cm}^{-2}$$

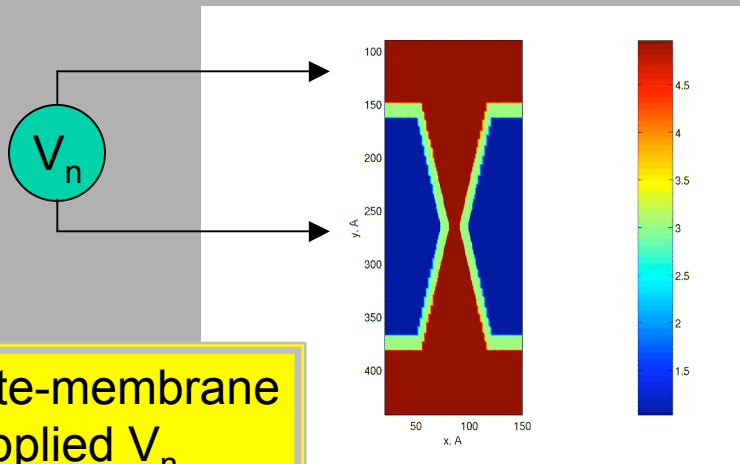
$$\sigma_4 = -0.032 \text{ Cm}^{-2}$$

[KCl]=1M

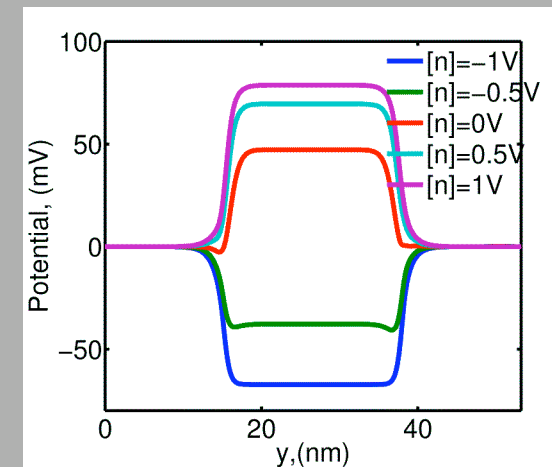
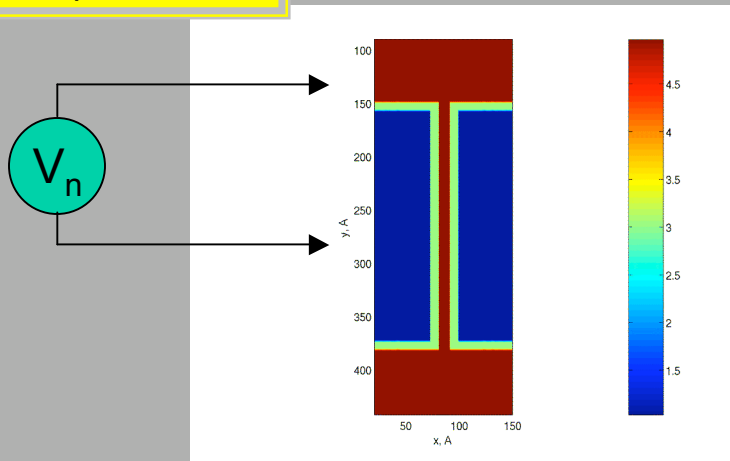
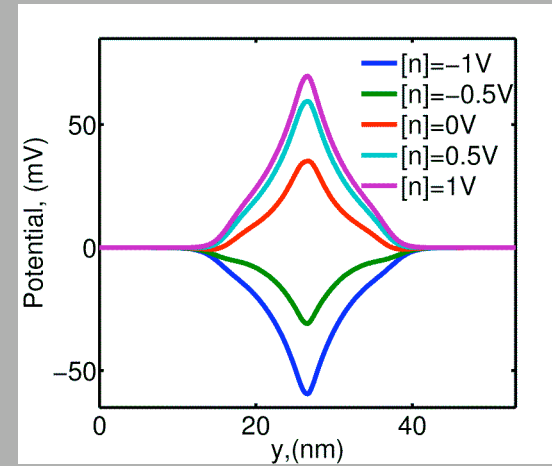
Gracheva & Leburton  
Nanotechnology 18, p.145704 (2007)



# Semiconductor Membrane: Electrical Tunability



Electrolyte-membrane bias is applied  $V_n$  ([n] in figures)



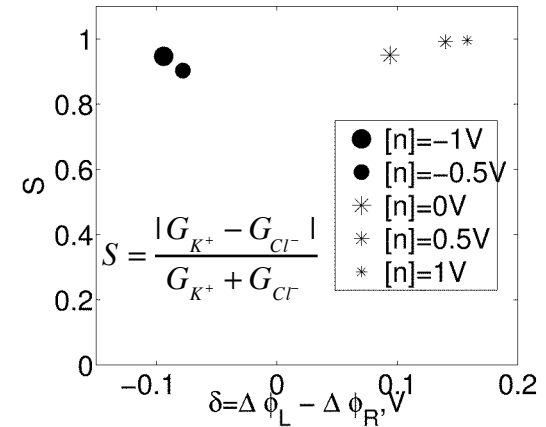
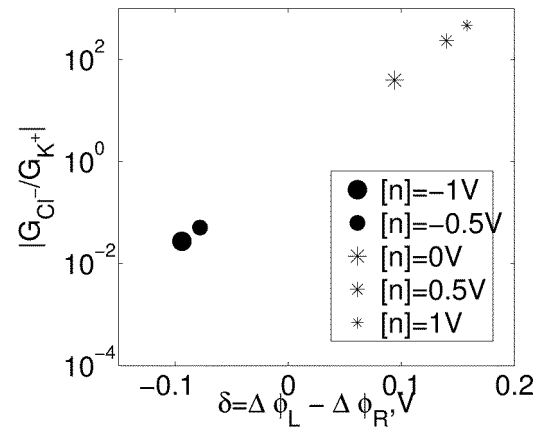
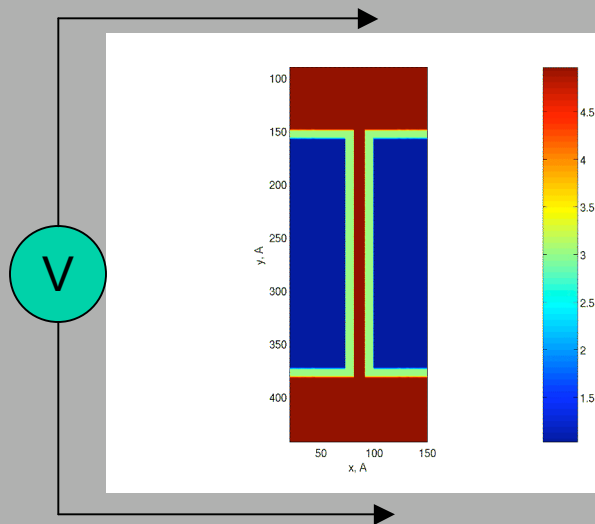
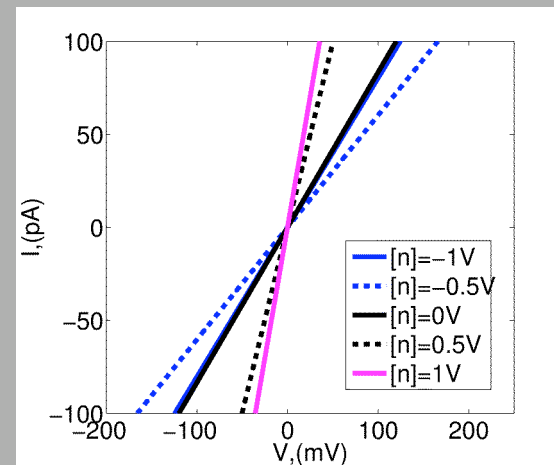
[KCl]=0.1M

# Controlled Ionic Conductance & Filtering (Linear Response)

Current-voltage characteristics:  
assume constant field in the  
nanopore

Conductance:  $G=dI/dV$

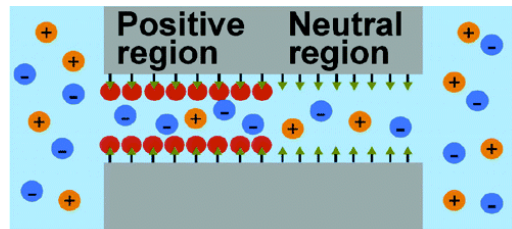
Selectivity:  $S=|(G_{Cl^-}-G_{K^+})/(G_{Cl^-}+G_{K^+})|$



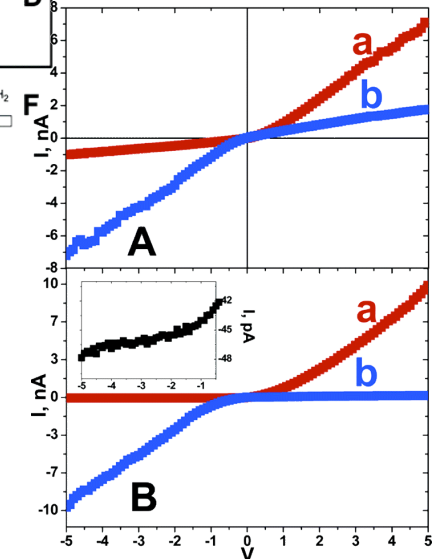
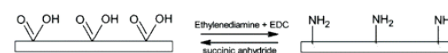
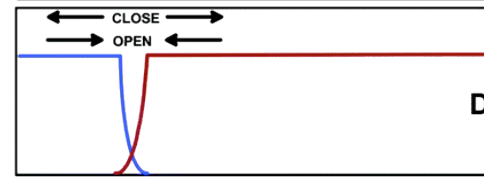
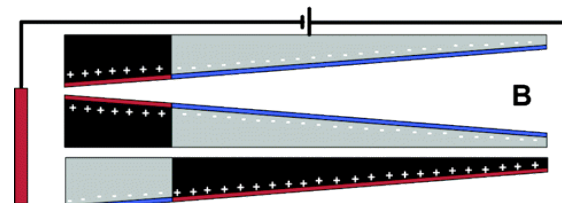
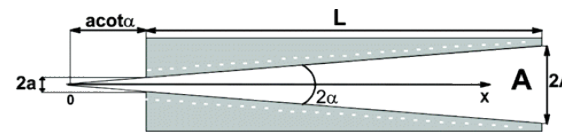
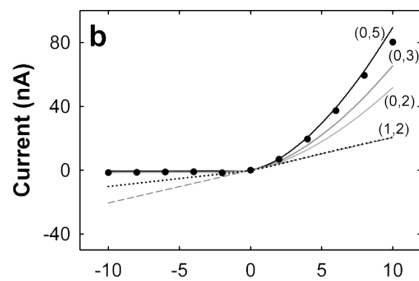
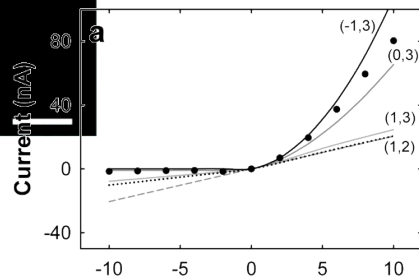
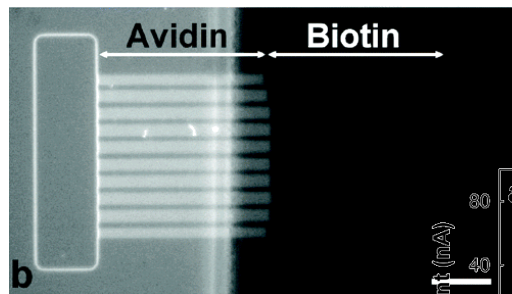
[KCl]=0.1M

Vidal, Gracheva & Leburton,  
Nanoscale Res. Lett. 2, p. 61 (2007)

# Toward Ionic Rectification: Static Dipolar Surface Charges



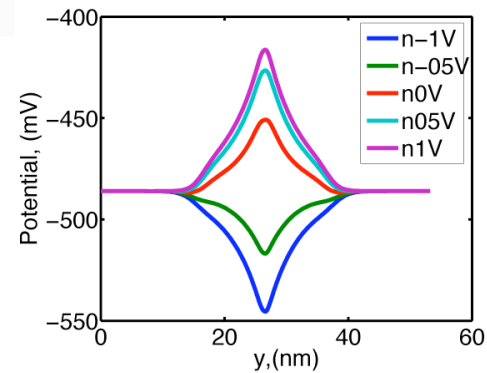
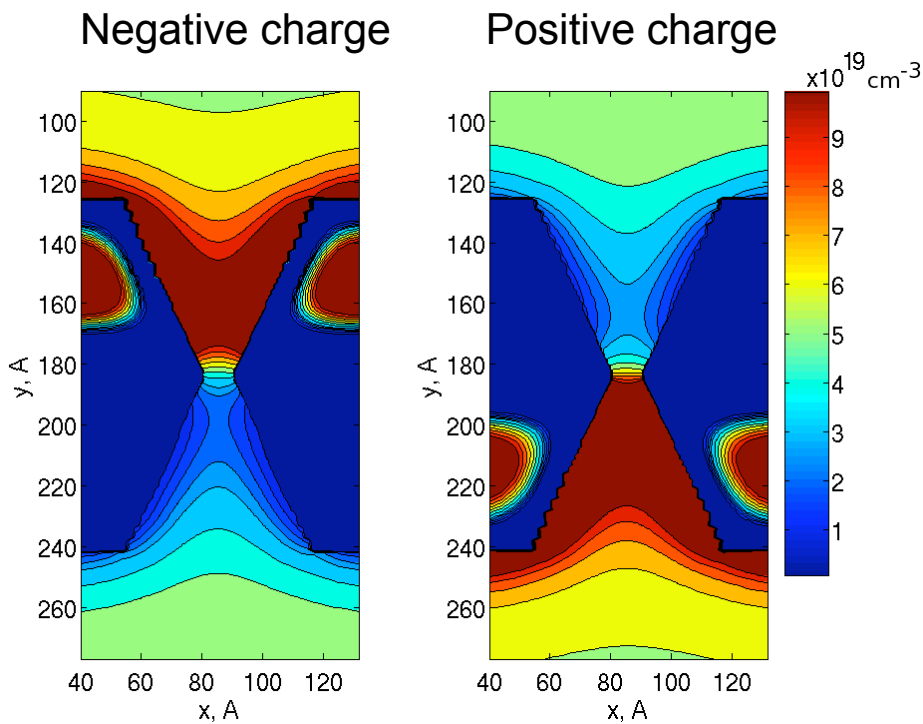
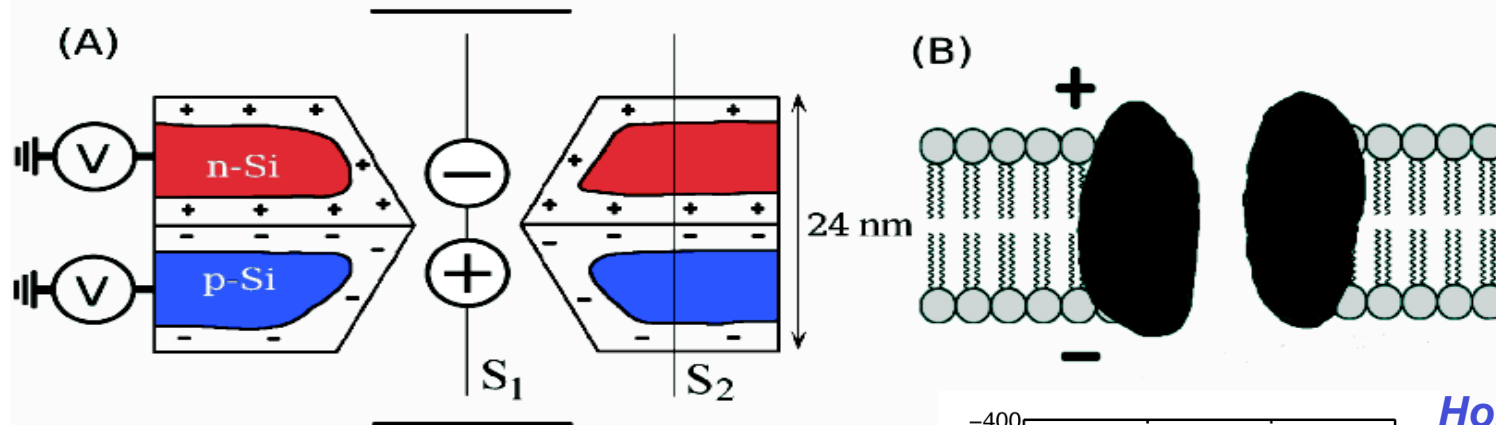
a Avidin Biotin



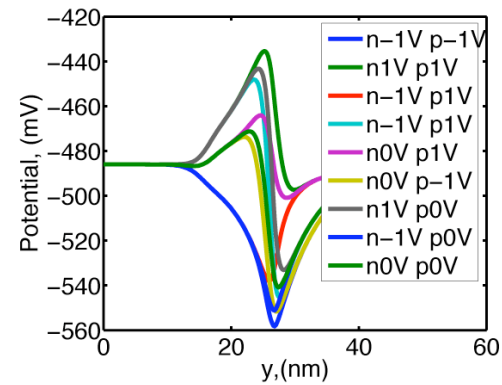
R. Karnik et al. Nanolett, 7, 547 (2007)

I. Vlassiuk et al. Nanolett, 7, 552 (2007)

# P-N Semiconductor Membrane



*Homogeneous n-Si membrane*



*Si PN membrane*

# Ionic Transport Model

Total ionic current:

$$I = \pi r^2 \sum_i z_i J_i$$

Ionic fluxes\*:

$$J_i = -D_i \frac{dc_i}{dx} - z_i D_i c_i \frac{F}{RT} \frac{d\phi}{dx}$$

$$J_i = \frac{z_i F}{RT} \frac{D_i \Delta\phi_D}{(d_2 - d_1)} \left[ \frac{c_i(d_1) \exp(-z_i F \Delta\phi_D / RT) - c_i(d_2)}{1 - \exp(-z_i F \Delta\phi_D / RT)} \right]$$

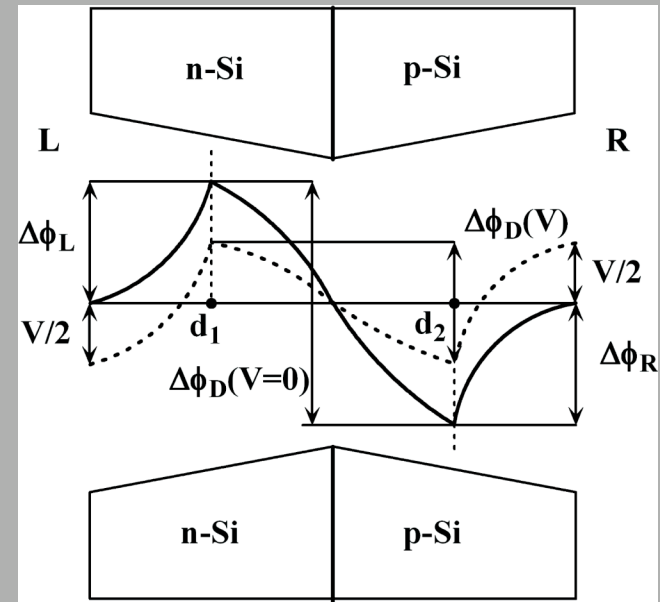
Constant field approximation on segment ( $d_1; d_2$ ):

$$c_i(d_1) = c_{i,L} \exp(-z_i F \Delta\phi_L / RT)$$

$$V = -(\Delta\phi_L(V) + \Delta\phi_D(V) + \Delta\phi_R(V))$$

$$c_i(d_2) = c_{i,R} \exp(-z_i F \Delta\phi_R / RT)$$

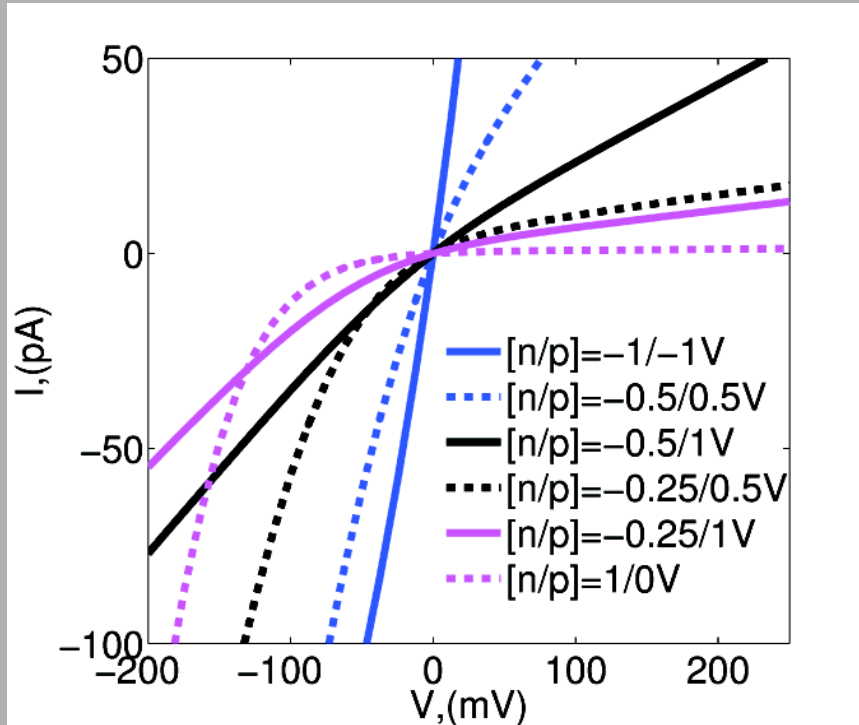
$$\Delta\phi_D(V) = \Delta\phi_D(V=0) - V \quad \leftarrow \text{assume}$$



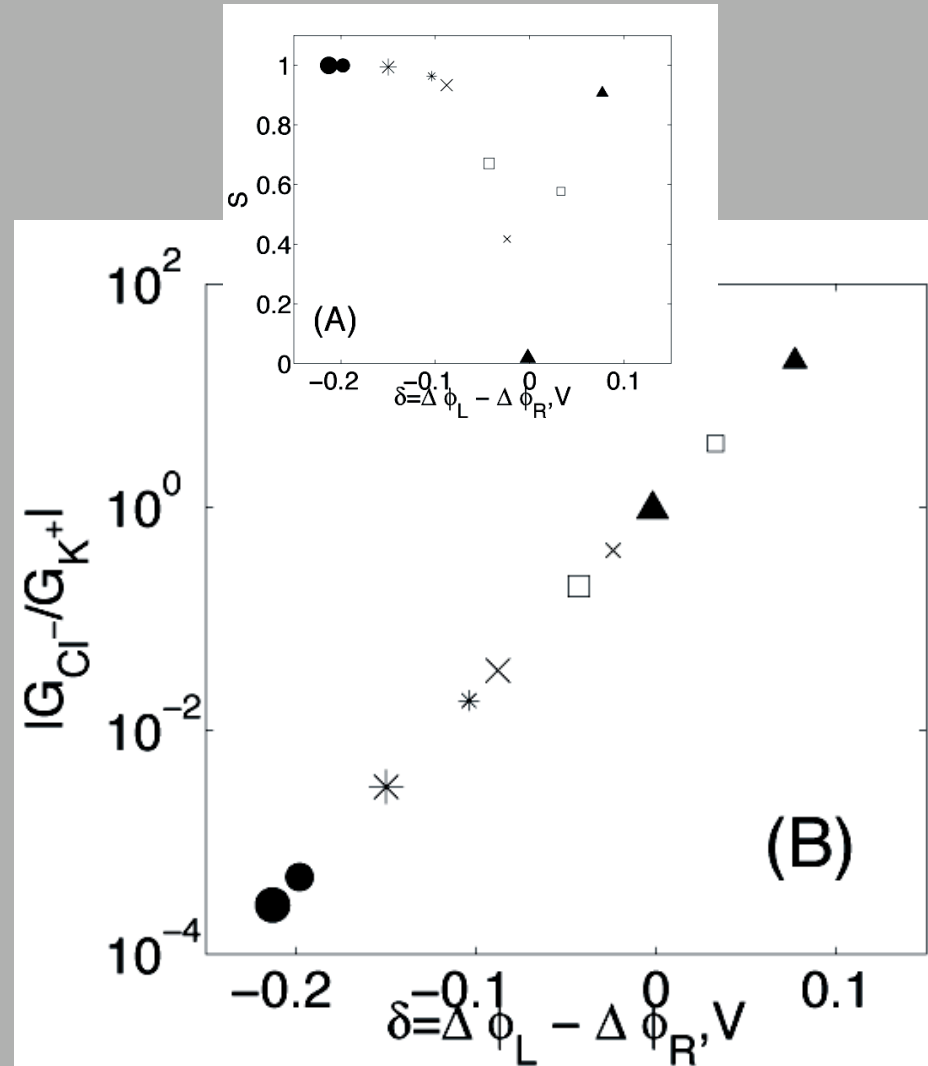
\*After Ramírez et al., Phys. Rev. E 68, 011910(1)-011910(8) (2003)

# Tunable Rectification and Ion Filtering

## Rectification



## Ion selectivity



# Future: fast DNA sequencing?

- Nano-bio-electronic scheme for DNA sequencing
- Integrated multi-scale MD-Self consistent Poisson approach.
- Individual bases electrically detectable, BUT noise and conformation analysis!!!
- Electrical tunability of semiconductor membranes for double layer control.
- PN membranes for tunable rectification and ionic filtering.

