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Vienna University of Technology

# AC Gentechnologie

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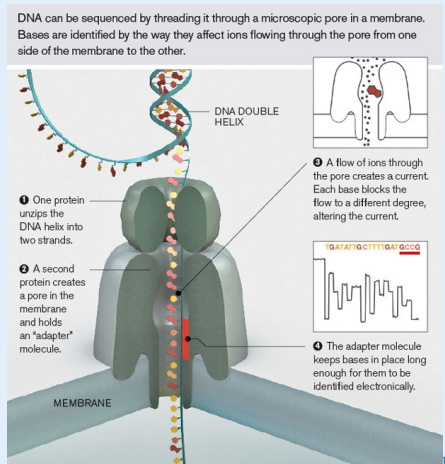
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# Nanopores as sensors

Nanopores can be used as molecular sensors using the principle of a Coulter counter.

Applications include

- ▶ next-generation DNA sequencing (Oxford Nanopore Technologies), [Quick et al. Real-time, portable genome sequencing for Ebola surveillance. *Nature*, **530**:228–232, 2016.]
- ▶ marker-free detection of single molecules. [Burns et al. A biomimetic DNA-based channel for the ligand-controlled transport of charged molecular cargo across a biological membrane. *Nature Nanotechnology*, **6**:152–156, 2016.]

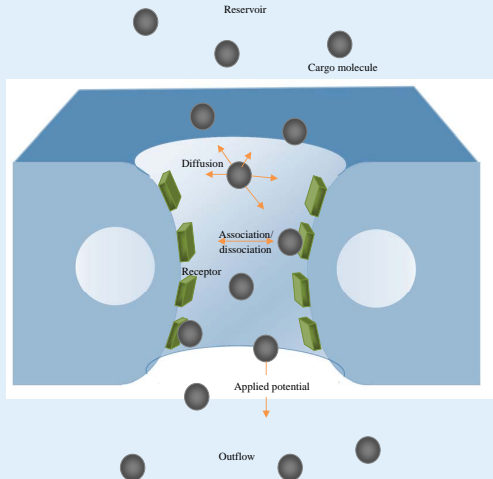


<http://www2.technologyreview.com/article/427677/nanopore-sequencing/>

# Movie: nanopore sequencing

Movie

# Nanopore sensors



# The basic continuum model: the drift-diffusion-Stokes-Poisson system

Drift-diffusion equations for ionic transport,  
(Navier-)Stokes equations for water transport,  
Poisson equation for electrostatic interactions.

$$\begin{aligned}-\nabla \cdot (A \nabla V) &= q(c^+ - c^- + c_m^+ - c_m^-), \\ \nabla \cdot J^- &= 0, \\ \nabla \cdot J^+ &= 0, \\ J^- &= q(D^- \nabla c^- - \mu^- c^- \nabla V - c^- \mathbf{v}), \\ J^+ &= q(-D^+ \nabla c^+ - \mu^+ c^+ \nabla V + c^+ \mathbf{v}), \\ \nabla P &= \eta \Delta \mathbf{v} - q(c_m^+ + c^+ - c_m^- - c^-) \nabla V, \\ \nabla \cdot \mathbf{v} &= 0.\end{aligned}$$

We use 2D and 3D FE calculations to solve this system.

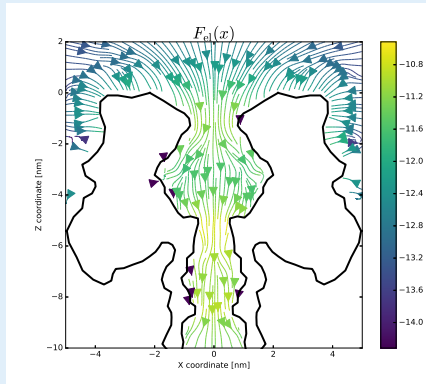
# The force on the target molecules

Force consists of electrostatic and drag contributions:

$$F(x) = F_{\text{el}}(x) + F_{\text{drag}}(x),$$

$$F_{\text{el}}(x) = \int_M E \rho,$$

$$F_{\text{drag}}(x) = \int_{\partial M} n \cdot (-\eta (\nabla u + \nabla u^T) + pI) .$$



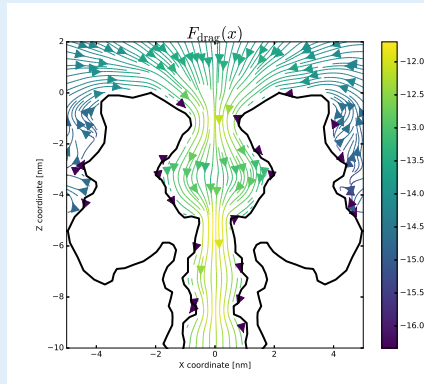
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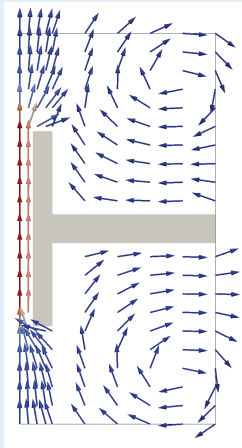
$$F(x) = F_{\text{el}}(x) + F_{\text{drag}}(x),$$

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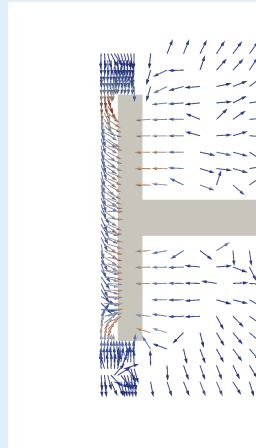
$$F_{\text{drag}}(x) = \int_{\partial M} n \cdot (-\eta (\nabla u + \nabla u^T) + pl).$$



# DNA-origami nanopore: the force acting on the target molecules



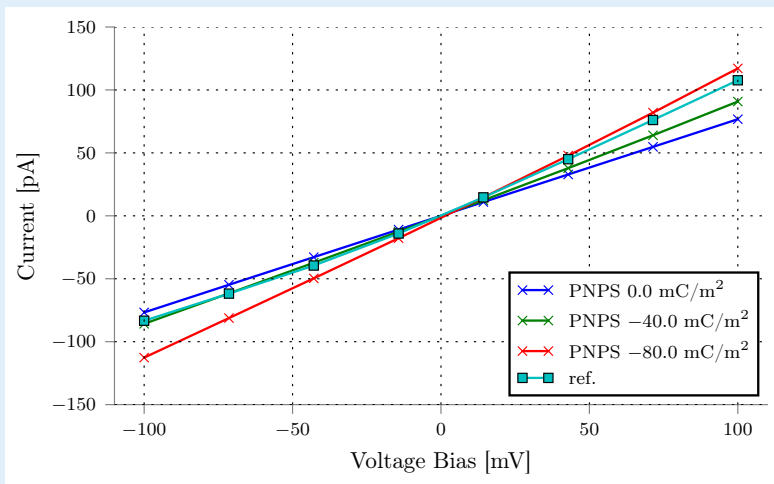
Velocity of liquid



Electrostatic and drag force



# Validation by comparison with experimental data



# Movie: nanopore as single-molecule sensor

Movie

# Continuum formulation of exit-time problem

From the Langevin equation

$$\frac{d\mathbf{x}}{dt} = \sqrt{2D}\xi_t + \frac{D}{kT}\mathbf{F}(\mathbf{x}),$$

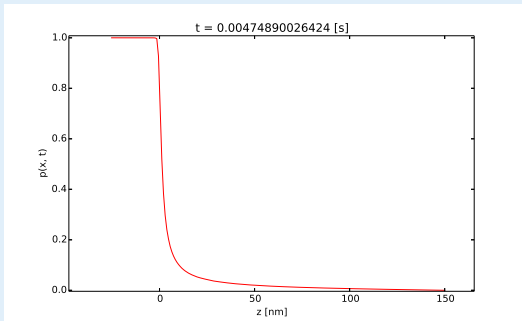
the backward Fokker-Planck equation

$$\partial_t p = \nabla \cdot (D\nabla p) + \frac{D}{kT}\mathbf{F} \cdot \nabla p$$

can be derived. Here  $p(\mathbf{x}, t)$  is the probability that particle has exited.

# Exit-time problem: continuum solution

Probability of **successful exit**: start point  $z$  nanometers above the pore and translocate pore within  $[0, t]$  without hitting boundary far away above pore.



Probability calculated here by solving the backward Fokker-Planck equation with Dirichlet boundary condition (1 means success, 0 means failure) and zero Neumann boundary conditions everywhere else.

Technologischer Fortschritt in der Gentechnologie betrifft mehrere Schritte:

- ▶ Lesen (d.h. Sequenzieren):  
seit Jahrzehnten möglich, wird noch billiger werden
- ▶ Verstehen (Zusammenhang Genotyp-Phänotyp):  
daran wird gearbeitet, z.B. im GTEEx-Projekt (genotype-tissue expression)
- ▶ Editieren:  
seit kurzem möglich, z.B. CRISPR
- ▶ Schreiben (d.h. Gensynthese):  
das nächste Ziel

# Ethische Fragen

- ▶ Für Fortschritte in der personalisierten Medizin ist es notwendig, große (anonymisierte) Datenmenge (inkl. DNA-Sequenzen) zusammenzuführen und auszuwerten.  
Soll dafür die Zustimmung der Patienten notwendig sein? Wem gehören die Daten?
- ▶ Die Kosten mancher Therapien werden substantiell sein.  
Wird man sich die Kosten leisten können? Wer wird bezahlen?
- ▶ Durch Schreiben synthetischer Gene wird man neue Organismen erschaffen können.  
Soll das Editieren und Schreiben von Genen erlaubt sein? In welchen Fällen und für welche Organismen soll es erlaubt sein?

# Danke für Eure Aufmerksamkeit!

## Fragen?

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