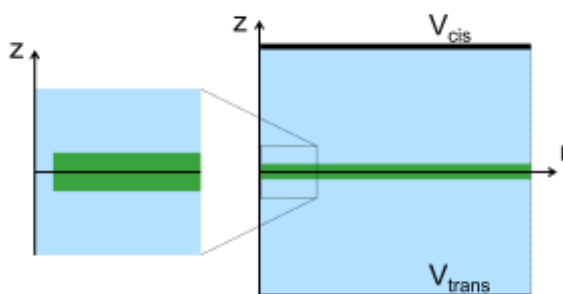


Supplementary Materials:

# Clog and release, and reverse motions of DNA in a nanopore

Tomoya Kubota<sup>1</sup>, Kent Lloyd<sup>1</sup>, Naoto Sakashita<sup>1</sup>, Seiya Minato<sup>1</sup>, Kentaro Ishida<sup>1</sup>, and Toshiyuki Mitsui<sup>1,\*</sup>

Figure S1. FEM based numerical simulation



As shown in Figure S1 for our nanopore simulation geometry, the thickness of the freestanding SiN membrane was set as 200 nm and the pore's inner radius was either 20 nm or 100 nm for 100 nm and 200 nm diameter pores, respectively. The simulation size was  $20 \times 20 \mu\text{m}$  which corresponds to  $0 \leq r \leq 20 \mu\text{m}$  with  $-10 \leq z \leq 10 \mu\text{m}$  in the cylindrical coordinate. As for boundary conditions, the locations where electrical potentials for the cis and trans electrodes are also shown as lines,  $z = 10 \mu\text{m}$  and  $z = -10 \mu\text{m}$ , respectively. The electric fields driving DNA molecules electrophoretically around a nanopore and the possible bulk fluid motions by electroosmosis can be estimated by three equations, Nernst-Planck equation (S1), Poisson equation (S2) and Navier-Stokes equation (S3),

$$\frac{\partial c_i}{\partial t} = -\nabla \cdot (\mathbf{v} - D_i \nabla c_i - \mu_i c_i \nabla \phi) \quad \text{S1}$$

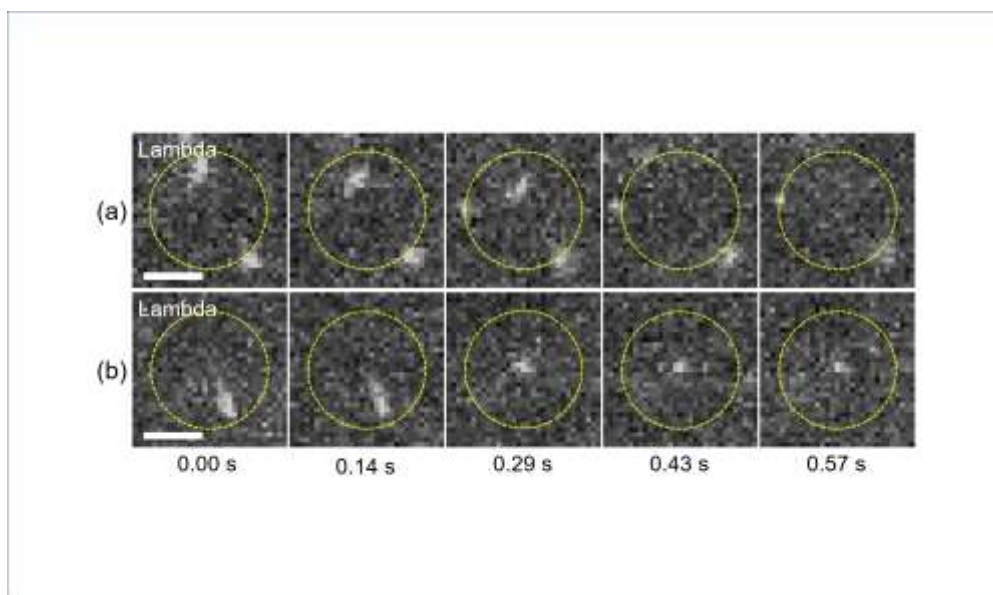
$$\nabla^2 \phi = -\frac{\rho_e}{\epsilon} = -\frac{e \sum z_i c_i}{\epsilon} \quad \text{S2}$$

$$\rho \left( \frac{\partial \mathbf{v}}{\partial t} + (\mathbf{v} \cdot \nabla) \mathbf{v} \right) = -\nabla p + \eta \nabla^2 \mathbf{v} + \frac{e \sum z_i c_i}{\epsilon} \nabla \phi \quad \text{S3}$$

where,  $c_i$ ,  $D_i$ ,  $\mu_i$  and  $z_i$  are the concentration, diffusion constant, ionic mobility and ionic valence for ion kinds  $i$ , respectively.  $\Phi$ ,  $\rho$ ,  $\mathbf{v}$  and  $p$  are the local electric potential, fluid density, fluid velocity and hydrostatic pressure, respectively.  $\rho_e$ ,  $e$ ,  $\epsilon$  and  $\eta$  are the charge density, elementary charge, permittivity and liquid viscosity, respectively. The ionic concentration was increased by 5 mM to compensate the ionic strength of TE buffer in solution. The velocity of the DNA motion driven by

electrophoresis can be estimated by  $v_{\text{ele}} = \mu_{\text{DNA}} \nabla \Phi$  where  $\mu_{\text{DNA}}$  is DNA's mobility. The velocity induced by the fluidic motion i.e. electroosmosis is  $v$  in Eq. S3. The net DNA velocity can be estimated by adding these velocities.

**Figure S2. Translocation and Clog of DNA**



Representative time-resolved fluorescence images of a lambda DNA molecule near a nanopore, displaying (a) translocation and (b) clog. Images are extracted from a sequence of frames recorded at 14 Hz. See also Video S5 lambda DNA translocate and Video S6 lambda DNA clog.

**Table S1. Physical properties of DNA**

DNA	DNA length [kbp]	Diffusion coefficient [ $\mu\text{m}^2/\text{s}$ ]	Mobility [ $10^{-8} \text{ m}^2/\text{Vs}$ ]
Phi X174	5.39	1.5 [1, 2]	3.8 [3]
10 kbp circular	10.0	3.8 [2]	
10 kbp	10.0	3.5	1.1 [4]
$\lambda$	48.5	0.47 [5]	4.2 [5]
T4	166	0.22 [6]	2.8 [7]

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